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Meet the editors



Muhammad Sarwar Khan is a distinguished Plant Molecular Biologist and started his career as a Batchelor and Master's student in horticulture. He earned his Ph.D. from the University of Cambridge, UK. Dr. Khan was awarded a prestigious fellowship to research at the Waksman Institute of Microbiology, Rutgers, The State University of New Jersey, by the Rockefeller Foundation. He has served as the founding Head of the Biotech

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Preface

Citrus is a nutrient-rich fruit crop, predominantly cultivated in tropical and subtropical regions of the world. The *Citrus* genus belongs to the Rutaceae family and consists of a variable number of species due to the admixture of wide morphological diversity, intra- and interspecific sexual compatibility, apomixis, and spontaneous mutations. Citrus fruits are highly nutritious and beneficial for health due to the presence of bioactive compounds that have antioxidant, antitumor, antiinflammatory, and blood clot-inhibiting characteristics. However, citrus production and quality are challenged with several biotic and abiotic problems. Conventional research has played a pivotal role in the improvement of citrus and the introduction of novel biotechnological approaches reduces the time involved in the development of varieties and fixes problems where traditional approaches have failed. Further, transgenic technology and omics approaches have great potential to improve this fruit crop and address biotic as well as abiotic problems.

Citrus - Research, Development and Biotechnology consists of fourteen chapters that are divided into three sections. Section I, "Citrus Genealogy, Classification, and Biotechnology", consists of Chapters 1 through 3. In Chapter 1, Dr. Khan highlights citrus genealogy, production, and crop management. Further, it demonstrates the nutritional and nutraceutical importance of citrus and the biotechnology interventions to improve citrus. In Chapter 2, Dr. Sindhu et al. examine the horticultural classification of *Citrus* cultivars and the sexual and asexual means through which these varieties have been evolved. In Chapter 3, Dr. Khan et al. describe the milestones achieved in citrus improvement employing conventional approaches as well as the achievements made through biotechnology interventions.

Section II, "Citrus Biotic and Abiotic Stress Management", consists of Chapters 4 through 8. In Chapter 4, Drs. Daniele and Etienne explain the importance of weather conditions in cropping systems and substantiate how pathogen and disease spreading is managed through the structural features of the cocoa-based agroforestry system (CBAFS) in the humid forest zones of Cameroon. In Chapter 5, Dr. Lahlali et al. describe two prevailing viroids—Citrus Exocortis Viroid (CEVd) and Hop Stunt Viroid (HSVd)—and stress the use of transcriptomic and proteomic approaches to fully analyze and understand the mechanisms of host-pathogen interactions. In Chapter 6, Dr. Iftikhar et al. highlight that the indexing of diseases caused by a virus and virus-like pathogens is essential for producing disease-free citrus nurseries. Further, they briefly describe the commonly used biological, serological, and molecular tests for the detection of citrus viruses and virus-like pathogens. In Chapter 7, Drs. Caicedo and Villamizar review the main structural and functional features of bacterial responsible for spreading disease and causing symptoms in a susceptible host, including bacterial attachment, antagonism, effector production, quorum sensing regulation, and genetic plasticity at phenotypical and genotypical levels. In Chapter 8, Dr. Naqvi et al. explain the effects of changing climate on citrus and highlight how agronomic, breeding, and biotechnological interventions can mitigate climate change effects on citrus.

Section III, "Citrus Nutritional and Nutraceutical Importance", consists of Chapters 9 through 14. In Chapter 9, Drs. Saeid and Ahmed highlight the intervention of novel approaches in converting citrus byproducts into valuable commodities. Further, they explain the primary and secondary research findings of citrus fruits, especially lemon and pomelo, their chemical properties, composition, and use in health and cosmetic needs. In Chapter 10, Dr. Khanikor et al. discuss the potential and effective use of green pesticides, developed from citrus essential oils, for indoor and outdoor insect management. In Chapter 11, Dr. Fontana stresses the recycling processes of a huge amount of citrus waste as fruit pomace and the outstanding chemical value of peel of Citrus sinensis, rich in useful chemicals, such as polyphenols, polymethoxylated phenols, glycosylated flavonoids, volatile and non-volatile terpenoids, enzymes, and pectin. In Chapter 12, Dr. Minamisawa highlights the importance of yuzu seed-derived nutrients and experimentally demonstrates that limonoid and spermine improve the proportions of beneficial bacteria and their metabolites in the intestinal flora. In Chapter 13, Cavaco et al. review the application of Vis-NIRS in the assessment of the quality and ripening of citrus fruit. In Chapter 14, Demir et al. highlight the use of electrochemical methods to analyze the antioxidant ingredients in food and fruit samples.

Citrus fruits, diverse in color and size, are highly nutritious and beneficial for health because of bioactive compounds such as carotenoids, flavonoids, and ascorbic acid, which have medicinal properties. Thus, this book serves as a guide for students and professionals of biotechnology, medicinal chemistry, and food technologists.

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Citrus Genealogy, Classification, and Biotechnology

Chapter 1

Introductory Chapter: Citrus for a Healthy Life

Muhammad Sarwar Khan

1. An overview

Citrus is an extensively produced fruit crop and is cultivated predominantly in tropical and subtropical regions of the world. The genus Citrus and related genera (Fortunella, Poncirus, Eremocitrus, and Microcitrus) belong to the family Rutaceae. Of these genera, Citrus is the widely grown genus and is well known for fruits like oranges, mandarins, lemons, limes, and grapefruits [1]. The classification of citrus is complexed however, the genus Citrus consists of more than 100 species. The number of species is variable and this species variation in a single genus is due to the admixture of wide morphological diversity, intra- and interspecific sexual compatibility, apomixis, and spontaneous mutations. However, intergenic hybrids such as citranges, citrumelos, and citrandarins, citremons, citradias, and citrumquats are also reported and are getting increasing importance. Several indigenous varieties are developed and consumed locally in specific regions. The citrus fruits are tangy with pleasant flavor and taste, a combination of sweet and sour flavors. Oranges and mandarins are predominant species of genus Citrus, marketed as fresh or processed juice [2].

The citrus plants in the orchards are confronted worldwide with increasing biotic and abiotic factors due to the changing climate. Amongst abiotic factors, fluctuating temperature and unexpected frosts are the main limiting factors whereas, bacteria, viruses, viroids, nematodes, fungi, and phytoplasmas are major biotic factors. Some factors result in a massive reduction in production and quality while others may destroy altogether the citrus industry. Citrus improvement through conventional approaches is discouraged due to the genetic and reproductive characteristics of the plant. The omics and biotechnology-based interdisciplinary interventions may allow combating such external factors and improving the health, nutritional quality of the fruit. The book, Citrus: Research and Development describe the citrus plant, the biotic as well as abiotic challenges, nutrients, and nutritional value, and nutraceutical applications to improve human health. Citrus production, management, detection, and documentation of citrus pathogens and their management, fruit nutritional quality, and potential use as nutraceutical is an interdisciplinary endeavor; therefore, it is difficult to cover all aspects of this subject in a single book. The editor of the book is conscious of the fact that there is considerable scope for improving citrus production and controlling the diseases and benefitting from the availability of chemicals of nutritional and nutraceutical importance, novel approaches for detection of such chemicals, enriching the genetic information through next-generation sequencing and improving the genome by incorporating new genes through genetic engineering and knocking out genes using CRISPR/Cas technology, and hence the information relevant to the topics is covered in the book.

2. Citrus genealogy

Citrus domesticated in Southeast Asia started several thousand years ago and was distributed to different regions of the world through ancient land and sea routes. The genealogy of the modern cultivated citrus is controversial because these are either selections from or hybrids of wild progenitors (**Figure 1**). The biological features and cultivation of Citrus have further complicated the lineage of modern species. Citrus is being cultivated either through clonal grafting or by asexual means of propagation to maintain the identified superior traits however, diversity within such populations is due to somatic mutations. Further, spontaneous mutants have been reported and are occasionally selected from limb sports or nucellar seedlings. The genetic diversity in citrus is the major impediment in the classification of the ever-increasing number of varieties. Several methods have been used to classify the citrus varieties however the methods proposed by Swingle [3] and Tanaka [4] are commonly adopted. Swingle's classification is based on native varieties rather than cultivated, and he placed two subgenera Papeda with six species and Citrus with ten species in the genus Citrus [3], and the rest as natural



Figure 1.

The genealogy and evolution of citrus fruits. Four ancestral species namely; Citrus reticulata, Citrus maxima, Citrus medica, and Citrus micrantha, depicted in the central circle, have contributed to the evolution of cultivated species. The figure is derived from Velasco and Licciardello, 2014 [8] and other published articles.

Introductory Chapter: Citrus for a Healthy Life DOI: http://dx.doi.org/10.5772/intechopen.99355

hybrids of native species. Whereas Tanaka [4] classified both indigenous and cultivated varieties as species and placed two subgenera namely; Archicitrus with 111 species and Metacitrus with 48 species, the list of species was swelled to 159 [5]. The intervention of biotechnology approaches like RAPD, RFLP, AFLP, SSR, SRAP, and more recently the next-generation sequencing and pan-genomics has made it easier to determine the genealogy of citrus, extensively reviewed elsewhere [6]. The genomes of clementines, mandarins, pummelos, sweet oranges, and sour oranges have been sequenced using Sanger whole-genome sequencing method [7]. The sequence data together with earlier similar work was a step forward in elucidating the phylogenetic history of citrus domestication and highlights the genetic basis of the diversity in the colors, flavors, sizes, and aromas of citrus fruits, which could be introduced in novel varieties [8]. Another exciting and challenging work has been published, again by Wu et al. [9], where they worked out that distant genera i.e., Fortunella, Eremocitrus, and Microcitrus are a part of the citrus monophyletic group, whereas a related genus Poncirus, originally believed to be a part of Citrus group, is declared as a distinct clade based on whole-genome phylogeny. These findings have challenged the earlier taxonomic and phylogenetic developments and have warranted reformulation of the genus Citrus.

3. Citrus production and management

Worldwide, citrus is cultivated for consumption as fresh or processed fruit. Major citrus-producing countries are; China, Brazil, USA, Mexico, India, Spain, Iran, Italy, Nigeria, and Turkey. Production and consumption trends are diverse in different regions and countries however, variably 147 million tones citrus is produced, annually [10]. As citrus is grown from subtropical to tropical and the Mediterranean regions of the world, hence, its production is dependent on soil and climate conditions. The global orange production is projected to grow 3.6 million metric tons from the previous year due to favorable weather in Brazil and Mexico. Only a slight increase in production, as well as consumption, is expected for mandarins/tangerines. Unexpectedly, global grapefruit consumption and exports will rise to their highest levels in three years due to favorable weather conditions and expanded areas in China and Mexico. More than 80% of the fruit is being processed for juice production in developed countries, and the juice demand is increasing day by day. The market and consumption trends of major types of citrus fruit as well as of juice are not affected even by the COVID-19 pandemic, this is perhaps due to the perception of the consumers that citrus fruits are immunity boosters being rich in vitamin C.

In addition to climate, plant–soil interaction affects citrus production by affecting the availability of nutrients to the plants [11]. For efficient nutrient availability, the practices of controlled release of fertilizers are preferred [12]. The published data confirm that the nitrogen uptake is greater by using controlled means, compared to conventional approaches, resultantly, the plant growth is improved [13]. Hence, technology-based precision management of orchards including the application of balanced fertilizers, herbicides, and pesticides is required for sustainable citrus production. However, there are several biotic and abiotic factors including diseases that affect the citrus industry. Amongst diseases, bacterial, fungal, and viral are constant threats and cause substantial economic impact in all growing areas around the world. Citrus greening is an extremely dangerous disease, caused by different species of the bacterium Candidatus Liberibacter including Candidatus Liberibacter asiaticus, Candidatus Liberibacter americanus, and Candidatus Liberibacter africanus. Of these three species, the Candidatus Liberibacter asiaticus (CLas) has become a serious threat to the citrus industry. The disease was reported from Brazil in 2004, from Florida, the USA in 2005, in 2007 from Cuba, and molecularl detection of the disease has been reported in 2007 from Pakistan, in 2008 from the Dominican Republic; in 2010 from Mexico [14–18]. However, the disease has been reported and described in China since 2019. The disease is transmitted by the psyllid Diaphorina citri Kuwayama (Hemiptera: Psyllidae), commonly called Asian citrus psyllid. As CLas is difficult to culture on artificial media hence its detection is possible through polymerase chain reaction (PCR) and particularly by quantitative real-time polymerase chain reaction (qPCR) targeting the 16S rDNA gene [14]. Recently, chloroplasts provide opportunities for pathogens to directly or indirectly target 'chloroplast immunity' as these organelles are the main sites for the synthesis of precursors of phytohormones, hence are coordinating plant defense responses. As hormonal crosstalk between host and pathogens is well established. The identification of chloroplast genes targeted by the CLas effectors will open the window to control this disease. The pathogen genome has been sequenced from Diaphorina citri from America [19] and the genome is composed of 1231639 bp with GC 36.5% contents. The pathogen genome has been sequenced from two strains from Pakistan. The CLas genomes of two strains (PA19 and PA20) were sequenced that is comprised of 1224156 bp and 1226225 bp, respectively with an average GC content of 36.4% [20]. The genome sequence of CLas from Thailand has also been reported with total GC contents of 36.4% and 1230623 bp genome. Several genes from the Candidatus spp. have been identified that interact with the genes of the plant defense system and scientists are working to identify the plant genes regulated by these pathogen effectors. The chloroplast being the main site of phytohormone precursor synthesis provides opportunities for pathogens to target, directly or indirectly, the 'chloroplast immunity'. As hormonal crosstalk between host and pathogens is now well established hence identification and editing of chloroplast genes using CRISPR/Cas technologies hold promise to control the disease. Similarly, the canker susceptibility gene, CsLOB1, of Duncan grapefruit has been knockout. The infection by Xanthomonas citri was significantly reduced with no disease development on plants [21].

4. Citrus nutritional and nutraceutical importance

Citrus fruits, diverse in color and size, are highly nutritious. They are beneficial for health, due to the presence of bioactive compounds such as carotenoids, flavonoids, and ascorbic acid [22]. These compounds have antioxidant, antitumor, antiinflammatory and blood clotting inhibiting characteristics [23, 24]. These fruits are also a rich source of vitamins and minerals like vitamin C, A, and B-complex [25–27], vitamin A benefits skin and vision whereas vitamin B-complex like thiamin, folates, and pyridoxine are required as external sources to replenish. Of minerals, potassium, magnesium, calcium, and sodium are present in citrus from very high to low levels in citrus fruits [28]. However, zinc, iron, and manganese are present in trace amounts (**Figure 2**).

Nutraceutical is a combination of two words; 'nutrition' and 'pharmaceutical', hence the word infers that the nutraceuticals could be regulated as dietary supplements, medicine, and food ingredients. The word, 'nutraceutical' was first coined by Stephen L. DeFelice in 1989 who explained nutraceuticals as; "Food, or parts of food, that provide medical or health benefits, including the prevention and treatment of disease" [29, 30]. However, the government of Japan started approving foods with proven benefits for the general public in the 1980s, reviewed elsewhere [31]. Nutraceuticals protect against diseases developed due to nutrient deficiencies Introductory Chapter: Citrus for a Healthy Life DOI: http://dx.doi.org/10.5772/intechopen.99355



Figure 2.

The citrus, a source of nutrients and nutraceuticals. Citrus species are a rich source of carotenoids, flavonoids, minerals, and vitamins that function as antioxidants, antitumor, and anti-inflammatory compounds.

and also have physiological benefits. These are used as dietary supplements and food ingredients. Being dietary supplements, these may contain vitamins, minerals, botanical extracts, essential amino acids, Poly Unsaturated Fatty Acids (PUFA), and enzymes that are probably deficient in most of our diets. Another category of nutraceuticals is nutrient-fortified food. For example, the iron-fortified wheat flour protects the wheat-dependent population from diseases that develop due to iron deficiency. A salient example is iron deficiency anemias. Other examples include purple cauliflower and purple potatoes, having additional anthocyanin content. Golden rice and Golden potatoes as well as Provitamin-A-fortified maize are the rich sources of Provitamin-A and carotenoids. These crops could reduce the disease development in humans caused by Vitamin A deficiency diseases e.g. night blindness, xerophthalmia, prevalent in Africa and Asia. Similarly, Quinoa is of great nutritional value. Laden with fiber, vitamins, and minerals, this plant also is rich in lysine, hence its proteins are nutritionally more complete than many vegetables [32]. Thus, quinoa holds great potential as a Nutraceutical, to curb the malnutrition rampant in many third-world countries, the likes of Pakistan, India, Nepal, Bangladesh, and several African nations.

Several plants, including food crops, have been reported that contain compounds to prevent diseases. Even in this era of rapid medicine evolution, cancer remains a major threat to the population and a leading cause of mortality in developed nations. Introduction of plants into lifestyle at an early age could reduce cancer risk up to 33%. For instance, blue maize is useful in preventing different types of cancers, such as colon cancer [33]. Several chemotherapeutic agents such as Taxol, Vincristine, and Vinblastine are derived from plants such as *Taxus brevifolia* and alkaloids of Vinca species. Nutraceuticals have also been shown to reduce the toxic effects of chemotherapeutic agents and radiation therapies [34].

Bacterial infections and the growing resistance to synthetic antibiotics is a serious concern. It has been proven experimentally that medicinal plants are effective against bacterial infections. In this era of technology development, the introduction of medicinal and nutritional traits transgenically into food crops is on the top priority of biotechnologists, hence engineering the citrus genome will be a better choice.

5. Citrus improvement through biotech approaches

Citrus fruits are highly nutritious and are beneficial for health, due to the presence of bioactive compounds such as carotenoids, flavonoids, and ascorbic acid. However, citrus production and quality are challenged with several biotic and abiotic problems. Biotech interdisciplinary interventions including transgenesis, genome editing, and OMICS could offer solutions to the issues of this fruit crop. Genetic transformation has been established in many citrus species thereby transgenic plants have been developed against bacterial, viral, and fungal pathogens. Equally, OMICS approaches; genomics, transcriptomics, proteomics, metabolomics, interactomics, and phenomics are exploited to improve the citrus fruits. Since, first attempt to manipulate the citrus genome remained unsuccessful hence, the protocols for efficient regeneration from explants like seeds, embryogenic cells, epicotyls, callus, nodal stem segments, and protoplasts, followed by transformation and selection have been optimized for different citrus species. Maximum regeneration potential has been observed in explant 'epicotyl' hence, the epicotyl has been used as an explant for the genetic transformation of citrus plants. Transgenically stable plants were recovered from Agrobacterium treated Duncan grapefruit epicotyls. The recovered plants were confirmed for transgene presence using PCR and Southern blotting techniques. Similarly, transgenic plants using epicotyl tissues as explants were developed from sweet orange, and citrange. The transformation efficiency remained as high as 93%. Hence, transgenic technology is proven as one of the most reliable interventions to genetically improve tolerance/resistance to abiotic/biotic factors in citrus [35, 36].

Using the technology, the nutrition and medicine-related traits have been successfully tailored in citrus fruits. For example, the expression of genes that encode enzymes like phytoene synthase, lycopene- β -cyclase, and phytoene desaturase of the carotenoid biosynthesis pathway have been modulated to supplement human nutrition with vitamin A and antioxidants. The Valencia orange is majorly grown for its juice but the quality of the juice is deteriorated due to the degradation of an enzyme, named thermostable pectin methylesterase (TSPME). Hence, the gene (CsPME4) that encodes TSPME was downregulated to improve the juice quality [37]. Further, an environmentally friendly technology named, 'chloroplast transformation' is available to develop transgenic plants [38–46]. This technology offers several superior advantages like overexpression of transgenes up to 70% due to

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polyploidy at organelle and genome (plastome) levels, accumulation of functional proteins, and natural containment of transgenes since plastids are transmitted to the next generation through ovary, rather than pollens that cause horizontal gene transfer, in most of the cultivated plant species. The chloroplast genome of several citrus species has been sequenced [47–51], thus necessary genetic information of the subcellular organelle is available to develop chloroplast transformation vectors and achieve chloroplast transformation, successfully. Therefore, the development of transgenic plants through chloroplast genome engineering is a promising way forward for cost-effective production of nutraceuticals.

6. Conclusions

Conventional research has played a pivotal role in the improvement of citrus. Enhanced heterozygosity has helped in the development of genetically diverse germplasm in most of the citrus species and numerous varieties have been released for commercial cultivation. However, with the advent of modern biotechnological tools, the period involved in crop improvement through indirect mutagenesis and polyploidization could be further reduced and enhance cost-effectiveness. Transgenic technology and OMICS have great potential to improve this fruit crop.

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Chapter 2

Horticultural Classification of Citrus Cultivars

Jagveer Singh, Vishal Sharma, Kuldeep Pandey, Shahnawaz Ahmed, Manveen Kaur and Gurupkar Singh Sidhu

Abstract

Globally, citrus fruits are grown over an area of 11.42 million ha with 179.0 million tons production. China with 82.7 m tons production is the major producer of citrus fruits followed by Brazil (18.14 m tons) and India (10.53 m tons) (FAOSTAT, 2019). All commercially used scion and rootstock cultivars belong to the genus Citrus, except kumquats, Fortunella spp., and *Poncirus trifoliata*, which are used as rootstock only all over the world. Worldwide citrus cultivars divided into four, reasonably-well-defined horticultural groups: the Sweet oranges, the mandarins, the grapefruits and the pummelos and the common acid members. The true or 'biological' citrus, including species of Citrus (C. reticulata, C. maxima and *C. medica*), share certain characteristics, however, these are clearly differentiated according to the morpho- taxonomic traits. Hundreds of different citrus cultivars are available. Many varieties were chance finds from natural populations, and not the product of intentional breeding efforts. Other varieties in common use have originated from planned citrus hybridization and breeding efforts from worldwide. Most of the readers will be well acquainted with the cultivated types of Citrus scion and rootstocks. This chapter provides ripening season information for worldwide, farmers/gardeners have had success with citrus in many different regions of world where tropical/subtropical climatic conditions occur.

Keywords: true citrus species and its relatives, commercial cultivars

1. Introduction

Generally, there is a strong demand for citrus varieties of superior eating and processing quality. A shortage of supply of consumer-preferred varieties and high prices are the dominant market forces responsible for the revitalisation of the fresh citrus sector. The general demand is for sweet, low acid fruit, with an aromatic flavor. The shortage of supply has meant the acceptance of a range of varieties, some with marginal quality, which it is expected will have a limited commercial potential. Growing citrus in your own backyard and field can be both enjoyable and rewarding! Beautiful green foliage, fragrant blossoms, and delicious, healthful fresh fruit readily available at your doorstep make citrus excellent garden trees. You can choose a citrus variety according to the climate in your area. While this chapter provides ripening season information for worldwide, farmers/gardeners have had success with citrus in many different regions of world where tropical/ subtropical climatic conditions occur. In general appearance and other respects, the citrus fruits of principal commercial importance fall into four, reasonably-welldefined horticultural groups: the Sweet oranges, the mandarins, the grapefruits and the pummelos and the common acid members. The true or 'biological' citrus fruit trees, including species of *Citrus (C. reticulata, C. maxima and C. medica)*, share certain characteristics, however, these are clearly differentiated according to the morpho- taxonomic key of Swingle [1, 2]. Wu et al. [2] found that several named genera (*Fortunella, Eremocitrus* and *Microcitrus*) are in fact nested within the citrus clade. These and other distinct clades that they have identified are therefore more appropriately considered species within the genus *Citrus*. The pulp vesicles contain droplets of oil, which are more abundant in Poncirus, Microcitrus, and the papedas. Fruits of the true citrus species are segmented and fruits of the genera other than *Citrus* are smaller than those of Citrus itself.

Fortunella and Eremocitrus have ovaries with three to five locules, each of which has only two ovules, whereas Citrus, Microcitrus, and Poncirus have ovaries with six to eight locules, each of which contains many ovules. Members of the true citrus fruit trees are generally cross and graft compatible with other members of the group [3–5]. Fortunella (Kumquat) trees, leaves, flowers, and fruits are generally smaller than those of Citrus. Kumquats are adapted to climates that are marginally cool for most of the other members of the subfamily Aurantioideae, they require less heat to achieve fruit maturity and have a certain level of winter dormancy [1]. Eremocitrus and Microcitrus are both endemic to the Oceania region. Both differ from Citrus in having dimorphic foliage and free stamens; however, Microcitrus has an ovary with four to eight locules, whereas Eremocitrus has an ovary with three to five locules. The cold hardiness of Eremocitrus stated in Swingle [1] and Swingle and Reece [6] is in error; Eremocitrus can probably tolerate temperatures as low as -5.5 °C, consistent with the original description of the genus in 1914 [3, 7, 8]. Microcitrus, on the other hand, is considered semixerophytic and able to withstand prolonged periods of drought [1, 6]. Trifoliate orange was considered as a mono-typic genus for many years, represented by Poncirus trifoliata [1], with distinctive trifoliate leaves (unique among the true citrus fruit trees) and deciduous growth habit. This gives to trifoliate oranges the highest degree of cold hardiness among the true citrus fruit trees, surpassing that of kumquats. The adaptation of Poncirus to cold conditions led Swingle [1] to speculate that the remote ancestor of the true citrus fruit trees originated in a tropical or semitropical climate. While the other genera of the true citrus fruit trees remained in these climates, *Poncirus* (or its ancestors) "migrated" to the temperate climate of Northeastern Asia, during which time it developed the adaptations to colder winters mentioned previously. In addition to cold tolerance, Poncirus exhibits many other characteristics that have been and continues to be used in citrus rootstock breeding, notably disease tolerance (including citrus tristeza virus immunity) and dwarfing. For a more complete utilization of Poncirus, the reader is referred to Krueger and Navarro [3, 9, 10]. Relatively recently, a new species, *Poncirus polyandra*, was published [11, 12], which differs from *P. trifoliata* by its larger leaves, some floral differences, and most notably, being evergreen. Perhaps, this latter characteristic is related to its habitat in Yunnan, the southernmost province of China. Clymenia is a very distinctive member of the other true citrus fruit trees. Clymenia was separated from Citrus by Swingle [1] based upon the structure of the pulp vesicles, which are short, plump, blunt, ovoid or sub-globose, sessile or very short stalked, and attached to the side walls of the 14–16 locules. Table 1 summarizes the correspondence between the proposed classification and the former most important ones of Tanaka [13], Swingle and Reece [6], and Mabberley [22] revised by Zhang and Mabberley [14]. Commercially

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Phylogenomic classification	Tanaka (1961) [13]	Swingle and Reece (1967) [6]	Zhang and Mabberley (2008) [14]	Common names (examples)	Phylo-genomic references
<i>C.cavaleriei</i> H. Lev. ex Cavalerie	C. ichangensis Swingle	C. ichangensis	C. cavaleriei	Adsae	[2]
C. maxima (Burm.) Merr.	C. maxima	C. maxima	C. maxima	Pummelos (Pink, Deep Red, Timor,)	[2, 15]
C. medica L.	C. limonimedica Lush.	C. medica	C. medica	Etrog citron	[2, 15, 16]
I	C. medica			Citrons (Corsican, Diamante, Buddha's hand, Humpang)	[2, 15, 16]
C. micrantha Wester	C. micrantha	C. micrantha	C. hystrix DC.	Small-flowered papeda, small-fruited papeda	[2, 15, 16]
<i>C. reticulata</i> var. austeraSwingle		C. <i>reticulata</i> var. austere	C. <i>reticulata</i> Blanco	Sun-Chu-Sha-Kat mandarin	[2]
<i>C. reticulata</i> var. tachibanained.	<i>C. tachibana</i> (Makino) Tanaka	C. tachibana	C. reticulata	Tachibana mandarin	[2]
C. × amblycarpa	C. amblycarpa	C. <i>reticulata</i> hybrid		Nasnaran mandarin	[15]
C. × <i>aurantiifolia</i> var. aurantiifolia	C. aurantiifolia	C. aurantiifolia	C. × aurantiifolia	Mexican, Key, West Indies limes	[2, 16, 17]
<i>C. × aurantifolia</i> var. macrophyllained.	C. macrophylla Wester	<i>C. aurantiifolia</i> (Christm.) Swingle		Alemow	[16]
C. × <i>aurantiifolia</i> var. auratained.	C. aurataRisso	C. <i>limon</i> (L.) Burm. f.	C. × aurantium L.	Adam's apple	[16]
	C. excelsa Wester	C. aurantiifolia		Excelsa and Nestour lime	[16]
С. × <i>aurantium</i> var. L. var. aurantium	C. aurantium	C. aurantium	C. × aurantium	Sour orange, Bouquetier	[2, 15, 16, 18, 19]
	C. myrtifoliaRaf.			Myrtle-leaf orange, Chinoto	[15]
C. × <i>aurantium</i> var. clementinained.	C. <i>clementina</i> hort. ex Tanaka	C. reticulata		Clementine	[2, 15, 18, 19]

Phylogenomic classification	Tanaka (1961) [13]	Swingle and Reece (1967) [6]	Zhang and Mabberley (2008) [14]	Common names (examples)	Phylo-genomic references
<i>C. × aurantium</i> var. deliciosained.	C. deliciosa Ten.	C. reticulata	C. reticulata	Willowleaf, Chios mandarins	[2, 15, 18, 19]
<i>C. × aurantium</i> var. erythrosained.	<i>C. erythrosa</i> hort. ex Tanaka	C. tachibana	C. reticulata	Fuzhu and San huhongchu mandarins	[19]
<i>C.</i> × <i>aurantium</i> var. kinokuniined.	<i>C. kinokuni</i> hort. ex Tanaka	C. tachibana		Kinokuni, Kishu, Huanglingmiao mandarins	[2, 19]
<i>C. × aurantium</i> var. nobilisined.	C. nobilis Lour.	<i>C. reticulata</i> hybrid	C. × aurantium	King mandarin	[2, 15, 19]
C. × <i>aurantium</i> var. paradisiined.	C. paradisiMacfad.	C. paradisi	C. × aurantium	Star Ruby, Marsh, Duncan, etc.	[2, 15, 17, 19]
С. × <i>аиганtium</i> var. paratangerinained.	<i>C. paratangerina</i> hort. ex Tanaka	C. reticulata		Ladu Mandarin	[19]
C. × aurantium var. sinensis L.	C. sinensis (L.) Osbeck	C. sinensis	C. × aurantium	Sweet oranges (Valencia, Washington Navel, Tarroco, etc.)	[2, 15, 18, 19]
<i>C.</i> × <i>aurantium</i> var. suhuiensisined.	<i>C. suhuiensis</i> hort. ex Tanaka	C. reticulata	C. reticulata	Szibat and Se Hui Gan mandarins	[19]
C. × <i>aurantium</i> var. tangerinained.	<i>C. tangerina</i> hort. ex Tanaka	C. reticulata	C. reticulata	Dancy, Beauty mandarins	[2, 15, 19]
С. × <i>аитанtium</i> var. temple ined.	<i>C. temple</i> hort. ex Yu. Tanaka	C. sinensis		Temple tangor	[19]
C. × <i>aurantium</i> var. unshiuined.	C. unshiu Marcow.	C. reticulata clone	C. reticulata	Satsuma mandarins	[2, 15, 17, 19]
C. × latifolia var. nov. 1				India lime	[16]
C. × latifolia var. nov. 2				Kirklime	[16]
C. × <i>latifolia</i> var. latifolia	C. latifolia	C. aurantiifolia	C. × latifolia	Bears, Tahiti, Persian limes	[16]
C. × limon var. bergamiained.	<i>C. bergamia</i> Risso and Poit.	C. aurantiifolia	C. × limon	Fantastico, Femminello, Castagnaro bergamots	[16, 17]
C. × <i>limon</i> var. meyeriiined.	C. <i>meyerii</i> Yu. Tanaka	C. limon	C. × limon	Meyer lemon	[16]

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Phylogenomic classification	Tanaka (1961) [13]	Swingle and Reece (1967) [6]	Zhang and Mabberley (2008) [14]	Common names (examples)	Phylo-genomic references
<i>C</i> . × <i>linon</i> var. limettioidesined.	C. limettioides Tanaka	C. aurantiifolia		Palestinian and Brazil sweet limes and Butnal sweet lemon	[16, 17]
C. × <i>limon</i> var. limettained.	C. limetta Risso	C. limon		Marrakech limonette	[16]
$C. \times limon$ var. limon (L.) Burm. f.	C. limon (L.) Burm. f.	C. limon	C. × limon	Lemons (Lisbon, Eureka, Verna, Luminciana, Interdonato, etc.)	[2, 16]
C. × <i>limonia</i> var. nov. 1				India sweet lime, Indian lemon	[16]
C. × <i>limonia</i> var. jambhiriined.	C. jambhiri Lush.	C. limon	C. × taitensisRisso	Rough lemon	[2, 16, 17]
C. × <i>limonia</i> Osbeck var. limonia	C. limonia	C. limon		Rangpur lime	[2, 16]
	<i>C. karna</i> Raf.			KhattaKharna lime	[16]
C. × <i>limonia</i> var. nov. 2				Voangiala	[16]
<i>C. x limonia</i> var. volkameriana Pasquale	C. limonia Osbeck	C. limon		Volkamer lemon	[16]
C. × <i>lumia</i> var. nov. 1				Bitrouni lime	[16]
C. × lumia var. nov. 2				Fourny hybrid	[16]
C. × <i>lumia</i> var. lumia	C. lumia Risso and Poit.	C. limon		Jaffa lemon	[16]
C. × <i>lumia</i> var. pyriformis ined.	C. pyriformis Hassk.	C. limon	C. maxima	Ponderosa lemon	[16]
C. × microcarpa	C. madurensis Lour.	<i>C. reticulata</i> hybrid	C. × microcarpa	Calamondin, Calamansi	[16]
C. × <i>pseudolumia</i> ined.				Borneo, Barum, Baboon lemons	[16]
Reference: [20, 21].					

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 Table 1.

 Correspondences between the new phylogenomic classification and the former classifications of Tanaka, Swingle and Reece, and Mabberley revised by Zhang and Mabberley.

grown citrus trees such as the varieties discussed in this publication are not grown from seed but are grafted or budded onto a seedling of a rootstock variety. Varieties that are used as rootstocks provide a number of important qualities to the entire tree such as disease tolerance, cold hardiness, soil adaptation, and, to a certain degree, tree size. In the world, most citrus nurseries do not label or identify the rootstock of a tree, but they do select rootstocks that protect trees from important diseases of commercial citrus and are adapted to a range of regions and soil conditions.

2. Varietal groups

Four main varietal groups are distinguished in the international market:

2.1 Citrus sinensis (L.) Osbeck

Sweet orange is the main group which is used both for fresh fruit and processing. It probably originated in China but its major center of diversification is the Mediterranean Basin (**Figure 1**). Major cultivars in this group are classified as navel oranges (Washington Navel, Navelina, Navelate, Powell, Rhode Navel, Cara Cara), blonde oranges (Shamouti, Valencia Late, Hamlin, Pineapple, Trovita, Salustiana, Delta Valencia, Pera), and blood oranges (Tarocco, Moro, Sanguinelli, Maltese).

2.1.1 Navel oranges

Washington (Riverside, Bahia, Baia or Baiana): Is considered to be a limb sport of a variety 'Selecta' in Bahia, Brazil. Tree medium in size and vigor, crown round topped, anthers are without pollen. Rind medium thick tender flesh deep orange, firm less juicy, rich in flavor and taste. Processing quality poor, ships and stores well. Seedless.



Sweet Orange

Figure 1. Sweet orange varieties.

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Navelencia: In growth characters is less vigorous than Washington. Flesh light in color (as rind) firm more juicy, flavor good more than Thompson. Fruit matures earlier than Washington and after Thompson, and hangs well on tree, almost seedless.

Thompson: Limb sport of Washington. Tree compact semi dwarf type, less in vigor than Washington. Rind and flesh less colored, rind smooth glossy pitted finely, flesh firm more juicy, taste and flavor good. Ripens early, seedless.

2.1.2 Blonde oranges

Shamouti (Palestine Jaffa, Jaffa, Chamouti): Tree upright moderate in growth and vigor, branches thick and thornless, petioles with narrow wing. Rind thick leathery smooth pitted, oil glands faint, flesh light orange, firm juicy fragrant and sweet in taste, peels easily. Shipping and storage very good. Mid-season cultivar. Seedless or nearly.

Valencia (Valencia Late, Hart's Tardiff, Hart Late): Has wide adaptation with alternate bearing, tendency to be heavy cropper. Tree large upright vigorous. Matures late. Storage and transportation qualities are very good and hangs well on trees. Seeds few to none.

Hamlin (Norris): Originated as chance seedling and named after the owner of the orchard AG Hamlin. Tolerate cold better than most oranges, productive. Tree medium large with moderate vigor. Matures very early in the season, seeds few to none.

Pineapple: Chance seedling very sensitive to frost, very productive and excellent for processing. Tree medium large with moderate vigor. A mid-season ripening cultivar. Has two limb sport which are seedless. Seedless Pineapple and Blaquemines.

2.1.3 Pigmented or blood oranges

This group of oranges differ from the common sweet orange in that the fruit generally has pink or red coloration on the rind, in the flesh and juice and also has distinct flavor.

Ruby (Ruby Blood): This cultivar was introduced from Mediterranean region. Plants are compact, not very large, moderate in vigor and production. Ripens in mid-season. Rich in flavor and few seeds.

Spanish Sanguinelli (Syn. Sanguinelli, SanguinellaNegra): Originated as a limb sport of Doblefina. Tree foliage light green, spineless, productive small medium in size. Fruit large very attractive with persistent style, late to mid-season in maturity, shipping and storage qualities very good.

Torocco (Tarocco dal Muso, Taracco di Francofonte): An Italian cultivar. Tree medium in size, irregular in bearing but moderate production. Fruits are quite large, variable in shape. Shipping and storage quality good and has mid-season maturity, does not retain quality if left for long on trees. Seeds few to none.

Doblefina (Oval Sangre, Blood Oval): Tree small poor in growth, branches spreading, crown open, foliage light green. Precocious and heavy bearer. Shipping, storage quality good but does not hang well on tree, almost seedless.

Blood Red (Blood Red Malta): Origin unknown probably came from Mediterranean basin. Grown widely and commercially in India and Pakistan. Good coloration is generally attained in the submountain region.

Maltaise Sanguine (Portugaise): Origin of this variety is uncertain but the Egyptian variety Baladi Blood introduced from Malta is said to be Maltaise Sanguine. Likewise Maltase Blood and Bloodred Malta seems to be a clone of this cultivar.Trees are very productive, medium large moderate in vigor. Rind soft easy to peel, taste and flavor excellent, seeds few, storage and shipping quality poor.

2.1.4 Sugar or Acidless oranges

Maltaise Meski: Originated in Tunisia, a non-acid orange cultivar. For all plant, foliage, flower and fruit characters it is like the parent cultivar Maltaise Blonde.

Shamouti Maski or Shamouti Moghrabi: It is a Labenese cultivar which is indistinguishable from Shamouti (Palestine Jaffa) baring that it is acid less (insipid in taste) and more seedy in nature.

2.1.5 Other sweet oranges

Jaffa (Florida Jaffa): It is a clone of Palestine beledi seedling group. Flesh pale orange, tender juicy with pleasant flavor, seeds few. Has good shipping quality but does not hang well on tree.

Joppa: Resembles Jaffa for number of characters and is different in that it starts bearing early and prolifically, branches are stiff and thornless, branchlets are stout. Flesh light orange, soft juicy with rich aroma. Mid-season cultivar, seeds few.

Foster: Tree medium in growth semi spreading foliage dense. Seeds large oval, maturity mid to late season.

Marrs (Marrs Early): Precocious and heavy bearer matures very early. Flesh orange well colored juicy, sweet in taste, acid very low. Fruits hang well on tree and maintain the quality. For good quality and high juice picking should be delayed. Seeds moderate in number.

Parson (Parson Brown): Originated as a chance seedling. It is an early maturing and relatively more seedy. Trees are large vigorous and productive. Rind pitted and pebbled moderately, flesh orange, firm juicy highly flavored.

2.1.6 Indian continents

In India, Mosambi and Sathgudi are invariably placed under sweet oranges as the acid content is very low.

Mosambi (Mosambique): A very popular variety grown commercially in India, early in maturity. Rind quite thick, stripes faint with longitudinal ridges and grooves. Flesh light yellow juicy, acid low, tastes insipid, seeds few.

Sathgudi: Origin is unknown and a popular cultivar grown extensively in India. Tree vigorous produces moderately. Flesh orange, juicy flavor fair sweet in taste, acid very low, mid-season cultivar, moderately seedy.

2.1.7 Sequenced oranges

Chinese box orange (*Severinia buxifolia* (Poir.) Tenore) is native to China and grows as a compact tree or a small shrub. Among the trees related to citrus is the hardiest one. It produces small fruits that have no commercial value and it is used as an ornamental species (IVIA-147).

Amber sweet orange, [(An unnamed hybrid of Clementine mandarin x Orlando tangelo) x unnamed midseason sweet orange seedling], is a variety released by the USDA because of its resemblance to sweet orange, early maturity and deep flesh color. It is the only such hybrid ever legally designated as a "sweet orange", so that its juice could be used to blend with true sweet orange juice, according to juice industry regulations in Florida. All other known sweet oranges are derived only by somatic mutation, not by sexual hybridization, so Amber sweet is not a true sweet orange [23, 24]. There is one true sweet orange (*C. sinensis*) from which many somatic mutants are derived, including Washington navel and blood
orange. The Amber sweet orange is a mandarin x sweet orange hybrid, and not a true sweet orange, as noted above.

2.1.7.1. Sour oranges

Wu et al. [2] reserved the name "sour orange" (*C. aurantium*) to refer to the genome from which cultivar Seville and other somatic mutants are derived. It is the maternal parent of lemon (*C. limon*). The two sour oranges from South China [24] represent two different genomes both unrelated to sour orange (*C. aurantium*).

2.2 Lemon and lime

Lemon and limes are included in the second group. Two main types of limes are distinguished: the small diploid and seedy lime (Mexican) and the big seedless triploid lime (Tahiti, Bears). Several lemon cultivars having major contribution in the world production include Lisbon, Verna, Eureka, Feminello, Fino and Primofiori.

2.2.1 Lemon (C. limon)

There are some distinct fruits in which lemon characteristics are evident, however the differences are to such a magnitude which warrants their separate characterization and classification. In this group, the most important are the karna, the galgal or hill lemon and jambhiri or rough lemon and all of them are widely grown in India. Meyor lemon and the limettas are also lemon like fruits, described below.

Hill lemon (Galgal) C. pseudolimon: An ancient Indian citrus whose origin is unknown and is commonly called as hill or kumaon lemon, grown extensively in the submountain areas along the foothills of Himalayas and in Punjab as a substitute for lime or lemon. It is indigenous to North India grown in sub-Himalayan region. Plant is tall, vigorous, upright and spreading with an irregular and loose crown, fruit ovate oblong, yellow, apex nipped, base rounded or nippled, rind medium, axis hollow, segments 7–11, seed 25–50. Ripening during October–December.

Karna (*C. karna* Raf): Karna Khatta, Karna Nimbu, Khatta Nimbu. A very old Indian citrus fruit of unknown origin, moderately polyembryonic. Considered to be a natural hybrid between rough lemon and sour orange, as the characters exhibited resemble the two species. Widely employed as a rootstock in northern India, second only to rough lemon. Flowers and fruits only once a year. Rind quite thick adhere tightly, golden yellow to deep orange, smooth or ribbed. Flesh orange, texture coarse, semidry, acidic in taste, flavor sour orange like. Seedy, cotyledons white.

Rough Lemon (*C. jambhiri* Lush.), Jatti Khatti, Lemon, Citronelle [Red rough lemon (C x jambhiri (Lush)) Wu et al. [2]. The species is regarded to be native of Himalayan foothills in India, where even today it grows wild. It was thought to be a natural hybrid between citron and lemon. However, Wu et al. [2] was reported that it originated from an F₁ cross *C. reticulata* x *C. medica* by whole genome sequence comparison and is not a true lemon. It is presumed that Portuguese while returning home introduced it in the southeast Africa. Later towards the end of the fifteenth or early sixteenth century it was brought to Europe from where it reached new world. Fruits are acidic, medium sized, shape variable, usually oblate to elliptic oblong. Rind lemon yellow to brownish orange in color, medium thick, surface typically deeply pitted, bumpy (sunken oil glands) deeply pitted or ribbed, separates readily. Flesh pale yellow to pale orange, acidic in taste, juice moderate, segments 10, hollow and large. Seeds many, small highly polyembryonic, cotyledons light green in color.

Eureka (*C. x limon* L. (Burm. f.)): Carvalho et al. [25] were sequenced cultivar Eureka and related somatic mutants. Its seed parent is sour orange and pollen parent

is an unknown citron. It is one of the most important commercial varieties around the world [2]. It is seedling selection of Sicilian lemons. Tree is medium, spreading and having few thorns. Its fruit color is lemon yellow, surface rugose, pitted, shape obovate, size medium, apex round, rind medium thin axis small, solid, segments 8–10, juice acidic with excellent flavor and quality. Eureka is heavy yielder and begins bearing at early age. It has tendency of tip bearing. Rind semi thick pitted, oil glands sunken. Fruiting more in winter, spring and early summer. Seeds few to none.

Lisbon: Originated in Portugal. Its appearance and yield is superior to Eureka. It is resistant to frost, heat and high wind velocity. Tree is large and vigorous with spreading shoots. It has upright thorny growth, lemon yellow fruit color, smooth surface, medium size, pitted rind, small axis, solid, 6–10 segments with 0–8 seeds. Rind quite thick adherence tight, pitted surface smooth and less ribbed than Eureka.

Lucknow seedless: It is hardy, medium, vigorous, spreading drooping, dense foliage, thorny, fruit color yellow, smooth, nippled apex, base round, thin rind, hollow axis, segments 10–12 maturity during November–February.

Plant lemon: Fruit size medium, juicy, heavy fruiting, tolerant to pests and diseases.

Villafranca lemon (CRC 280): It belongs to Eureka group and was introduced into Florida from Europe in about 1875. Originated in Sicily. Commercially not very important compared to Eureka and Lisbon. Tree characters resembles to that of Lisbon, but the plants are more erect or upright in growth, foliage less and with few thorns. In fruit characters, it is similar to Eureka but fruits more during winter like Lisbon.

Nepali oblong (Assam, Nimber or Pat Nebu): An ever bearing lemon cultivar. Plant medium sized, hardy, spreading drooping with irregular crown, fruit shape oblong, color lemon yellow, segments 10–12, the fruit ripen during December– January. Rind relatively thick, greenish yellow, glossy and smooth. Flesh fine grained, greenish yellow, juicy, not too acidic, seeds none to few.

Meyer lemon: Flowers throughout the year, but more so in spring. Tree semi dwarf, thornless, spreading, cold resistant, fruit color light orange, smooth surface, finely pitted, shape obolate or oblong base rounded, rind thin, axis small, 8–10 segments and 8–12 seeds. Rind thin, adhere tightly, surface smooth yellowish orange to orange, flesh light orange yellow, soft, very juicy and typical lemon flavor.

2.2.2 Limes

Like the citron and lemon, the limes likely have originated in north eastern India. Limes are generally of two forms- Small fruited acid limes (*C. aurantifolia*), West Indian lime and the large fruited acid limes (*C. lantifolia*). In its natural habitat, several forms are recognized which differs markedly in size, form, shape, spine and seediness character etc. The West Indian or Mexican lime is the Kagzinimbu and has number of vernacular names. Australian desert lime (Eremocitrus glauca (Lindl.) Swingle, C. glauca (Lindl.) Burkill) is native to Australia and produces fruits of sour taste that can be used as condiment. It is drought tolerant and has very few soil requirements (UCR-12B-38-01). Eremorange, Australian desert lime hybrid (Eremocitrus glauca x C. sinensis) (SRA-871). Australian finger lime (Microcitru saustralasica (F. Muell.) Swingle, C. *australasica* F. Muell), native to Australia, develops elongated finger-shaped fruits of different colors. Juice vesicles that can be broken down and separate very easily are of sharp acid flavor. It is used as a food seasoning (UCR-18B-16-04). Australian finger lime is an accession that we find has Australian round lime admixture. BC2 backcross. (SRA-1002). Australian round lime (*Microcitrus australis*, (Swingle), *C. australis* (A. Cunn. ex Mudie)) native to Australia produces rounded green fruit although at full maturity they become yellow. The pulp has low cohesive juice vesicles as the Australian finger lime. It is used as a food seasoning (UCR-18A-32-01).

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Acid lime (C. aurantifolia (Christm.) Swingle.):

West Indian (Mexican sour lime): It is native of India and widely cultivated in the tropics. Rouiss et al. [26] reported that it is a natural hybrid between micrantha and citron. It is tenderest species among all the citrus species. Tree medium sized, hardy, semi vigorous, upright growth, thorny, fruit round to oblong, yellow apex rounded and slightly nippled, base round, rind thin, papery, 8–10 segments and seeds both. Rind thin, surface fine, smooth, adhere tightly, greenish yellow. Polyembryonic with few seeds, aroma characteristic, flesh greenish yellow, fine, soft juicy, very acidic.

Tahiti lime (Persian lime) C. lantifolia.

Cold resistant, same as that of lemons. It is large fruited acid lime. Flowers throughout the year more in spring. Purple pigmentation present on shoot and flowers. The plants are large, spreading, cold resistant, thornless, fruit large, seed-less, triploid and produce non-viable pollen. It is considered as hybrid between lime and lemon. Fruit color orange yellow, smooth surface and 8–10 segments. It is late variety. Rind thin lemon yellow, adherence tight, flesh greenish yellow, tender juicy acidic. Usually 10 segments.

Rangpur lime (C. limonia Osbeck):

Rangpur lime (*C. x limonia* (Osbeck)) produces non-commercial small and very acidic fruits of orange color. It is indigenous to India and is commonly used as rootstock. Rangpur lime is mainly grown for home consumption and ornamental purpose [2]. Fruits are used for making limeade. It is also known as Marmalade orange. It has loose rind, easily separable segments and pulp is light orange yellow. It is very cold hardy.

Sweet lime (C. limettioides):

In north-eastern India to which it is native it is said that sohsynteng of Assam is the acid form of this fruit. Like acid limes, the Indian sweet lime is the mithanimbu and number of its forms differing in fruitfulness, fruit shape, size and with or without nipple etc. are easily recognized. Generally, sweet lime is grown as a rootstock and for its non-acidic fruits.

Vikram: It was developed at MAU, Parbhani, fruit medium size, heavy fruiting, fruit color golden.

Pramilini: It was also developed at MAU, Parbhani, high yielder, golden fruit color, tolerant to canker.

Sai Sarbati: It was developed at MPKV, Rahuri, high yielder, suitable for summer cropping, tolerant to canker and tristeza.

Jai devi: It was developed at FRS, Periakulum, high yield, juicy, thin peel and pleasant aroma.

2.3 Mandarins

The easy peeling mandarins are becoming more important in the fresh fruit market. Principal in importance in the Orient are the Mandarins, a large, distinctive, and highly varied group that includes some of the finest and most highly reputed citrus fruits. These fruits are commonly referred to as loose-skin oranges. Clementines are the most important mandarins in the Mediterranean Basin, while Satsumas predominate in Japan. Other commercial mandarins include intraspecific or interspecific hybrids such as Fortune, Kinnow, Minneola and several chance seedlings such as Ponkan, Ellendale, Ortanique, Murcott, and Nadorcott (**Figure 2**). In the United States, where the name tangerine first came into common usage, mandarin and tangerine are used interchangeably to designate the whole group. Since mandarin is the older and much more widely employed name, its use is clearly preferable. Presumably because of the orange-red color of the Dancy variety, which originated in Florida and was introduced in the markets as the Dancy tangerine, horticulturists have tended to restrict the use of term tangerine to the mandarins of similar deep





color. Tangerine is applied more strictly to those varieties which produce deep orange or scarlet fruits. Mandarin is known as the mikan of Japan, the suntara or sangtra (numerous modifications) of India, mandarino of Italy and Spain and the mandarine of French-speaking countries.

Due to remarkable diversity of mandarins and the writer's lack of firsthand knowledge of many of the Oriential members, considerable difficulty was experienced in developing a satisfactory horticultural classification for this group. Webber (1948) has separated the mandarin oranges into (a) King group (b) Satsuma group (c) Mandarin group (d) Tangerine group (e) Mandarin-Lime group (f) Mitis group. In this treatment, therefore, the mandarins are presented as the following classes.

2.3.1 The Satsuma mandarins

The Satsuma mandarins (unshiu) mandarin, cv. Owari (UNS) (*C. unshiu* [(Mak.) Marc]; *C. reticulata* (Swingle)) is a commercial midseason, sterile and parthenocarpic, easy peeling mandarin. Satsumas are a group of commercial varieties with relatively high tolerance to low winter temperatures. It is a Japanese variety that was introduced in Florida in 1876. It is also resistant to canker, gummosis, scaly bark and melanose. Plant is thornless having spreading growth habit, orange fruit color, rough surface, oblate to spherical shape, medium size, thin and easily separable rind, flavor rich and seedless. Ripens early than any other oranges as its heat requirement is very low. Number of segments is 10–12, axis hollow and capillary membranes are loose. Fruits should be picked quickly when mature otherwise quality deteriorates and it stores well.

2.3.2 The King mandarins

The King mandarins (C. nobilis (Lour.), C. reticulata (Swingle)) is thought to be a natural tangor, i.e., a hybrid between mandarin and orange, that originated in Vietnam. However, this conventional wisdom is evidently wrong, as Wu et al. [2] was reported that whole genome sequence analysis shows that sweet orange is not a direct parent of King mandarin. King mandarin was first introduced from Cochin China into California in 1882. The king is a prolific bearer, frost resistant and produces high quality fruit. Fruits have had much commercial interest since they are large, develop good flavor when ripe and are of late harvest.

2.3.3 Tachibana mandarin

Tachibana mandarin (*C. tachibana* (Mak.) Tanaka, *C. reticulata* (Blanco)) is thought to be native to Japan and surrounding islands. It develops easy peeling, small fruits of pale-yellow-orange color and acid flavor. Although taste is not completely unpleasant the fruit is not palatable.

2.3.4 Sunki mandarin

Sunki mandarin (sour mandarin, suanju) (C. sunki (Hayata, Hort. ex Tanaka, C. reticulata (Blanco)) produces easy peeling, very acidic small fruits, of an attractive orange color. Its fruits are not palatable and the plants are used as rootstocks.

2.3.5 Cleopatra mandarin

Cleopatra mandarin (C. reshni (Hort. ex Tanaka), C. reticulata (Blanco)) is native to India. It produces unpalatable, small and very acidic fruits. It is widely used as a salt tolerant rootstock and also as an attractive ornamental because of the deep red color of the peel. Plant is thornless with dense top. Fruits are produced singly or in bunch, fruit color is dark orange red, shape oblate, flattened at both ends, size is small with12–15 segments.

2.3.6 The Mediterranean mandarins

The Mediterranean mandarins (C. deliciosa Tenore), which is of principal importance in the Mediterranean Basin. It differs from other mandarins as seeds are plumb and spherical, leaves are small narrow lanceolate and aroma of oil glands and juice is very aromatically flavored.

Willow leaf mandarin (C. deliciosa): The tree is willowy in growth, almost thornless, fruits usually born singly at the tip of slender branches. Fruit color orange, surface smooth, glossy, frequently slightly lobed, necked base, apex depressed, wrinkled, rind thin with 10–12 segments. It is an early variety. Trees are cold hardy but at the ripening time rind separates rapidly and lacks storage ability.

2.3.7 The common mandarins

The common mandarins (*C. reticulata* Blanco), which have worldwide importance and are represented by numerous varieties.

2.3.8 The small-fruited mandarins

The small-fruited mandarins, which are of considerable importance in the Orient and consist of many varieties.

2.3.9 Clementine (Algerian Tangerine)

Clementine (Algerian Tangerine) C. clementina Hort.ex Tanaka: It is a tangerine and is probably an accidental hybrid of the mandarin and sour orange which originated in Algeria. Fruit color is deep orange, shape globose to elliptical, size medium

with depressed apex, rind thick and segments are 8–12, adhered slightly. It is an early variety. Cotyledons are green in color and seeds are monoembryonic.

2.3.10 Kishu mandarin

Kishu mandarin (Kinokuni mandarin) (*C. kinokuni* Hort. Ex Tanaka). The seeded form of this small tangerine grows in southern China and also in Japan, where it was introduced. We sequenced the seedless mutant known in Japan as Mukakukishu; sweet, juicy, and easy to peel, it is appreciated because of its pleasant taste and wonderful aroma. Whole genome sequence comparison shows that it has the same base genotype (i.e., is a somatic mutant of) Huanglingmiao1 mandarin.

2.3.11 Dancy mandarin

Dancy mandarin, Dancy tangerine (*C. tangerine* (Tanaka), *C. reticulata* (Swingle)) is an easy peeling commercial late harvesting variety of excellent color and good size and perdurability on the tree. Originated in 1867 from a chance seedling. In USA, dancy is the best known and highly prized of all the mandarin oranges. Tree is large, nearly thornless and has upright growth. It has tendency towards alternate bearing and seeds are polyembryonic.

Fallglo [Hybrid of Bower mandarin (Clementine mandarin x Orlando tangelo) x Temple tangor, a presumed mandarin-sweet orange hybrid of unknown parentage], a seeded, early maturing and large fruited mandarin hybrid, developed by the USDA and produced primarily in Florida, USA [24].

2.3.12 Calamondin (C. madurensis (Lour.))

Calamondin (C. madurensis (Lour.)): Tanaka has recognized it as a loose skinned orange group. It has less value as a fruit but hangs well on tree and is widely used as an ornamental fruit. It is very cold resistant true citrus fruit and as hardy as Satsuma. Fruit color is orange to deep orange, smooth and glossy surface, pitted shape oblate, size small with flattened base having 7–10 segments.

Sun Chu Sha Kat mandarin (*C. reticulata* (Blanco), *C. reticulata* var. austere (Swingle), *C. erythrosa* (Tanaka)) is characterized by small flowers, small but narrow leaves and small fruits. These are broader than long, peel color may change from yellow to deep red and taste is acidic or acidic-sweet. It is used as rootstock.

Changsha mandarin (*C. reticulata* (Blanco)) produces small, juicy, puffy, brilliant orange-red and seedy fruit. The taste is sweet or acidic-sweet. The tree is rather tolerant to frost and yields heavy crops. It is also grown as an ornamental (UCR-12B-23-07).

Wilking [It is a sister hybrid of Kinnow mandarin both having King mandarin x Willowleaf mandarin parentage], developed by the University of California, Riverside in 1915. Fruit are small, quite fragrant and richly aromatic. Because it produces monoembryonic (zygotic) seeds, it has been used in breeding programs, but not grown commercially to any great extent [24].

Feutrell's early: It is an old variety of New South Wales. Its parents are unknown. The fruit characteristics indicate that it may be a natural tangore and those of the three suggest the possibility that medaterian or Willow leaf might have been the mandarin parent.

Coorg orange: It is an important variety of South India particularly in Coorg and Wynad tracts. Fruits are medium to large, bright orange color, oblate to globose in shape, finally papillate and winkled, glossy with 9–11 segments.

Deshi mandarin (Pathankot): This variety is mainly grown in Punjab hills. The tree is large with semi-upright growth habit with compact foliage and spineless.

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Fruit is ovoid to sub globose, color uniformly cadmium, surface pitted, semi glossy and finally wrinkled, rind medium, adherence slight with 7–10 segments.

Khasi mandarin: It is commercial variety of Assam. Fruit is depressed, globose to oblate, orange yellow to bright orange, surface smooth, glossy, base even or obtuse, rind thin, soft and 8–10 segments.

Nagpur santra (Ponkan, Warnucro): This variety occupies premier position in Indian market and is one of the finest mandarins grown in the world. It is also known as Ponkan. Tree is large, vigorous, spineless with compact foliage. Fruit size is medium, cadmium color, smooth surface, glossy, rind thin, soft, and slightly adhered with 10–12 segments.

Kinnow mandarin: It is first generation hybrid between the King and Willow leaf mandarin that was developed by H.B. Frost at the California Citrus Experiment Station in 1915. It was introduced into Indian Punjab from USA. Tree is vigorous, large, top erect, dense symmetrical with few scattered thorns. Fruit color resembles of King, deep yellowish orange, surface smooth, glossy, very shallow pitted, shape slightly oblate, size medium with flattened base, rind thin, peel tough and leathery, 9–10 segment easily separable and 12–24 seeds. It is a late variety, cold resistant, has strong alternate bearing tendency and seeds are polyembryonic.

PAU Kinnow-1: It is bud mutant of Kinnow mandarin and is low seeded compared to Kinnow.

Tengors: The mandarin-like fruits include the synthetic tangors; the so-called natural tangor, Temple. Kiyomi [Hybrid of Miyagawa-wase satsuma mandarin x Trovita sweet orange], developed by the Okitsu Branch Fruit Research Station, now known as the Okitsu Citrus Research Station, National Institute of Fruit Tree Science. This is a large fruited juicy tangor, with aroma closely resembling sweet orange, and is seedless in the absence of cross pollination. It produces abundant monoembryonic (zygotic) seeds when cross pollinated and has been used as a scion breeding parent in Japan and elsewhere [24].

Temple mandarin: It is a hybrid between Tangerine and sweet orange. Temple mandarin is most beautiful and highly flavored fruit of the citrus group. Tree is medium, thorny, spreading with deep orange to reddish fruit color, rugose glossy surface, medium to large, depressed or nearly flat apex, loose rind, solid axis with 10–12 segments and orange pulp. It is late in maturity.

Dweet: Evolved from a cross between Mediterranean Sweet Orange and Dancy Tangerine. Late in maturity. Tree character intermediate of the two parents. Fruits do not hang well on tree. Fruit reddish-orange in color, globose-oblate, neck distinct, medium to large, surface pebbled. Rind adherence tight and peels very poorly, also puffs, flesh orange, firm very juicy, flavor rich, seeds numerous.

Mency: Originated from a reciprocal cross of Dweet. Fruits ripen early, do not hang well on tree, susceptible to sunburn, small, reddish orange, somewhat oblate and necked, surface pebbled due to oil glands. Rind adherence is not tight and peels easily. Flesh is orange colored and juicy with acidic flavor. Seeds are many.

Tangelos: The mandarin-like fruits include the synthetic tangelos; the so-called natural tangelo, Ugli. Hybrids of mandarin, grapefruit and pummelo are designated as tangelos. They exhibit characters of both the parents.

Clement: It is a hybrid between Duncan grapefruit and Clementine mandarin. Tree and foliage characters are intermediate and productive. Fruit is subglobose to oblong, light orange yellow in color and medium large. Rind is quite thick, pebbled and peels easily. Flesh is dull yellow soft and moderately sweet. Fruit is early to medium in ripening.

Minneola: It is hybrid of Duncan grapefruit and Dancy tangerine. Less cold hardy than Orlando, requires cross pollination for regular and higher yields. Maturity midseason. Rind deep reddish orange, smooth, medium thick, adherence moderate, surface pitted. **Orlando:** A hybrid of Duncan grapefruit x Dancy tangerine. Early in maturity. Fruit broad oblate to subglobose without neck, medium large. Rind, thin orange in color, adherence quite tight, does not peel easily, pebbled. Flesh orange, very juicy tender, somewhat sweet seedy.

Seminole: Parentage is same as that of Orlando and Minneola. More fruitful. Late in maturity, trees are productive, medium large, leaves small to medium, rounded and cupped. Fruit broad oblate deep reddish orange in color. Rind adhere moderately, thin, pebbled, peelable, axis usually hollow. Flesh dark orange, juicy soft, somewhat acidic in taste, seedy.

Page: An early ripening cultivar obtained from a cross between Minneola tangelo x Clementine mandarin.

2.4 Grapefruit and Pummelo

The last group is grapefruit which is divided into the yellow flesh cultivars (Marsh, Duncan) and the red flesh cultivars ('Hudson', 'Star Ruby', 'Ray Ruby' 'Rio Red'). In the Southeast Asia and the Pacific, pummelo (*C. maxima*) and many traditional local mandarin cultivars are still important in the domestic market.

2.4.1 Citrus paradisi Macf. Grapefruit

It is closely related to pummelo, originated in Barbados (West Indies), as old records refer to 'forbidden fruit' (**Figure 3**). It has become popular as a breakfast fruit because of its typical flavor and mild bitterness of the juice. Grapefruit is regarded to be either an interspecific hybrid of pummelo and sweet orange or a hybrid or a mutant of pummel Wu et al. [2]. Seeds are polyembryonic. The fruit is also called as small shaddock, obtained from pummelo, and the name has been derived from the fact that it bears in clusters, and flower resemble that of grape. Like pummelo, grapefruit is also of two types- the common and pigmented grapefruits. The varieties obviously differ in a number of characters i.e., maturity time, seedless or seedy and flavor etc. Important cultivars grown worldwide are described briefly. It is a hybrid between a pummelo and sweet orange. The Cocktail grapefruit is not a true grapefruit.

Common Grapefruit Varieties:

Marsh: (Syn. White Marsh, Marsh seedless, (*C. x paradisi* (Macfadyen)). It is most extended varieties of grapefruit, originated as a chance seedling around 1860



Grapefruit

Figure 3. Grapefruit varieties.

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in Lakeland, Florida. It is a late-ripening, self-incompatible variety that shows long tree storage capability and very good behavior during postharvest. It comes under pallid pulp group of grapefruit. It is bud sport from cultivar Marsh. Fruit color light yellow, surface smooth, shape oblate to globose, size medium to large, basal area small, rind thin, segments 12–14 and 2–5 seeds. It is a late cultivar. Cultivars more significant value is its seedlessness character and its late ripening. Fruit medium sized, spherical or oblate, light yellow, areole ring almost absent. Rind surface even, smooth semi thick, flesh light yellow, soft very juicy, with pleasant flavor, but less than that in seedy cultivars. Has very good storage quality and fruit hangs for long on the tree. Seeds few to none.

Duncan: It was developed as chance seedling in Florida. It is the hardiest variety for cold, fruit color yellow, surface smooth, shape oblate to globose, size large, basal area depressed, apex round, rind medium thick, firm, axis medium in size segments 12–14 and 25–50 seeds. Plants are large, very vigorous and productive early to medium in ripening and also among the few which are cold hardy. Rind surface even/ smooth, glossy, medium in thickness. Flesh light yellow/ buff, soft very juicy, flavor very characteristics strong and pleasant.

Pigmented Grapefruit Cultivars

Foster (Foster Pink): It belongs to pink or red pulp group and originated as bud sport of Walters grapefruit by R.B. Foster in 1906–1907. Fruit color is light yellow, surface smooth, oblate or globose shape, size medium large, base rounded, apex round, rind medium thick with 12–15 segments and 50 or more seeds. It matures during November–December. Cultivar is said to be the first pigmented grapefruit variety that ripens in mid-season, originated as a limb sport of cv. Walters. Fruit medium large in size, round flat to spherical, light yellow overlaid by pink blush, which is also seen in the albedo (mesocarp), areole inconspicuous, furrows at base are small and diverging. Rind smooth, medium thick, flesh light yellow, pink under favorable conditions, soft, flavor good.

Red blush (Syn. Ruby, Red Marsh, Red Seedless): It belongs to pink or red pulp group. It originated as bud sport from Thompson. Deep red color which uniformly distributed throughout pulp. Fruit resembles Thomson for most of the characters except intense pigmentation of the flesh, albedo/ mesocarp also pink with bright red blush on the fruit surface. Hangs well on the tree as the parent cultivar.

Thompson (Pink Marsh): It originated from bud sport to marsh. Fruit color light yellow, surface smooth, 10–12 segments with 2–5 seeds. Plants are very productive large with vigorous growth. Fruits medium sized, spherical-oblate, areole indistinct light yellow, rind relatively thick, smooth and tough. Has excellent storage and shipping quality, fruits hang well for long period of time. Seeds few to none.

Saharanpur Special: Fruit round to oblate, empire yellow, surface small, segments 10–15 and 50–100 seeds. Fruits ripen in November to February.

2.4.2 Citrus grandis (L.) Osbeck. Pummelo, shaddock

Seeds are large yellowish and ridged, monoembryonic. It is native of Polinasia and Malaysia and commonly grown in South China. Fruit is pyriform, largest fruit size among citrus fruits, rind thick, juice is acid bitter, juice sacs easily separable. Seeds are monoembryonic. Fruits are of two types (a) elongated pear shaped with neck (b) Oblate or globose, flattened and neckless. In India there is no improved cultivar except Nagpur chakotra.

In all likelihood is indigenous to the Malayan and East Indian archipelago. In respect of fruit characters, the pummelos fall into two major groups the pigmented and the non-pigmented types. The pigmentation in the former types is caused by carotenoid lycopene and varies from pink to deep red some of which are very attractive with very high in flavor. The non-pigmented (common pummelo) are very variable in number of characters, have moderate to high acid contents and mostly seedy. Description of some known pummelo cultivars is described briefly.

Non-Pigmented Varieties

Banpeiyu: The variety is considered to have originated in Malayan region and is very popular in Japan and Taiwan. Rind thick tightly adhering to vesicles, surface smooth. Flesh pale yellow, soft and juicy, segments many (15–18) central axis solid and large, carpellary membranes thin and tough. Mid-season ripening cultivar, flavor excellent and pleasant taste of acid and sugar, storage quality very good, seedy.

Hirado (Hirado Buntan): Originated as a chance seedling at Nagasaki Prefecture, Japan and rated very highly resistant to cold. Fruit large bright yellow, glossy, oblate both ends slightly depressed, surface smooth, rind medium thick, flesh light green, yellow soft semi dry, flavor pleasant with balanced blend of acid and sugar with some bitterness, segments many, carpellary membranes thin and tough. Seedy.

Kao Phuang: One of the most famous cultivars grown in Thailand. Mid to late in ripening, fruits hang well on tree for long. Tree upright very vigorous. Rind semi thick adherence medium with the vesicles which are firm large and separates easily, juicy, flavor good, capillary membranes semi thick, axis small and compact. Seedy.

Pigmented Varieties

In its place of origin-Orient a number of pink and red fleshed types are present some of which are pomologically described below.

Chandler: Is a hybrid cultivar obtained from a cross between Siamese Sweet x Siamese Pink. Flesh in pink and in other characters it is intermediate between the parents. Rind smooth medium thick, vesicles moderate in juice flesh firm later turns soft, flavor pleasant sub acidic in taste, seeds many.

Ogami (Syn. Egami): This cultivar has performed very well in Florida and other parts of the world. Fruit quite large broadly round flat (oblate), rind smooth shiny and moderate in thickness, pink to deep pink in color, flesh deep pink, pigmentation extends far into the albedo (mesocarp), seedy.

Siamese Pink (Siam): Considered to be one of the finest cultivars in respect of its strong flavor, matures late. Rind adheres tightly medium thick, smooth and glossy. Segments many, carpellary membranes quite thick and tough but open at axis when mature. Almost seedless.

Siamese Sweet: Commonly called as sweet pummelo practically non-acid. Plants are dwarf type, branches drooping, leaves distinctly round pointed. Vesicles separate easily, are large crisp, lack in juice, sweet in taste.

3. Some wild and semi wild species

As north eastern region of India has been recognized as one of the major centre of citrus origin, a number of species are said to be native to this place **Table 1**.

C. indica (Indian wild orange): Specie is found growing in many parts of Assam, Nagaland, Meghalaya and other north eastern parts of India in wild form. Fruit small broad obovoid or subpyriform, appear singly on terminal twigs, about 2 cm in diameter, pedicel very small. Rind very thin, red in color, segments few, orange red in color, inedible, vesicles spindle shaped, very soft, pulp slimy, acidic in taste with unpleasant flavor. Seeds are very large, smooth, monoembryonic and occupy major portion of fruit.

C. assamensis (Adajamir): A very distinct specie known for its peculiar aroma that resembles to ginger or eucalyptus smell that emits from the crushed leaf or from the fruit. A new specie which was indigenous to Assam region. Fruit medium

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small, spherical to round in shape, surface smooth, very acidic juice hence has very limited use as fresh fruit.

The other species *C. latipes*, *C. ichangensis* and *C. macroptera* are usually placed under subgenus Papeda which has number of true wild species of citrus.

C. macroptera (Malanesian Papeda): Almost all the species of Papeda group, this species is most promising as a rootstock. Widely distributed in Indo-China, Thailand, Philippines, New Guinea, New Caledonia and Polynesia. All plants parts are with amber colored glands, spine axillary straight. Flowers are like that of orange. Fruit globose, pale yellow, small (2.5 in dia), smooth, glands small, segments 10–12 with 1–2 seed/vesicle. Juice scanty, highly acidic.

C. latipes (Khasi Papeda): Cold hardy. Native of northeastern Khasi hills (India), and Northern Burma. A small spiny shrub or a small tree (10–15 feet), twigs angular with sharp stout spines (1.2–2.5 cm), reduced or absent on flowering twigs. Fruit medium sized, globular, rind somewhat thick, leathery, segments 9, quite large pulp, vesicles few, spindle shaped, well developed. Seeds many 5–7 in each vesicle, round in shape.

C. ichangensis (Ichang Papeda): Native of south western and west central China. This specie is the most cold hardy of all evergreen types. Differs from other Papeda in that it has large flowers, stamens are connate. Small shrub or a tree (12–15 feet), twigs angular, spines stout. Fruit small (3–4 cm dia) glossy, peel rough, bumpy, medium thick, segments 7–9, each with many seeds large and thick, blunt at both ends, monoembryonic.

Micrantha, Biasong (*C. micrantha* (Wester)) it is thought to be native of the Southeast of the Philippines. It produces small, bitter and inedible fruit with a skin comparatively thick and broadly winged leaves (SRA-1114).

Citron (*C. medica* L).

Citron is considered as native of India having probable region of origin in South Western Asia. Citron is often ranked as the first specie to be cultivated in the western world and in China.

Major citron cultivars are divided into two groups:

- Acid cultivars: Diamante is commercially grown in Italy and Etrog is the main cultivar of Israel.
- Nonacid cultivars: Corsican-well established in California. Sarcodactylis is well known in Japan and China, often referred as fingered citron, because fruit split into number of segments (like finger) with very less pulp.

This citrus fruit is most likely to have originated in the north eastern India and the areas nearby. As the fruit is inedible and tree lacks ornamental value, one ponders why the Romans and Greeks liked it so much. This was the first citrus fruit which was known to the Romans, further the fruit fragrance is soft, penetrating and lasting. Features of some citron cultivars grown extensively are described.

Diamante (Cedro Liscio): A very important and commercial cultivar of Italy, origin is unknown. New growth, buds and flowers are distinctly purple pigmented. Fruit is usually large, long oval to ellipsoid, basal cavity furrowed, apex pointed-nipple like and lemon yellow in color. Rind is very thick, fleshy surface lobed or ribbed and smooth. Flesh is crisp, dry, acidic in taste and seeds are many.

Etrog (Atrog, Ethrog): New growth, flowers and flower buds all are purple pigmented. Trees are small, productive and moderate in growth. Fruit is small to medium, ellipsoid, neck distinct, nipple at apex sharp, style persistent and lemon yellow. Rind is fleshy, thick, surface bumpy/ribbed, flesh firm and crisp, acidic in

taste and low in juice content. Citrus is a highly heterogeneous group and its various species hybridize freely with each other in nature.

Humpang citron (*C. medica* L.) fruit is large, oblong or oval, of green color when growing but generally yellow when ripe. The surface usually is smooth, the rind and the albedo are very thick and the segments are filled with acidic pale greenish pulp-vesicles. Citrons were the first citrus fruit to reach the Mediterranean region and are cold sensitive, monoembryonic, unpalatable and very fragrant.

Mac Veu citron (*C. medica* L., *C. lumia* Risso & Poit). Similar to Humpang citron.

Corsican citron (C. medica L.) is an acidless citron of unknown origin.

Buddha's hand citron var. Sarcodactylus (*C. medica* L. (Noot.) Swingle) produces a very characteristic fruit usually without pulp and split into a number of finger-like sections. This fingered citron is well-regarded because of its fragrance for perfuming rooms and clothing. It is also grown as a dwarf plant for ornamental purposes.

Fortunella margarita Fortunella/Kumquat.

Kumquats are part of genus *Fortunella*, and in Chinese language the word kumquat means 'Gold orange'. This distinct group of citrus has been named after Robert Fortune, the well-known English horticulture worker.

Other species of *Fortunella* are:

- F. crassifolia Swingle (Meiwa kumquat)
- F. margarita Swingle (Oval or Nagami kumquat)
- F. japonica Swingel (Round or Marumi kumquat)
- F. hindsii Swingle (Hongkong wild kumquat)

Pomological description of important Kumquats is briefly summarized below:

Meiwa- Large round Kumquat (*F. crassifolia* Swing.): It is less cold hardy compared to Nagami but considered to be the best as fresh fruit. This specie is the Ninpo, Neiha or Meiwa Kinkan of Japan. A natural hybrid of oval and round Kumquats.

Nagami or Oval Kumquat (*F. margarita* [Lour] Swing): Popularly called as Naga or Nagami Kinkan of Japan. The characteristic features are that fruits are oval, oblong or obovate in shape having fewer number of segments usually 4–5, rind deep orange, both flesh and rind are richly flavored. Tree and leaves are larger in size. Since trees are small and show slow, cold-tolerant growth it is also used as an ornamental. It produces fertile hybrids when crossed with species of the genus *Citrus*.

Marumi or Round Kumquat (*F. japonica* [Thumb.] Swing): Maru or Marumi Kinkan of Japan. Tree is low in vigor with some thorns. Leaves are small and apex is not sharp pointed. Fruit is small round to slightly oblate-obovate. Rind is thin and sweet in taste, segments vary from 4–7.

Poncirus and hybrids.

Trifoliate orange, Poncirus Pomeroy (*Poncirus trifoliata* (L.) Raf.) shows trifoliate leaves and deciduous behavior, two dominant characters that are not present in citrus. The tree also has high resistance to cold. Its fruit has no commercial value and the plant is commonly used as rootstock like its hybrids, especially the citranges, Carrizo and Troyer.

Citranges: It is hybrid between *Poncirus trifoliata* and *C. sinensis* and is hardy than sweet orange. It is nearly deciduous, fruit color varies from yellow to deep orange, surface rough, wrinkled, ribbed or smooth, rind thin, juicy pulp, highly

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acidic. It is used as dwarfing rootstock for grapefruit, Satsuma, sweet orange and lemon.

Varieties: Coleman, Etonia, Mortan, Rusk, Cunnigham, Rustic, Sanford, Savage, Troyer.

Citrangequats: It is trigeneric hybrid of Citrange (Sweet orange x Trifoliate orange) and Kumquat. Citranges are subjected to spring frost injury due to its tenderness while Kumquat is tolerant to the spring frost because of their winter dormancy.

4. Sequenced genome of Citrus spp. and their varieties

Xu et al. [27] reported Valencia sweet orange (*C. sinensis* cv. Valencia): A model of the sweet orange origin. With pummelo as the female parent crossed with mandarin, the interspecific hybrid was backcrossed with mandarin and produced the ancient sweet orange. Haploid Clementine, Clementine mandarin, Ponkan mandarin, Willowleaf mandarin, W. Murcott mandarin, Chandler pummel, Low-acid pummel, Sweet orange, Seville sour orange [18] **Table 1**.

5. Conclusion

Classification of the Citrus varieties has long been debated by taxonomists and botanists. The reticulate evolution combined with partial apomixis has led to very different classification systems. Earlier classification was very difficult due to lacking of genomic study but recently, phylogenomic data revealed the origins and admixtures of modern cultivars and wild types. Coupled with reproductive biology, phylogenomy supports the inclusion of all true citrus of the Swingle system plus Oxanthera in the genus Citrus. The variety rank is defined by the old independent reticulation events from which groups of cultivars were differentiated by asexual mechanisms. It provides an unambiguous conceptual framework for Citrus classification based on the phylogenomic and genetic data. However, today, the available genomic data remain available for further study and further WGS studies are needed to establish a definitive classification of the Citrus varieties. Commercially grown citrus varieties discussed in this text, are not grown from seed but are grafted and budded onto a seedling of a rootstock.

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Conflict of interest

The authors declare no conflict of interest.

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Chapter 3

Citrus Biotechnology: Current Innovations and Future Prospects

Ghulam Mustafa, Muhammad Usman, Faiz Ahmad Joyia and Muhammad Sarwar Khan

Abstract

Citrus is a valuable fruit crop worldwide. It not only provides essential minerals and vitamins but is also of great commercial importance. Conventional research has contributed a lot to the improvement of this fruit plant. Numerous improved varieties have been developed through conventional breeding, mutational breeding, polyploidization and tissue culture yet pathogens continue to emerge at a consistent pace over a wide range of citrus species. Citriculture is vulnerable to various biotic and abiotic stresses which are quite difficult to be controlled through conventional research. Biotechnological intervention including transgenesis, genome editing, and OMICS offers several innovative options to resolve existing issues in this fruit crop. Genetic transformation has been established in many citrus species and transgenic plants have been developed having the ability to tolerate bacterial, viral, and fungal pathogens. Genome editing has also been worked out to develop disease-resistant plants. Likewise, advancement in OMICS has helped to improve citrus fruit through the knowledge of genomics, transcriptomics, proteomics, metabolomics, interactomics, and phenomics. This chapter highlights not only the milestones achieved through conventional research but also briefs about the achievements attained through advanced molecular biology research.

Keywords: citriculture, conventional research, transgenesis, genome editing, multi-OMICS

1. Introduction

Citrus is one of the most diverse members of the family *Rutaceae* and is the leading tree fruit crop in the world. Citrus comprises different species of edible fruits like mandarins (*Citrus reticulata* Blanco), sweet oranges (*C. sinensis* Osbeck), grapefruit (*C. paradisi* Macf.), acid limes (*C. aurantifolia* Swingle), and sweet limes (*C. limettioides*), lemons (*C. limon* Burmf.) and their hybrids including tangerines, tangelos, tangors, etc. [1].

Citrus is being widely cultivated in the sub-tropical, tropical, and temperate regions of the world. Global citrus production is 157 million tons per annum from an area of 15 million hectares. About 50% of the area and production of citrus is being contributed by the northern hemisphere of the world. China (28%) and the Mediterranean regions (25%) are the major contributors to global citrus production followed by Brazil (13%). China is leading in grapefruit and mandarin production. Among Mediterranean countries, Spain is leading in global citrus

production (6 million tons) including mandarins, oranges, limes, lemons and exports. Brazil is leading in global fresh sweet orange and its juice production. Mexico and India are major lime producers [2]. Pakistan's share in global citrus production is quite low (1.6%) which includes mandarin and sweet oranges as major species whereas limes, grapefruit, and lemons have less production and are dealt as minor species. The global citrus industry is facing many biotic (Citrus greening, Citrus tristeza virus, sudden death, citrus canker, and Phytophthora) and abiotic stress (salinity, drought, and temperature

fluctuations) which have a direct impact on fruit crop production and yield [3].

2. Origin and diversification

Citrus and other genera including *Poncirus*, *Clymenia*, *Fortunella*, *Eremocitrus*, and *Microcitrus* belong to tribe Citreae and sub-tribe Citrineae and are considered as true citrus [4]. Classification in citrus has been controversial since ancient times due to vast morphological diversity, interspecific and intergeneric sexual compatibility. However, molecular biology tools have revealed four species including mandarins, citron (*C. medica*), pummelos (*C. grandis* Linn.), and wild cultivar of papeda (*C. micrantha* Wester) as the true parental species that have contributed to the development of other species during the process of evolution [5, 6]. Based on phylogenetic and genomic studies it is revealed that mandarin originated in China, Vietnam, and Japan whereas citron was originated in northeast India and China. Pummelo originated in Indonesia and Malay whereas *C. micrantha* was originated in the Philippines [6]. Other citrus species including sweet oranges, grapefruit, lime, lemon, sour oranges, and hybrids (tangelos and tangors) have developed from these ancestral species through random hybridization and natural mutation events [7].

Among citrus genetic resource centers, major collections are found in the USA, China, Spain, France, Japan, and Brazil where a large number of wild species, their relatives, old and new varieties, and breeding lines are conserved [8]. In Pakistan, citrus genetic resources are conserved mainly in the field as orchards or germplasm units in Sargodha, Faisalabad, and Sahiwal in different academic and research institutes.

3. Conventional approaches for crop improvement

Citrus breeders have been using different approaches for their improvement including conventional breeding, mutation breeding, polyploidization and *in vitro* culture tools particularly somatic hybridization which has played an essential role in developing new somatic hybrids. These techniques have contributed towards the selection and development of new potential cultivars and are still being used as important fundamental tools for the development of genetically diverse germplasm which could be further screened and characterized using modern breeding technologies.

3.1 Classical and mutation breeding

Though conventional breeding has limitations in citrus due to its complex reproductive behavior, nucellar embryony, long juvenility, sterility, sexual incompatibility, and endogametic depression [9, 10]. However, still, many hybrids have been developed by conventional breeding and recovered using *in vitro* tools.

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Mutation breeding has played a pivotal role in fruit crop improvement including citrus and has developed several mutants with improved phenotypic and genotypic traits [11]. Spontaneous or induced mutants do not have intellectual property rights (IPR) related issues that have to be faced in the case of conventional breeding and transgenics [12]. Both spontaneous and induced mutations have enhanced genetic diversity in existing varieties and have provided the raw material for making selections for the novel horticultural traits [13]. About 3365 mutant varieties belonging to 170 plant species have been released including citrus and 20 other fruit species [14]. Among continents, Asia is leading with 2052 mutants released followed by Europe (960 mutants). Among countries, China (817), Japan (479), India (341) and the USA (139) are leading in mutant development whereas Pakistan has released 59 mutants in different crops [15]. In citrus, a total of 15 mutants have been released since 1970 including mandarins and clementine (6), sweet oranges (6), grapefruit (2), and lemon (1) [10]. Pakistan has registered a single mutant variety in citrus, a Kinnow mandarin induced mutant having less number of seeds and named it as "NIAB Kinnow" in 2017.

The rate of spontaneous mutations has been much higher in citrus compared with other fruit crops, however, due to random genetic alterations it has been difficult to identify and utilize such mutants [16, 17]. Induced mutations using different irradiation sources including gamma rays (physical mutagens) and various chemical compounds have enhanced the frequency of genetic variability. Physical mutagens or ionizing radiations have been more commonly used for inducing genetic diversity, chromosomal aberrations, and point mutations. About 70% of the mutant varieties have been developed using physical mutagens [18]. In fruit crops, physical mutagens have altered key horticultural traits like seedlessness, precocious bearing, and dwarfism [19–21]. Other traits include fruit ripening time, fruit skin and flesh color, fruit aroma, self-compatibility, pathogen resistance, and fertility restoration in sterile hybrids. Among physical mutagens, gamma rays have been most used for the development of mutants due to their shorter wavelength and greater penetration [22], however, the ion beam is getting more popular and is being widely used due to its greater efficiency and precision compared with gamma rays [23]. Among chemical mutagens, ethyl methanesulfonate (EMS), diethyl sulfate (DES), ethylenimine (EI), sodium azide (SA) has been most frequently used for reliable and gene-specific mutations. A comprehensive review of the role of mutation breeding in mandarins and lime crop improvement has been discussed [24, 25]. Irradiation and chemical mutagen treatment of seeds and budwood have been commonly used by breeders for inducing variation followed by selection and clonal propagation. Mutation breeding applications have been reported in different fruit crops including papaya, peach, pear, grapes, sweet and sour cherries, banana, plum, almond [26], apple [27], and rough lemon [28]. Natural bud mutants include Washington navel orange, most of the early grapefruit varieties including Marsh, Foster, Shamber, Salustiana sweet orange, and Shamouti orange have originated as bud sports. Now there are several commercial seedless varieties including Daisy SL, Kinnow SL, Fairchild SL, and Tango that have been developed from their seedy parents through mutation breeding and are being commercially cultivated [29]. Other commercial mutants in citrus include sweet orange varieties Jin Cheng [30], Kozan [21], and NIAB Kinnow mandarin [31]. In grapefruit, Rio Red and Star Ruby are two induced mutants that have obtained commercial significance due to their better fruit color and seedlessness, respectively [32]. These are leading grapefruit varieties in Texas, USA. Star Ruby is the leading variety in Turkey, South Africa, Australia, and Spain. Rio Red is the main cultivar in China, India, and Argentine [33]. In Pakistan, Shamber is the main grapefruit variety that needs to be replaced with other potential candidate varieties like Star Ruby, Rio Red, and Flame [10].

Conclusively mutation breeding has shown its enormous potential in citrus crop improvement particularly in economically important horticultural traits. However, it is a slow and long-term process and takes more time to detection of desirable phenotypic variability. Utilization of modern breeding tools including molecular markers, advanced methods for phenotypic screening like Targeting Induced Local Lesions IN Genomes (TILLING) [34], using targeted mutagenesis and genome editing technologies [35] like Clustered Regularly Interspaced Short Palindromic Repeats (CRISPR/Cas9) could enhance the efficiency and cost-effectiveness of variety development having novel traits in citrus.

3.2 Ploidy manipulations

Polyploid organisms have a greater number of chromosomes compared with their diploid progenitors. Breeders have utilized polyploidization for the investigation of inheritance patterns in genes of interest. Polyploids have shown tremendous success in nature due to higher heterozygosity, less inbreeding depression, and more tolerance to biotic and abiotic stress conditions compared with their diploid progenitors [36–38]. The duplicated genes may evolve new functionalization during evolution [39]. Polyploids have been reported in many fruit crops including grapes, apples, strawberry, and citrus [40, 41], however, the frequency of spontaneous polyploid events is quite low and breeders prefer to induce hyperploidy using different chemicals.

Among chemical mutagens, colchicine is mostly used for the induction of polyploids due to its more reliability, higher efficacy, and cost-effectiveness. Colchicine is an alkaloid derived from *Colchicum autumnale* (meadow saffron). It is used for inducing chromosome doubling or developing tetraploids by restricting the chromosomal segregation at metaphase in meiosis [42, 43]. Other methods of polyploid induction include interploid hybridization [44], unreduced gamete formation [45, 46], and endosperm culture [47, 48].

Members of the subfamily Aurantioidae including Citrus, Fortunella, and *Poncirus* are mainly diploid having chromosome number 2n = 18 [49]. The occurrence of spontaneous polyploids in citrus is known since the 1940s [50]. Important spontaneous polyploids include triploid Tahiti lime [51], Triphasia desert lime [52], Clausena excavata [53], tetraploid mandarins [54], sweet oranges [41, 42] and grapefruit [43]. In spontaneous polyploids, triploids and tetraploids are believed to be formed by doubling of chromosomes in nucellar cells and fertilization of the unreduced gametophytes [55, 56]. Polyploids have been induced using colchicine in several citrus species and tetraploids produced have been used for interploid crossing to develop triploid progenies that are usually seedless due to irregular distribution of the chromosomes during cell division particularly gamete formation and formation of unreduced gametes. In interploid crossing, the formation of tetraploids in addition to triploids indicates the predominant formation of the unreduced (2n) gametes which may be formed by the first division restitution (FDR) or second division restitution (SDR) during meiosis. Production of 2n gametes was predominantly via SDR in lemon [44, 45] and monoembryonic Orah mandarin [57]. Higher tetraploid: triploid ratio in the progeny of the interploid hybridization indicates greater production of the 2n megagemtophytes in that cultivar which is promising to produce a greater number of polyploids.

3.3 In vitro culture: somatic hybridization

Plant tissue culture tools offer advantages related to efficient regeneration, propagation, and crop improvement in citrus and other horticultural crops.

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Endosperm cultures have been used for the development of triploids in citrus [48]. In interploid and wide hybridizations, the progeny may be sterile or have underdeveloped or shriveled seeds with viable embryos. The embryo rescue technique has shortened the breeding cycle and many plants have been recovered from these embryos through *in vitro* culture in different citrus species [45]. Similarly, micrografting is another tool in which a miniature bud is grafted under aseptic conditions on *in vitro* raised rootstocks and micrografted plants have been reported in many citrus varieties [58]. Micrografting is also useful for the production of virus-free citrus plants.

Another highly promising and most widely used approach is somatic hybridization which is utilized to overcome sexual incompatibility and to enhance genetic variability by combining nuclear and organelle (chloroplast and mitochondria) genomes followed by their characterization for hybrid confirmation and variability assessment [59]. The organelle genomes are known to encode genes related to photosynthesis and male sterility and new hybrids could be developed having novel genetic recombinations. Somatic hybrids may be developed through electrofusion of plant embryogenic protoplasts predominantly with mesophyll protoplasts. The plant progeny having nuclear origin could be characterized and separated using flow cytometry and molecular markers [60].

Protoplast fusion of distantly related citrus species bypasses the biological barriers and develops allopolyploids that could not be obtained through classical breeding. Somatic hybridization is an important tool and has been widely used in citrus scion and rootstock breeding. The first intergeneric allotetraploid somatic hybrid of Trovita sweet orange and *Poncirus trifoliata* was reported by Ohgawara et al. [61] followed by several interspecific and intergeneric hybrids in citrus from the USA [62], Japan [63], and other citrus-producing countries. Triploids were also reported from interspecific and intergeneric somatic hybridization of Citrus species, kumquats (*Fortunella japonica*), and *Poncirus trifoliata* by protoplast fusion [64]. Fusion of protoplasts from the haploid lines and diploid cultivars may also yield triploids and hundreds of triploids and tetraploids were developed and planted for field evaluations [65]. Polyethylene glycol may also be used to induce regeneration in the fused protoplast cultures as reported in Willow leaf mandarin (embryogenic parent) and Duncan grapefruit and sweet orange (mesophyll parents). The regenerated plants were identified as alloplasmic cybrids [66].

Polyploids developed through somatic hybridization have also shown enhanced tolerance to abiotic and biotic stress conditions. Allotetraploids of cv. FlhorAG1 (FL-4x) developed by somatic hybridization of diploid *Poncirus* and *Citrus* showed greater tolerance to cold and higher light conditions compared with parents (diploid) and their tetraploids [36]. Kumquats (*Fortunella* species) chloroplast have demonstrated higher resistance to canker in diploid kumquats and their tetraploid somatic hybrids developed with other citrus species including grapefruit [37].

4. Innovative approaches/technologies

4.1 Transgenesis

Since the advent of recombinant DNA technology, transgenesis has proved its significance, and 190.4 million hectares of transgenic crops were grown in more than 29 countries in 2019. They have significantly contributed to food security, climate change mitigation, sustainability thus uplifted the lives of 17 million biotech farmers worldwide. The first transgenic plant was developed in the 1980s and was available as commercial food in the 1990s. More than 400 transformation events

have been approved so far wherein 356 events have been approved for crop plants, 23 for ornamentals, 22 for fruit plants, and 2 for trees. Hence a wide range of plant species (maize, cotton, canola, papaya, rice, tomato, sweet pepper, squash, popular, petunia, sugarcane, alfalfa, and citrus) have been engineered for various valuable traits i.e. insect resistance, herbicide tolerance [67], abiotic stress tolerance, improved nutritional value, and disease resistance. In addition to nuclear transformation plastid genome has also been targeted and has proved to be of more value as multiple genes can be introduced at a specific target site, the transgene is contained owing to maternal inheritance, and hyperexpression of the transgene, etc. [68, 69].

Citrus is an economically important fruit crop worldwide. It not only provides essential minerals and vitamins but is also of great commercial importance. Conventional research has contributed a lot to the improvement of this fruit yet serious problems are evolving which are difficult to tackle with these conventional approaches [70]. Juvenility, sexual incompatibility, high heterozygosity, apomixes, large plant size, and nucellar polyembryony, and certain other biological limitations hinder the improvement of these plant species through conventional breeding. Genetic manipulation through advanced innovative techniques is a potential approach to improve crop plants as well as fruit species. Though citrus species are recalcitrant to transformation and subsequent rooting, yet consistent efforts by the researchers have resolved these bottlenecks and proficient protocols have been established. Likewise, various transformation methods i.e. Agrobacteriummediated transformation [71], biolistic transformation [72], and chemically assisted uptake of recombinant DNA by protoplasts [73] have been attempted to introduce genes of agronomic value as well as to strengthen it against bacterial, viral, and fungal pathogens (Figure 1).

Genetic manipulation of vegetatively propagated crops like citrus is very tricky as the expression of transgenes over a long period during numerous cycles of graft propagation should be stable.

The first attempt to produce transgenic citrus was made in the 1980s wherein protoplast transformation was attempted but it was not successful. The first authentic report was published by Kaneyoshi et al. [74] who reported transforming NPT II and GUS genes into trifoliate orange through Agrobacterium. Epicotyls of the aforementioned citrus species were used to transform with the selectable marker gene as well as reporter gene and more than 25% transformation efficiency was achieved. Likewise, Yao et al. [72] reported the first successful transformation through gene gun. They transformed tangelo (*C. reticulata* × *C. paradisi*) embryogenic cells.

Since genetic transformation has successfully been performed in different species and hybrids including Carrizo citrange, Washington naval orange, Poncirus trifoliata, Sour orange, Mexican lime, sweet orange, Citrus reticulata [75], and a valuable rootstock, swingle citrumelo. Similarly, protocols have been optimized for the genetic transformation of different citrus species by using different explant tissues including seeds, embryogenic cells, epicotyls, embryogenic cells, callus, nodal stem segments, and protoplasts. The most responsive explant tissue has been epicotyl from the *in vitro* germinated seedlings and is preferably used for genetic transformation research. Duncan grapefruit was successfully transformed through Agrobacterium for the first time using epicotyl and confirmation of the transgene (NPTII and GUS) integration was carried in the resultant 25 transgenic plants by histochemical staining, PCR, and Southern blot hybridization. Transgenic grapefruit, sweet orange, and citrange plants were developed using epicotyls as target explant whereas selection was carried out on kanamycin [76]. Epicotyl has also been used for Agrobacterium-mediated transformation of citrange and sweet orange [77]. In addition, callus, as well as suspension cultures derived from different parts of flower and seed, have also been attempted to transform. The transformation

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Figure 1.

Schematic sketch showing the importance of conventional and advanced innovative approaches for the improvement of different citrus species.

efficiency attained, in this case, was lower than 0.5%. Genetic transformation has also been optimized in pomelo (*Citrus grandis*) and sour orange wherein internodal stem segments were used as explants and a promising transformation efficiency was achieved (91%) [78].

The biolistic transformation has also been performed successfully in tangelo (*C. paradisi* Macf. x *C. reticulata* Blanco) using nucellar embryogenic cells raised from the suspension culture and more than 15 stable whereas 600 transient transgenic lines were attained per bombardment. The transformed calli cells showed proficient growth on kanamycin selection medium and positive GUS activity but were not able to regenerate into plants. Calli treatment with 0.3 M osmoticum sorbitol and 0.3 M mannitol appeared to have positive effects for enhanced transformation efficiency for stable and transient transformation. Thin epicotyl segments of germinated seedlings were also targeted through the biolistic gun and more than 93% transformation efficiency was attained for transient transgene integration in *C. citrange*. The incubation of explant on culture medium before bombardment appeared to have profound effects on transformation efficiency which was further improved [79].

Since transgenic technology is the most reliable intervention having the massive potential to improve the citrus crop. The introduction of alien genes is only possible through this technology so any of the desired traits can be engineered. Recent research indicates that citrus growing farmers are facing severe problems due to biotic and abiotic stresses i.e. salinity, cold, drought, and diseases. Hence, the development of improved citrus varieties is direly needed to get a quality crop. Various citrus species have been engineered with alien genes to combat abiotic stresses including salinity and drought. Expression of HLA2 gene, isolated from yeast imparted salinity tolerance and resultant transgenic plants were able to tolerate a higher level of the salts as compared with non-transformants [80]. PsCBL and PsCIPK derived from *Pisum sativum* were transformed into *Citrus sinensis* and *Citrus reticulata* by targeting calli derived from mature seed. Bacterial strain LBA4404 was used to induce infection in the target calli cells. The putative transformants showed better performance as compared with control plants for salinity and drought tolerance when tested under *in vitro* conditions [81].

Citrus paradisi was transformed to improve carotenoid content by manipulating the genes involved in carotenoid biosynthesis i.e. phytoene synthase, lycopene- β -cyclase, and phytoene desaturase. The multigene transgenic citrus plants were aimed to supplement human nutrition with vitamin A along with antioxidants. Similarly, fruit juice quality has been attempted to improve in Valencia orange, a valuable variety that is majorly grown for its juice. Degradation of TSPME (thermostable pectin methylesterase) deteriorates the quality of the juice. This TSPME is encoded by the *CsPME4* gene. The protoplasts were isolated from embryogenic suspension cultures and transgene was introduced through the PEG mediated transformation method [82] aiming at down-regulation of the *CsPME4* resulting in the citrus with improved juice quality.

Citriculture is prone to be infected by a wide range of diseases that are controlled by chemicals, a drastic non-environmentally friendly strategy. Different types of viral, bacterial, and fungal pathogens infect these plants resulting in drastic losses to crop production and quality of the produce. A range of transgenic citrus lines have been developed varying from fully resistant to susceptible to the diseases. Coat protein (p25) from the CTV (Citrus Tristeza Virus) was expressed in Mexican lime and 33% of the transgenic plants were found to be resistant, neither symptoms appeared nor the viral load was detected. Accumulation of siRNA (small interfering RNA) in the transgenic lines resulted in resistant phenotype and plants were able to withstand viral infection [83]. Expression of viral coat protein (part of *p23* gene and the 3UTR), in the sense and antisense orientation also delayed viral infection in grapefruit [84].

Phytophthora is a noxious fungal pathogen that has been reported to infect a wide range of citrus species. Among these *Phytophthora parasitica* and *Phytophthora citrophthora* cause more severe damage in the citrus orchards and nurseries all over the world [85]. Expression of *bo* gene (bacterio-opsin) in Rangpur lime rootstock showed an elevated level of tolerance against *Phytophthora parasitica* infection. It was observed that expression of the aforementioned gene led to induce expression of defense-related proteins; chitinase, salicylic acid, and glucanase. Hence plants with an elevated level of transgene expression showed greater resistance against the oomycetic fungi including *Phytophthora parasitica*. Transgenic citrus plants expressing the tomato *PR5* gene showed an enhanced survival rate in the presence of pathogen (*P. citrophthora*). Transgenic grapefruit plants were able to better withstand citrus scab infection when transformed with the *attE* gene encoding for antimicrobial peptide [86].

Transgenic technology has also played a pivotal role to tackle another noxious disease in citrus i.e. Huanglongbing (HLB) which is supposed to be caused by phloem-restricted Gram-negative bacteria; *Candidatus Liberibacter* americanus, *Candidatus Liberibacter* asiaticus, and *Candidatus Liberibacter* africanus [87]. Various genetic strategies have been tested to develop HLB-resistant citrus lines

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with decreased susceptibility to the pathogen. These include the expression of anti-microbial peptides from a bacterial, fungal, plant, or animal origin and engineering host-pathogen interaction pathways. The expression of antimicrobial proteins under phloem-specific promoters has been an effective strategy to control this phloem-resident pathogen. Overexpression of synthetic cecropin B gene under phloem specific promoter resulted in reduced bacterial population after one year of inoculation and no disease symptoms appeared even after two years of inoculation [88]. Overexpression of modified methionine under double 35S promoter also appeared to have an inhibitory effect on bacterial growth and lowered down the CLas (*Candidatus Liberibacter* asiaticus) titer in the roots of transgenic Carrizo citrange (rootstock). Further, newly emerging leaves from the rough lemon, grafted on this transgenic rootstock, also had a non-detectable bacterial titer. Expression of AtNPR1 and chimeric proteins (ThioninLBP and Thionin1-D4E1) demonstrated elimination of CLas providing tolerance against HLB infection [89].

Another economically important disease, the citrus canker has also been addressed through transgenesis resulting in enhanced tolerance against *Xanthomonas citri*. Engineering sweet orange genome with *Xa21* gene showed a significant reduction in disease severity upon inoculation in three lines Hamlin, Pera, and Natal. Expression of the *Xa21* gene under its promoter appeared to be more effective in disease resistance when expressed in highly susceptible Anliucheng sweet orange [90]. Transgenic Carrizo citrange and sweet orange plants were developed by introducing RpfF from *X. fastidiosa* which encodes for a quorum-sensing factor and can disrupt bacterial communication by reducing activation of virulence factors, thus enhancing the ability to tolerate pathogen infection. Similarly, the expression of AMP sarcotoxin from flesh fly also enhanced tolerance against *X. citri* [91].

4.2 Genome editing

Genome editing through CRISPR-Cas9 has emerged as a breakthrough technology for the precise modification and manipulation of targeted genomic DNA. It has extensively been exploited by several research groups [92] and certain successes have been achieved. Three major sequence-specific engineered nucleases that have so far been used for genome editing are CRISPR-Cas (clustered regularly interspaced short palindromic repeats), TALENs (transcription activators such as effector nucleases), and ZFNs (zinc finger nucleases). Among these, the CRISPR-Cas9 editing system has been established in many plant species through gene activation, repression, mutation, and epigenome editing in wheat, rice, maize, tomato, potato, carrot, apple, grape, and citrus. Even a few of the genome-edited crops have been approved for commercial cultivation. Through this technology, field crops as well fruit crops can not only be manipulated for improved agronomic traits but can also be manipulated for improved nutritional value [93].

For citrus, genome editing research is at infancy, yet few successes have been achieved by editing its genome for enhanced resistance against diseases. The CRISPR-Cas9 system was firstly used to target the *CsPDS* gene in Duncan grapefruit and sweet orange. The target gene was successfully modified through a transient expression method, Xcc-facilitated agroinfiltration [94]. The modified *CsPDS* sequence was not detectable in the leaves of sweet orange indicating that CRISPR/Cas9 has induced the desired mutation successfully.

Most of the research studies were carried out to target the *LOB1* (LATERAL ORGAN BOUNDARIES 1) gene which has been characterized as a citrus susceptibility gene for *Xanthomonas citri*. The said gene has been explored to be upregulated by TAL (transcription activator-like) effector PthA4 which binds to the EBE (effector

binding elements) in the promoter region of LOB1 thus activates expression of this canker-susceptibility gene [95]. Mutation in the single allele of effector binding elements of the *LOB1* gene resulted in minor alleviation of canker symptoms in Duncan grapefruit. Anyhow, a mutation in effector binding elements of both of the alleles of LOB1 promoters alleviated canker symptoms to great extent thus showing a high degree of resistance in Wanjincheng orange [96]. Another research group explored that editing the coding region of LOB1 in Duncan grapefruit, through the CRISPR-Cas9 system also provides resistance against *X. citri* infection.

A marker gene for pathogen triggered immunity (CsWRKY22) was knocked out in Wanjincheng orange and the resultant mutant plant showed a decreased level of susceptibility to the canker disease [97]. In addition to the CRISPR-Cas9 system, another improved genome editing system (CRISPR/Cas12a (Cpf1) has also been used to edit the *CsPDS* gene in the Duncan grapefruit gene. It appeared to be a more efficient editing system with lower off-target effects thus will prove a great milestone in citrus genome editing [98]. These studies indicate that CRISPR-mediated genome editing can be a promising pathway to generate disease-resistant citrus cultivars [99].

5. Multi-Omics technology: An integrated approach and useful strategy for the improvement of the citrus crop

MultiOMICS including genomics, transcriptomics, proteomics, metabolomics, interactomics, and phenomics approaches have massive potential for citrus improvement just like other crops and fruits. In all disciplines of OMICS, various techniques can be utilized for genome analyses, transcripts, proteins, metabolites, and interactions between different molecules to indicate the molecules which may result in crop improvement.

5.1 Genomics

The field of genomics is a highly applicable part of Omics technologies. It is based on sequencing technologies and the analysis of subsequent genome sequences. Many advanced techniques in genomics for example sequence determination DNA, marker-assisted selection, and transition from marker-assisted selection to genomic selection assist in quick varietal development. Genome sequencing technologies have brought about a revolution in the field of biology. It has also transformed the citrus breeding that helps to understand a relationship between the genetic makeup and response towards various abiotic and biotic stresses like Alternaria brown spot [100].

A specific pathotype of *Alternaria alternata* (Fr.) Keisel is a disease with heavy losses [101]. It causes necrosis and resultant lesions on the surface of fruits and young leaves. It leads towards defoliation and fruit drop [102]. Thus, exploitation of innate genetic resistance appears to be the most applicable and effective strategy of disease control. Currently, the control is primarily based on the application of 4–10 sprays of toxic and environmentally injurious fungicide per year [103]. Such limitations are compelling farmers to find alternative cultivars with resistant ones [104].

Usually, the female parent transmits the 2*n* gamete in 2x × 2x citrus crosses [105–108]. Cuenca et al. [109] recognized ABS resistance locus containing genomic region by using BSA-genome scan combined with HTA based. Trait segregation in crosses between two heterozygous ABS-susceptible or between heterozygous ABSsusceptible parents and resistance was used to confirm the recessive inheritance of the ABS resistance in triploid populations. ABSr locus was first located at 10 cM from a

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centromere based on segregation of 368 SDR 2*n* gametes. A genomic region containing several markers with a high probability (> 99%) of association with phenotype variation was identified on chromosome III by performing BSA over 93 triploid hybrids from a Fortune × Willow leaf population. This identified region contains 25 significant SNP markers within an interval of 13.1 cM. The size of the genomic region among these two markers is 15 Mb. Linkage genetic mapping was performed on identified genomic regions by developing new SNP and SSR markers. A 268-diploid mapping population was performed by Cuenca et al. [110] from a heterozygous-susceptible × resistant hybridization. Fine mapping was performed to confirm the location of *ABSr* locus in a region of 1.1 cM between the markers SNP05/SNP06/SNP07/AT21 (at 0.7 cM) and SNP08 (at 0.4 cM). Another region containing eight genes with NBS-LRR repeats was identified by the SNP08 marker and considered ABS resistance genes.

In citrus plant, molecular markers are linked to some agronomic traits, e.g. SSR markers are linked to Citrus tristeza virus resistance from Poncirus trifoliate, PCR assay for the anthocyanin content of pulp [111], AFLP markers are associated with polyembryony [112] and RAPD markers are associated to dwarfism and fruit acidity [113]. Some other characteristics such as salinity tolerance and nematode resistance are linked to QTLs [114]. The selection of resistant genotype at the early growth stage was improved by the newly developed SNP08 marker. This marker was mapped at 0.4 cM from the ABD resistance gene and it has role in avoiding the selection of susceptible varieties. On the other side of the gene, some new markers were also identified at 0.7 cM from the ABSr locus. Combining these new markers with SNP08, the probability of selection of resistant genotype was increased by 0.0028%. This marker appeared to be very helpful in the selection of resistant and susceptible genotypes and for analyzing the resistant germplasm to configure the ABS genes. So, it is a very valuable tool for the selection of susceptible heterozygous cultivars which may be used as breeding parents allowing manipulation of genetic diversity in citrus and prevents susceptible homozygous genotypes.

About 40 mandarin genotypes (susceptible and resistant) were tested by the SNP08 marker and were used as breeding parents. An ultimate association was observed between response to *Alternaria* infections and SNP08 marker. Recently SNP08 is used in breeding programs of citrus performed at CIRAD and IVIA for the selection of ABS-resistant citrus genotypes. About 2187 resistant hybrids were selected from 4517 total hybrids rising from 10 different parental combinations by using the SNP08 marker since its development. This analysis was very helpful to prevent the growth of more than 2000 susceptible lines which were removed at the early growth stage after selection so, a lot of time, cost, personnel, and resources were saved.

5.2 Proteomics and metabolomics approaches

Proteomics is the comprehensive analysis of all the proteins found in a cell. It includes the identification of proteins, their location in the cell, their interactions with other proteins and other biological components in the cell, and most importantly post-translational modifications that a protein undergoes in the cell [115]. Metabolites are referred to as the last product of any biological activity in a cell and are found in very small quantities [116]. Metabolites are small molecules including intermediates of various metabolic reactions, signaling molecules, hormones, and other regulatory products found in a cell. Hence, metabolomics is defined as the study of metabolites of a cell [117, 118]. It is estimated that around five thousand metabolites are found in any cell depending upon the physical and chemical complexity of that cell [119].

Huanglongbing (HLB) is considered one of the most devastating citrus diseases that affect not only the production but also the quality of citrus fruit and its juice.

Using a combination of proteomics and metabolomics approaches it was found that in symptomatic fruit, the expression of proteins found in the cytoplasm for glycolysis, in mitochondria involved in the tricarboxylic acid (TCA) cycle, and in chloroplasts for the synthesis of amino-acids was downregulated. Similar downregulation was observed for genes involved in terpenoid metabolism for example valencene, limonene, 3-carene, linalool, myrcene, and aterpineol in fruit found on infected trees. Similar phenomena were observed for sucrose and glucose. Hence, the off-flavor found in symptomatic fruits was linked to the downregulation of the above genes and a decrease in the levels of the abovementioned secondary metabolites [120].

In another study, comparative iTRAQ proteomic profiling was carried out using the fruits of sweet orange which was grafted on sensitive and tolerant rootstocks infected by CaLas. The results showed that symptomatic fruit on sensitive rootstock exhibited a greater number of differentially expressed (DE) proteins as compared to the healthy fruit on a similar rootstock. It was also found that the expression level of various defense-related proteins was reduced in symptomatic fruit on sensitive rootstock, particularly the proteins related to the jasmonate biosynthesis, is signaling, protein hydrolysis, and vesicle trafficking. Hence, it was concluded that the down-regulation of these proteins is likely to be linked with the sensitivity of citrus to the CaLas pathogen [121].

5.3 Interactomics and metabolomics and phenomics

Interactomics bears a broad scope as it may cover a complete set of interactions in a cell [122]. It covers every type of interaction among interacting molecules including proteins and other molecules. It is a well-known fact that the Protein–protein interactions are major of all cellular processes [123].

To designate the complete phenotype of a plant, the term phenome is used. Similarly, a phenotype encloses a group of traits that are liable to be distinguished either by utilizing modern science analytical techniques or by a naked eye evaluation. These traits can also be attributed to being an interaction between external factors (environment) and Genotype. David Houle also termed phenomics as the collection of data from varying backgrounds and dimensions in a single entity [124]. Phenomics involves both "extreme phenotyping," referring to a comprehensive selection of a wide range of valid and correct phenotypes, and "phenome analysis" indicating towards an analysis of specimen and correlation between syndication of genotype and phenotype.

Plant phenomics utilizes screening of large populations to analyze genetic mutations found in the population for a specific trait (drought, salinity, or high-temperature stress tolerance). Various types of imaging techniques are employed in the phenotyping of plants for various growth and developmental processes. The techniques include visible-light imaging [125], Thermographic imaging [126], Hyperspectral imaging, Chlorophyll fluorescence, X-ray, MRI, PET [127].

Using the phenomics approach and tools, we can study the traits regarding plant growth, leaf growth, root growth, and architecture of soil/root interaction, etc. This extensive use of phenomics and its integration into OMICS is the need of the hour to combat food security issues and overcome adverse effects of climate change on crop production.

6. Conclusions

Conventional research has played a pivotal role in the improvement of citrus. Enhanced heterozygosity has helped in the development of genetically diverse

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germplasm in most of the citrus species and numerous varieties have been released for commercial cultivation. However, with the advent of modern biotechnological tools, the period involved in crop improvement through indirect mutagenesis and polyploidization could be further reduced and enhancing cost-effectiveness. Transgenic technology and OMICS have great potential to improve this fruit crop. MultiOMICS, integrative-OMICS, or panOMICS technologies may result in better crops having better agronomic traits, enhanced yield potential, and less prone to insect pests. It will ultimately lead towards food security and poverty alleviation. Various OMICS technologies have been used for crop improvement, yet their integrated use will further strengthen the application of this robust technology. Still, there are many challenges associated with tolerant varieties which need to be fine-tuned. Moreover, three thousand reports of enhanced drought and salinity tolerance in wheat, sorghum, canola and rice are present but none of them is in use by farmers. A fundamental reason for this is that salinity and drought are complex multigenic traits. So, to induce tolerance in plants every gene needs to be fine-tuned precisely. However, their evaluation in the field is a long way, and distribution at the commercial level is also a hurdle in their production.

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Citrus Biotic and Abiotic Stress Management

Chapter 4

Integrated Management Approach to Citrus Fungal Diseases by Optimizing Cocoa-Based Agroforests Structural Characteristics

Ndo Eunice Golda Danièle and Akoutou Mvondo Etienne

Abstract

The health and productivity of citrus are generally jeopardized by a host of diseases, for which the environmental conditions of the cropping system are critical drivers. Several studies conducted on various diseases of perennial crops have shown the involvement of the structural futures of the cocoa-based agroforestry system (CBAFS) in the spread of pathogens and the epidemics development. This chapter highlights the effect of the CBAFS's structural characteristics on the intensity of three citrus diseases in the humid forest zones of Cameroon. The involvement of CBAFS structural characteristics in diseases regulation is demonstrated. In particular, the spatial structure of citrus in agroforests shows an effect on the spread of diseases. Moreover, distribution of citrus in the CBAFS, with minimum spacing of 12 m between citrus trees, limits the damage caused by Pseudocercospora leaf and fruit spot disease (PLFSD) and citrus diseases caused by Phytophthora (CDP). Dense shading helps to minimize the intensity of diseases such as CDP and PLFSD and Citrus scab disease. This work may make it possible to contribute to the development of an integrated management tool for citrus diseases in an associated crop context.

Keywords: Integrated disease management, cultural practices, citrus, fungal diseases, shade trees, spatial structure

1. Introduction

Conventional and intensive agriculture has enabled a considerable increase in agricultural production since the 1950s. However, the resulting heavy ecological balance sheet discredits this unsustainable model of agriculture [1]. Thus, although the preferred strategies of intensive agriculture have shown undeniable benefits, their use is becoming increasingly worrying both for agriculture itself and for the environment and human health. This is partly due to the excessive use of pesticides and other chemicals [2–6]. The improvement of intensive production systems towards new models of sustainable agriculture, favoring the development of effective means of combating diseases that are sustainable and environmentally friendly, is becoming a matter of urgency. Tropical agroforestry systems, thanks to their high biodiversity and structural diversity, represent a privileged way out of the agroecological transition. Several studies have demonstrated the contribution of the structural characteristics of these systems in the integrated management of pests and diseases of perennial crops, particularly citrus [7–9].

Citrus represent a fruit crop of prime importance in socio-ecological terms in Cameroon [10–12]. Their significance lies in the fact of their high-quality nutritional value and their contribution to the diversification of producers' incomes in rural areas. Citrus are also important in the local pharmacopeia [13–15]. They are also known for their role in restoring ecological balances after deforestation [16]. However, despite the favorable agroecological conditions throughout the country, the number of production basins identified and even the density of trees in farms, production remains poor [12, 17, 18]. A diversity of diseases affecting citrus in the country humid zones is the main constraint to their production [8, 11, 19, 20].

Pseudocercospora leaf and fruit spot disease (PLFSD) caused by *Pseudocercospora angolensis*, citrus scab disease caused by *Elsinoe* spp.; and citrus diseases caused by Phytophthora (CDP) caused by *Phytophthora* spp., are the main soil diseases on citrus in Cameroon (**Figure 1**) [7, 21–25]. Damages caused by these diseases result in heavy crop losses [26]. Sorting deviations of up to 100% of the production can be recorded in the case of PLFSD, if no treatment measures are taken [27]. Concerning citrus scab, severe attacks on young *C. volcameriana* plants, for example in the nursery, result in their death [20, 28]. CDP significantly limits citrus production in the plots where it is present [29–34]. In addition to reduced yields from the beginning of infection, the economic viability of orchards is reduced following tree death [7, 32, 35]. This strong and constantly changing diseases pressure, which not only causes enormous economic losses, but also leads (in the case of PLFSD) to quarantine and a ban on the export of citrus products to other production areas [25].

A variety of strategies are used to control citrus diseases. These include the practice of sanitation measures, the use of resistant cultivars and varieties, grafting, organic or mineral soil amendments, the use of plant extracts, biological control in a systemic approach and, above all, chemical control through fungicides [32, 36–40]. However, the high costs of these methods, development of resistance to chemical inputs, emergence of new diseases and growing concerns about environmental and soil health make these methods inadequate [41–45]. In addition, most of these techniques are unsuitable for the socio-cultural and even technological context of small-scale producers in inter-tropical regions. The development of targeted control protocols, taking into account the local socio-ecological context and existing production systems is therefore imperative. This chapter highlights the effect of the



Figure 1.

Symptoms of citrus diseases in Cameroon. C. paradisi fruit torn following a severe attack by Pseudocercospora angolensis (a), C. volcameriana fruit covered with scab spots due to Elsinoe spp. (b), lesions resulting from crown attacks due to Phytophthora spp. on C. sinensis tree(c) and dieback due to various diseases of an C. sinensis (d).

structural characteristics of the cocoa-based agroforestry system on the intensity of the three main citrus diseases in the humid forest zones of Cameroon.

2. Complex cocoa-based agroforest systems and structural characteristics

In Cameroon, citrus are mainly grown in cocoa-based agroforestry systems (CBAFS) [7, 46]. These are complex, highly biodiverse, natural forest-like cropping systems (**Figure 2**) [30, 47, 48]. In this system, several interactions of different nature and intensity can take place depending on the species present, their sizes and their positions [9, 49, 50]. One of these interactions is the action of diseases. The structural characteristics of CBAFS can contribute to control of these [20, 30, 51, 52]. Studies in these cropping systems and on various pathosystems have shown that spatial structure of species is important in reducing diseases development [49, 52]. Indeed, spatial structure has a twofold effect on the pathogen: firstly, the high plant biodiversity into CBAFS makes it possible to dilute the pathogens resource and thus reduce their presence and damage [53–55]. Secondly, multi-species agroecosystems are recognized for the high diversity of vertical and horizontal structure affects diseases mainly through the microclimatic weathering mechanism [51, 52].

Previous works have supported interactions between individuals of a host population of pathogen and associated plants within intercropping systems [7, 8]. This type of interaction is likely to influence the presence of diseases. The action of shade trees on the understory microclimate decreases with decreasing distance between trees and pathogen transmission decreases as the distance between host individuals increases [51, 52, 57–59].



Figure 2.

Illustration of a cocoa-based agroforestry system planted with citrus trees (a) and the horizontal structure of its plant population (b). 1 = cocoa trees, 2 = various forest tree species, 3 = various other fruit tree species, 4 = citrus and 5 = palm trees.

3. Effect of spatial structure of citrus into cocoa based agroforest systems on citrus diseases

The spatial structure of a plant community is the vertical and horizontal arrangement of constituent elements [51, 60, 61]. It reflects therefore the local environment around each individual [55, 62]. Within agroforest, non-host plants

mainly perform a physical barrier effect on diseases [27, 51, 54]. The effect of the citrus spatial structure within CBAFS on the three main diseases affecting them in Cameroon has been assessed through various studies. A network of 27 plots in the three study sites was set up in Obala, Muyuka and Bokito sites. These are located in three ecological conditions into humid forest zone of Cameroon. CBAFS with at least 12 citrus trees in the plot area were selected. Each plot area was a square of at least 2500 m² (50 X 50 m). Plots were chosen in villages among the most productive areas and also representative of the study zone in terms of system diversity and variability of citrus species produced.

The analysis of the spatial structure of the citrus sub-population was done by the Ripley method [62]. Following the method illustrated in Ngo Bieng [55], a typology of spatial structure was build based on the spatial structure of the citrus trees in the study plots. In a first step, the horizontal spatial structure was characterized on the citrus trees in each plot, using the L(r) modified Ripley function [20, 61, 63]. The L (r) function is based on the calculation of the expected number of neighbor trees (**Figure 2**), within a distance \leq to r of any point of the study pattern. This method enables to distinguish three types of tree spatial patterns: regular when L (r) is <0, aggregated when L (r) is >0, and random when L (r) =0. This function characterizes the neighborhood structure around a point. It is used for a simple, homogeneous and isotropic point process of density λ [64].

$$\mathbf{K}(\mathbf{r}) = \lambda^{-1} \mathbf{E}(\mathbf{r}) \quad L_{(r)} = \sqrt{\frac{K(r)}{\pi}} - r \tag{1}$$

Subsequently, a hierarchical cluster analysis based on the Euclidean distance between the values of the L(r) function of the citrus trees in the different plots was made. It resulted in clusters of plots with a similar spatial structure, based on their trend to regular, random or aggregated spatial structure. This analysis was done with *ads* and *ade4* package R 3.2.2 software (**Figure 3**). Symptoms of CDP, PLFSD, and citrus scab were assessed by the visual recognition method. The intensity was assessed using a scale from 1 to 4.

From this study it emerges that, the spatial structure has a significant influence on the intensity of the diseases observed. The analysis of variance and the mean comparison test reveal that plots in which citrus have an aggregated spatial structure, have a high intensity of citrus scab disease and PLFSD. On the other hand, plots in which citrus fruits have a regular spatial structure show a significantly low intensity of these same diseases (**Table 1**).



Figure 3.

Hierarchical classification of plots according to the spatial structure of citrus trees in the experimental plots.

Maladies fongiques	Aggregated spatial structure	Random Spatial structure	Regular Spatial structure	Anova/Tukey test	Df	F value	Pr(>F)
Pseudocercospora leaf and fruit spot disease	1.55 ± 0.36^{b}	1.43 ± 0.28^{a}	1.41 ± 0.26^{3}		2	10.08	5.14e ^{-7***}
Citrus scab disease	1.22 ± 0.56^{b}	1.11 ± 0.38^{ab}	1.08 ± 0.34^{a}		2	5.102	0.00638**
Phytophthora foot rot disease	1.59 ± 0.93^{a}	$1.97 \pm 0.94^{\rm b}$	1.78 ± 0.91^{ab}		2	6.297	0.002**
In the same column, values with same letter ι	are not significantly different	(Tukey HSD test P < 0.05). ***	indicates highly significant.				

Table 1. Effect of the spatial structure of citrus trees in cocoa based agroforests systems on diseases.

It is thus demonstrated that the aggregate spatial structure of citrus in CBAFS has a negative effect on diseases observed. These results are similar to those obtained by Ndo *et al.* [7, 8, 30] in the particular case of PLFSD. In addition to that, the involvement of spatial structure in the spread of various diseases has been shown [8, 53, 54]. Indeed, the aggregation of host populations favors the dispersion of diseases, while regularity would reduce it [65]. In addition, it is recognized that transmission of the pathogen decreases as the distance between host individuals increases [52, 58]. On the other hand, it has been shown that aggregation of host populations can reduce the incidence of pathogens [66]. Because the transmission of the pathogen between aggregates decreases with increasing distance between aggregates. These aspects would therefore explain the low intensity of CDP observed in plots where citrus fruits have an aggregated spatial structure.

4. Effect of shade intensity management on CDP and PLFSD

Depending on the situation of citrus in the CBAFS and in relation to the upper stratum, three levels of shading (dense, moderate and no shading) were defined. Shade trees play various roles in tropical agroforests. They can improve adverse weather conditions by modulating temperature variations [51, 52, 57, 59]. Shading has been recognized as one of the factors that can influence PLFSD dissemination [54, 67, 68]. Given that shading favors climatic conditions for the development of certain citrus pathogenic fungi such as *P. angolensis* or many *Phytophthora* spp. (high relative humidity (>60%) and cool temperature conditions (<25°C)) [18, 28, 69]. It is assumed that within a plot, trees under shade would have a higher incidence of the disease than those in full sunlight. On the other hand, given the role of shade trees in improving climatic and nutritional conditions, the growth of trees under these conditions can be improved, as well as their vigor and response to disease. In addition, shade trees can act as a barrier against wind and rain (the main factors in the spread of conidia) and slow the progress of the epidemic.

An experiment carried out in a fruit trees orchard in Foumbot in the Western region of the country demonstrated the effect of shade trees on the PLFSD epidemic. The trial was set up in the Institute of Agricultural Research for Development (IRAD) experimental orchard. This orchard comprises collection plots of mango (*Manguifera indica*), avocado (*Persea americana*) and various citrus trees separated from each other by fallow plots often reserved for annual crops. This experimental plot enabled to compare tree shading situations ie dense shade (under mango trees), light shade (under avocado trees) and full sun light (fallow plot). One-year old pomelo seedlings have been placed in three levels of shade i.e.: under mango trees (dense shade), under avocado trees (light shade) and on fallow land (no shade).

The results of this experiment showed that the higher the shade index (under mango trees), the lower the PLFSD severity. When the shade intensity is lower (under avocado trees), disease severity is also lower, however the differences are not significant (**Figure 4**). These results suggest that shading should be sufficient to significantly reduce PLFSD incidence. Otherwise too little shading will not have significant effect on disease severity. But in the meantime, the plant must receive sufficient sun radiation for good growth. So, it is necessary to determine an optimum shading that allows a good compromise between plant growth and reduces PLFSD incidence. This optimum can vary according to the climatic and sanitary conditions of the plantations under consideration [18, 20, 52].



Figure 4.

Graphical representation of the percentage of diseased leaves for each pomelo plant and the shade indices of the different shade trees during the first observation date (A) and the second date (B) in the Foumbot plot.

Citrus		Shade condition	15	Anova/	Df	Fvalue	Pr (>F)
diseases	Full sunlight	Light shade	Dense shade	Tukey test			
Citrus scab disease	1.08 ± 0.34 ^a	1.11 ± 0.41 ^a	1.18 ± 0.50^{a}		2	2.257	0.106
<i>Phytophthora</i> foot rot disease	1.92 ± 0.91 ^b	1.98 ± 1.00 ^b	1.56 ± 0.86 ^a	_	2	11.18	1.8 e ^{-6***}

In the same column, values with same letter are not significantly different (Tukey HSD test P < 0.05). $\stackrel{\cdots}{\cdots}$ indicates highly significant.

Table 2.

Effect of shade on citrus diseases.

Another experiment conducted in 26 cocoa-based agroforestry systems showed the effect of shading on the spread of CDP and citrus scab disease. In this study, a total set of 476 citrus were observed under three shading conditions. Depending on the tree diversity and population, the various scenarios of shade density have been coded as follows: (1) "*dense shade*" when the citrus tree were placed under a direct and thick shading of the upper stratum; (2) "*light shade*" when the citrus tree received a mean shading of a higher stratum and finally (3) "*full sun*" when the citrus fruit did not receive a shade of a top stratum.

Results showed that, there is a variation in the citrus diseases intensity depending on whether they are located under dense shade, light shade or in full sunlight. The mean comparison test thus revealed significant differences in the intensities of CDP, according to the three citrus trees situations depending on shade. Citrus trees under dense shade are significantly less affected by CDP compared to those under light shade and those in full sunlight (**Table 2**).

Results therefore showed that, the shade had a significant effect on diseases. This shading effect was positive on the intensity of CDP. In general, in CBAFS, shade reduces diseases intensity [52, 70]. Indeed, for pathogens, spore dispersal and germination are the two main phases of their life cycle. Shading promotes spore germination while the sensitivity of the dispersion of pathogen spores to microclimate depends on how it is dispersed [7, 59].

5. Mode of dispersal of citrus infectious pathogens and CBAFS structural characteristics

The majority of fungi require moisture for infection and production of conidia [9, 71, 72]. These conidia may be disseminated by wind or soil water runoff for telluric fungi like *Phytophthora*. Local dispersal is primarily favored by rain-splash as well as some insects moving on trees [24]. This mode of dissemination determines the spatial distribution of each disease.

In the case of PLFSD, the analysis of its spatial distribution indicates that the disease is distributed in clusters, and that above 12 m there is no spatial dependency. In fact, the disease spreads from one tree to its closest neighbors depending on wind speed and/or rainfall intensity. Infection will depend on the presence and quality of the host. If neighbors are susceptible hosts, infection continues and the epidemiological cycle continues. Otherwise, the course of the disease that usually have this mode of spread. However, Brown and Bolker [65] pointed out that the aggregation of host populations favors the dispersal of the diseases while its regulation reduces it. That is, the further away the trees are from each other, the slower the transmission of the disease [22].

With regard to CDP, the spread of *Phytophthora* in the field is primarily ensured by the use of infected plant material [32]. However, mechanical means of dispersal of *Phytophthora* have been illustrated. Cases of transmission from an infected root to a healthy root following their respective growing zones have been reported. The inoculum can also be spread by run-off water. Splashes can promote the spread of the inoculum from the soil to the aerial parts of the plant. This mode of disease spread may be promoted by aggregation of the host species of this pathogen. This hypothesis was confirmed by Akoutou *et al.* [7, 30]. These studies showed that citrus with an aggregated spatial structure were more attacked by CDP in contrast to those with a regular spatial structure. In fact, root diversity in the rhizosphere could limit contact between roots of the same species. Host trees planted at wide spacings and having non-host trees between them are less likely to come into contact and this would help to limit the spread of the inoculum. This effect of dilution of the pathogen's resource can also be applied to the mode of transmission through diseased fruits and contact with parts of the plant close to the soil.

In addition, it was shown that environmental factors play a critical role in the development, severity, dispersal and conservation of inoculum in the epidemiology of *Phytophthora* disease. The increase in temperature favors population growth of species such as *P. parasitica* and *P. palmivora*. Hot, dry climates are favorable to *P. parasitica*. These observations corroborate the conclusions drawn on the effect of shading on CDP development. Indeed, the cooler environmental conditions in the understory created by shade trees would make the habitat unfavorable for the pathogen. Citrus planted under dense shade would therefore be less exposed to the inoculum, which is therefore more intense in plot areas of the plot where there is no shade.

6. Conclusion

This study highlighted the effect of shading trees on citrus in agroforestry plots. In such plots citrus trees are mixed with plants belonging to different tree. Spatial structure has a significant influence on the observed diseases intensity. Plots in which citrus have an aggregated spatial structure have a high intensity of studied diseases, while plots in which citrus have a regular spatial structure are significantly

less attacked by these diseases. Optimizing the structural characteristics of CBAFS could lead to the development of integrated control strategies against fungal diseases. These management strategies will be adapted to local agroecological contexts, respectful of the environment, and applicable by smallholders.

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Chapter 5

A Current Overview of Two Viroids Prevailing in Citrus Orchards: Citrus Exocortis Viroid and Hop Stunt Viroid

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Abstract

Citrus exocortis viroid (CEVd) and hop stunt viroid (HSVd) are the main viroids circulating in all citrus-growing areas worldwide, and causing two well-known diseases on citrus trees; exocortis and cachexia, respectively. These viroids are small, covalently closed single-stranded RNA, allocated to the Pospiviroidae family. CEVd is the first viroid being described on citrus trees in 1948 in California. It is considered the largest citrus viroid at 371 nucleotides. It causes bark scaling disorder on the rootstock of citrus trees grafted on trifoliate orange and its hybrids and can cause dwarfing of trees grown on these rootstocks. HSVd was first observed in 1945 in Florida. It consists of 299 nucleotides. Stunting, chlorosis, bark gumming, stem pitting, decline, and depressions in the wood are the main symptoms of HSVd in mandarin and its hybrids. The introduction and propagation of infected budwoods are the main causes of viroids spread in citrus orchards. These agents are mechanically sap-transmissible and spread by contaminated tools. Neither seed transmission nor vectors have been reported for both viroids. Root transmission, though possible, would be overshadowed by mechanical transmission. Rapid and sensitive molecular-based detection methods specific to both viroids are available. Both diseases are controlled by using viroids-free budwoods for new plantations, launching budwood certification programs, and establishing a quarantine system for new citrus varieties introduction. The most important achievements in CEVd and HSVd researches are outlined in this chapter. This would help to provide a clearer understanding of the diseases they cause and contribute to the development of better control strategies.

Keywords: CEVd, HSVd, citrus, *Pospiviroidae*, transmission, diagnostic, interactions, synergy, antagonism

1. Introduction

Viroids are circular, highly structured, single-stranded RNA (ssRNA) phytopathogens. Although they do not code for any peptide, these enigmatic pathogens have evolved the capacity to replicate within cellular organella, the nucleus and chloroplast for Pospiviroidae and Avsunviroidae, respectively [1-4]. Viroid replication is ensured through an RNA-based rolling-circle mechanism [1]. Intriguingly, viroids can induce severe diseases in susceptible host plants similar to those caused by numerous plant viruses [5–7]. From the seven known citrus viroids only, two, namely, CEVd and HSVd, have been reported to be associated with citrus diseases that can pose significant economic risks to global citrus production. These diseases are exocortis and cachexia, respectively [8]. Since their original description in 1948 and 1950, respectively, both diseases have been reported to be present in almost all citrus-growing areas of the world, as well as in early citrus budwood registration programs [9, 10]. Given the importance of and rapid research progress in citrus virology in recent years, this review emphasizes recent findings related to CEVd and HSVd, the most serious viroids associated with citrus. It comprises reviews and research articles covering broad research areas on the characterization of both viroids and their symptoms, the development of reliable and rapid diagnosis methods, and management strategies. A brief snapshot of the present situation of CEVd and HSVd in the Mediterranean region, with an emphasis on their spread in citrus-growing areas of Morocco, is included.

2. Citrus exocortis viroid

2.1 Taxonomy

Citrus exocortis is a destructive disease infecting citrus species [9, 11]. The agent of this disease, citrus exocortis viroid [Pospiviroidae; Pospiviroid; CEVd], is a small, covalently closed ssRNA of about 371 nucleotides (nts) [12-14]. CEVd molecules can exist as either linear or circular [9]. As all viroids allocating to the Pospiviroid genus, CEVd lacks RNA self-cleavage activity and has a central conserved region (CCR), composed of two sets of conserved nucleotides in the upper and lower strands of its rod-like secondary structure, and a terminal conserved region (TCR) [15]. The rod-like secondary structure of CEVd takes the form of a model of five structural-functional domains. The latter is the Central (C), the Pathogenic (P), the Variable (V), the Terminal Left (TL), and the Terminal Right (TR) domains [16]. Based on their biological properties, CEVd sequences have been classified into two groups by using tomato as an experimental host: severe "Class A" and mild "Class B". Both classes of sequences differ by a minimum of 26 nts. These mutations affect two genomic regions, designated P_L and P_R , located respectively within the P and the V domains [17–19]. It is important to emphasize that CEVd strains of both classes cause distinct symptoms in gynura (Gynura aurantiaca (Blume) DC.) [20]. However, they induce only subtle differences in trifoliate orange (*Poncirus trifoliate* L. Raf.) used as a rootstock and a similar overall performance of the infected trees [21]. Sequencing of additional CEVd isolates revealed that further strains different from those of "Class A" and "Class B" existed. Furthermore, it seems that the sequence/pathogenicity relationship was more complex than originally anticipated [22]. Infectivity assays carried out with chimeric cDNA clones suggested that $P_{\rm L}$ is the pathogenicity-modulating domain. Although it remains to be explored how this domain modulates pathogenicity (i.e. stunting and epinasty). The role of the P_{R} domain is not known. However, infectivity assays suggested that it may influence the efficiency of viroid infection or replication in the plant [18]. Further infectivity assays of CEVd chimeras and another viroid of the Pospiviroid genus, tomato apical stunt viroid [TASVd], have been done to identify the role of individual structural domains. Firstly, it has been demonstrated that symptom severity is modulated by the TL and the P domains. Secondly, it has been shown that the V and the TR domains are involved in regulating viroid replication and/or accumulation [23].

2.2 Symptoms and economical impact

Citrus exocortis could affect a various part of the tree including the rootstock (at bark and wood levels), scion, leaves, and fruits, thus causing different types of damages such as bark scaling and cracking, bumps, severe stunting, low fruit-bearing, the poor appearance of the canopy [21, 24–27], and poor tree performance [28]. CEVd-infected trees in the orchard show typical symptoms. The most characteristic one is bark scaling on trifoliate orange rootstock, yellow stem blotch on trifoliate orange and its hybrids and Rangpur lime (*Citrus limonia* Osb.), and stunting on trifoliate orange or its hybrid rootstocks [9, 11]. It is important to note that classic exocortis symptoms are not always closely associated with all CEVd isolates. For instance, only transient flaking (Washington Navel orange L) or a fine reticulum of surface cracks (Washington Navel orange 3536) on the trifoliate orange rootstock have been observed on two CEVd-infected trees in Australia [28]. Additionally, no bark scaling symptoms have been observed in CEVd-infected Washington Navel orange trees grafted on Carrizo citrange although these trees presented lesions and blisters in the roots [26]. It is important to emphasize that bark scaling symptoms could be caused by viroids other than CEVd. Indeed, it has been proved that, in certain viroid combinations, synergistic effects occur and cause exocortis scaling symptoms in the absence of CEVd [24]. Furthermore, in Australia, a large number of trees showing exocortis-like symptoms including dwarfing and/or bud union abnormalities produced only mild epinasty when grafted on the indicator plant Etrog citron. The presence of viroids other than CEVd has been highlighted [28]. Bark scaling symptoms could be also the consequence of tree exposure to abiotic stresses such as sunburn [29]. A reduction in vegetative growth has been observed on commune clementine (*Citrus clementine* Hort. ex Tanaka) trees infected by CEVd as it has been determined by the height and rootstock and scion circumferences [21]. Similar but milder symptoms have been reported in CEVd-infected Washington Navel orange trees grafted on Carrizo citrange [26].

The major susceptible citrus rootstocks, which show exocortis bark scaling symptoms, are trifoliate orange and its hybrids, Palestine sweet lime (Citrus limettioedes Tan.) or Rangpur lime [11]. Trees grown on trifoliate orange are the most severely affected, with symptoms of bark scaling and severe stunting usually developing when the trees are around 4 years old [29, 30]. Cracking and peeling of the bark below the bud union appear when bark scaling occurs on these rootstocks [30]. Symptoms of exocortis have been also reported on citrange and Swingle citrumelo rootstocks. However, unlike trifoliate orange, bark scaling symptom does not always occur on trees grown on citrange rootstocks. Trees grafted on these rootstocks exhibit symptoms somewhat late and the level of tree stunting is usually less severe than that on trifoliate orange. On another susceptible rootstock, CEVd-infected trees showed symptoms of stunting, yellowing of the canopy, and general tree decline, and occasional flaking of the rootstock bark. On these trees, fruit quality is not affected. However, tree yield is severely reduced since the viroid causes tree stunting [30]. The time required for disease expression by citron scions is believed to be directly associated with the inherent vigor of the rootstock, the environmental temperature, and cultural practices [31]. CEVd does not induce any symptoms in most sweet orange (Citrus sinensis (L.) Osbeck), mandarin (Citrus reticulate Blanco), and grapefruit (Citrus paradisi Macfad) scion cultivars. However, when CEVd-infected budwoods are grafted on one of the previous susceptible rootstocks, distinct symptoms may appear [11].

The type and severity of symptoms induced by citrus exocortis disease depend not just on the selected rootstock as described above, but also on the amount of viroid present in the scion and the infection with other citrus viroids. High temperatures can also accelerate the development of symptoms [30]. The results of a long-term field trial carried out with clementine trees grafted on the trifoliate orange rootstock revealed that CEVd-induced effects might be both reduced or increased when CEVd-infected trees were exposed to mixed viroid infections. In other words, several interactions among viroids including CEVd have been revealed through comparative assays between symptoms developed on trees infected with CEVd alone or co-infected with other viroids. The most clear-cut interaction occurs between CEVd and Citrus viroid IV [*Pospiviroidae*; *Cocadviroid*; CVd-IV]. This interaction is manifested by the attenuation of bark-scaling or bark-cracking symptoms as a result of the occurrence of antagonism between both viroids. CVd-IV limits the negative effects of CEVd on tree performance. The reduction of tree size and fruit yield occurs mainly in trees infected with combinations containing CEVd or CVd-III and, to a lesser extent, those containing Citrus bent leaf viroid [*Pospiviroidae*; *Apscaviroid*; CBLVd] [24].

Numerous field trials have been conducted on different citrus species, varieties, and rootstocks under three different agroecosystems, to evaluate the effect of CEVd on vegetative growth and yield (Table 1). The first field trial has been conducted to assess the effect of CEVd infection on commune clementine trees grafted on Pomeroy trifoliate orange. CEVd-infected trees have been periodically monitored for a period of 12 years (from 1990 to 2002) for symptom expression, growth, and fruit yield. CEVd-infected trees showed a significant reduction of growth and yield, which became increasingly apparent over time with infection. Cumulative yield varied from 291,1 to 570,3 kg in 2001 for CEVd and the control, respectively. This equated to 50% cumulative yield lost. This yield attenuation was associated mainly with the loss of large fruit production. Indeed, it has been shown that CEVd reduced fruit production significantly for calibers 2 to 5. Cumulative weights were smaller than the control for caliber 0–1 and small calibers 6 and 7–8, with some significant difference [21]. The quality of fruits from CEVd-infected orange trees (Washington Navel) grafted on Carrizo citrange rootstock has been evaluated from 2004 to 2007. The results of this experiment showed that the quality of the fruit was not affected by CEVd infection [26].

2.3 Transmission and epidemiology

All citrus viroids are distributed primarily by the introduction and propagation of infected budwoods and subsequently by mechanical transmission, and CEVd is no exception [11]. Mechanical transmission of CEVd has been already reported. It took place on secateurs, tools, knives, and hedging equipment [9, 11, 27, 29] especially from lemon (*Citrus lemon* Bum. f.) to lemon [11]. Further, it has been shown that CEVd can survive for 8 days on steel knife blades. CEVd infectivity was not affected over a wide range of time intervals between knife contamination and transfer to citron or by 2 sequential transfers by this method. CEVd spread to susceptible hosts by contaminated tools was accomplished from numerous tested citrus species of great economic importance such as lemon, sweet orange, grapefruit, tangerine, and a trifoliate hybrid [31]. Another transmission assay carried out under greenhouse conditions showed that CEVd can be mechanically transmitted from citron to healthy citron [32, 33] and gynura herbaceous plant [33] by a single slash with a knife blade [32, 33]. CEVd-retransmission from infected gynura back to citron was successful [33]. Natural grafts of citrus roots seem to be associated with CEVd propagation. That is the case for example for a budwood multiplication block of an Australian nursery where the propagation of CEVd by natural grafts of roots induced the infection of some healthy lemon mother trees on which neither hedging nor pruning operations took place before

Combination		Effect on			References
Scion	Rootstock	Height	Canopy	Cumulative yield	
CEVd					
Clementine	Orange Pomeroy trifoliate	SE* (Reduction of almost 14% for CEVd-117)	NSE	SE (Reduction of almost 50%)	[21]
Oranger Washington Navel	Carrizo citrange ^c	SE (Reduction of almost 10% and 15% for CEVd-129 and CEVd-117, respectively)	NSE (Reduction of almost 17% and 27% for CEVd- 129 and CEVd-117, respectively)	NSE (Low reduction of about 2% and 10% for CEVd- 129 and CEVd-117, respectively)	[26]
Orange Maltaise demi	Soor orange (<i>Citrus</i> <i>aurantium</i> L.) ^a	NSE	NSE	NSE (Reduction of almost 20%)	[46]
sanguine –	Alemow <i>Citrus</i> macrophylla Webster ^b	NSE	NSE	NSE	
_	Carrizo citrange ^c	NSE	NSE	NSE	
_	<i>Citrus volkameriana</i> Ten. And Pasq. ^c	NSE	SE (Reduction of almost 33%)	NSE (Reduction of almost 20%)	
_	Cleopatra mandarin (<i>Citrus reshni</i> Hort. ex Tan.) ^c	NSE	NSE (Reduction of almost 25%)	NSE (Reduction of almost 20%)	
-	Swingle citrumelo (Citru) ^c	NSE	NSE	NSE	
_	Rangpur lime (<i>Citrus limonia</i> Osb.)°	NSE	NSE (Reduction of almost 28%)	NSE (Reduction of almost 20%)	
_	Trifoliate orange ^c	SE (Reduction of almost 25%)	SE (Reduction of almost 49%)	NSE	
HSVd					
Clementine	Orange Pomeroy trifoliate ^c	Little or no real impact	NSE	SE* (Reduction of almost 34% for CVd-IIc)	[21]
Orange Washington Navel	Carrizo citrange ^c	NSE (Reduction of almost 15% for CVd-IIc)	NSE (Reduction of almost 8% for CVd-IIb)	No effect	[26]

Combination	ı	Effect on			References
Scion	Rootstock	Height	Canopy	Cumulative yield	
Orange Maltaise demi	Soor orange (<i>C. aurantium</i>) ^a	NSE	NSE	NSE (Reduction of almost 20%)	[46]
sanguine	Alemow C. <i>macrophylla</i> Webster ^b	SE (Reduction of almost 33%)	SE (Reduction of almost 77%)	SE (Reduction of almost 76%)	
	Carrizo citrange ^c	NSE	NSE	NSE	
	<i>C. volkameriana</i> Ten. And Pasq. ^c	NSE	SE (Reduction of almost 30%)	NSE (Reduction of almost 20%)	
	Cleopatra mandarin (<i>C. reshni</i> Hort. ex Tan.) ^c	NSE	NSE	NSE (Reduction of almost 20%)	
	Swingle citrumelo (Citru) ^c	NSE	NSE	SE (Reduction of almost 36%)	
	Rangpur lime (<i>C.</i> <i>limonia</i> Osb.)°	NSE	NSE	NSE (Reduction of almost 20%)	
	Trifoliate orange ^c	SE (Reduction of almost 26%)	SE (Reduction of almost 45%)	SE (Reduction of almost 66%)	

SE: Significant effect. NSE: No significant effect.

^aSusceptible to citrus tristeza virus [Closteroviridae; Closterovirus; CTV] but viroids tolerant. ^bSusceptible to CTV stem-pitting and cachexia.

^cCTV tolerant.

^{*}A function of the used viroid isolates.

Table 1.

Results summary of the known field trials carried out in three citrus-growing countries of the Mediterranean area to evaluate the effect of CEVd and HSVd on vegetative growth and yield of different citrus scion and rootstock combinations.

their removal. This may be mainly linked to the fact that citrus trees were planted close to each other (within 2 m). The role of root grafting in CEVd transmission was assessed by excavating root systems [29]. CEVd root transmission, though possible, would be overshadowed by mechanical transmission. CEVd is not known to be a vector- or seed-transmitted [9, 11]. The role of gots as possible vectors of viroids, including CEVd, has been investigated. The experiment was carried out by rubbing healthy citrus plants with goat horns previously rubbed for 24 h on infected Etrog stems. Results highlighted the detection of CEVd in the tested plants. Therefore, transmission through gots could have facilitated the long-range spread of CEVd among both cultivated and wild plants and *vice versa* and also among graft-incompatible plants [34].

3. Hop stunt viroid

3.1 Taxonomy

Cachexia is a destructive disease infecting citrus species [11]. The agent of this disease, hop stunt viroid [Pospiviroidae; Hostuviroid; HSVd], is a small covalently closed ssRNA of about 300 nts [3, 35]. HSVd is a single member of the genus Hostuviroid [36]. HSVd molecules can exist as either circular or linear [35]. HSVd isolates are divided into five groups: three major and two minor groups. The first groups, composed of "plum-type", "hop-type" and "citrus-type", are composed of isolates from a limited number of isolation hosts. As to the second group, it has been suggested that they are the results of the occurrence of recombination events between members of the main groups [37]. Like CEVd, HSVd takes the form of a model of five structural-functional domains within the rod-like secondary structure: C, P, V, TR, and TL [15]. However, HSVd has a genus-specific CCR and a terminal conserved hairpin (TCH) and lacks a TCR [10]. It is worth mentioning that two HSVd-related Group II citrus viroids that differ by a "cachexia expression motif" have been described. It includes a cachexia disease non-pathogenic variant (CVd-IIa) and two pathogenic variants (CVd-IIb and CVd-IIc) [38-40]. Electrophoretic profiles obtained with single-stranded polymorphism (SSCP) allowed deciphering the variability among and within cachexia-inducing sources of citrus isolates of HSVd. SSCP allowed discrimination between non-cachexia and cachexia sources of HSVd. Sequence analysis showed that the V domain was extremely conserved among all the cachexia variants. Indeed, 5 nts differences, affecting both the upper (3 nts) and the lower (2 nts) strands of the V domain, were identified as the most characteristic motif allowing the discrimination between cachexia and non-cachexia sequences. It has been suggested, therefore, that the 5 nts affect the organization of a short helical region and two flanking loops of the V domain, thus modifying the three-dimensional geometry of the molecule [41]. Subsequently, it has been shown that only a single change in HSVd modulates citrus cachexia symptoms [38].

3.2 Symptoms and economical impact

Cachexia could affect a various part of the tree including the trunk, bark, twigs, branches, leaves, and fruits, thus causing different types of damages such as bark and trunk gumming with a rough and rugose appearance, bark-cracking, moderate and severe tree stunting, chlorosis, decline and death of severely affected trees, brown stipple spotting on the underside of the leaves, and the appearance of small pits on the wood [9, 21, 24]. Cachexia disease mainly affects some mandarins and their hybrids such as tangelos, and *Citrus macrophylla* Wester. Most other citrus species seem to be symptomless unless grafted on susceptible rootstocks [10]. Cachexia-inducing variants were proven to cause gummy bark disease of sweet orange [42, 43] and split bark disorder of sweet lime (*Citrus limetta* Risso) [44]. HSVd variants have been reported to induce yellow corky vein disease of Kagzi lime (*Citrus aurantifolia* (Christm.) Swingle) [45] and sweet orange [44] in India and Iran, respectively. It was subsequently shown that cachexia and a similar disorder previously described in Palestine sweet lime, known as xyloporosis, are caused by the same type of HSVd variants [40].

As mentioned before, for CEVd, the type and severity of citrus cachexia symptoms depend also on the presence of other citrus viroids in the tree. The results of a long-term field trial carried out with clementine trees grafted on the trifoliate orange rootstock revealed that HSVd-induced effects might be both reduced or increased when HSVd-infected trees were exposed to mixed viroid infections. The most clear-cut interaction occurs between HSVd and CVd-IV. This interaction is manifested by a slight increase in fruit yield and reduction of scion circumferences [24].

The same field trials described before to evaluate the effect of CEVd on vegetative growth and yield (**Table 1**) were used for the same purpose for HSVd. Little or no effect in vegetative growth has been observed on commune clementine trees infected by HSVd as it has been determined by the measure of height and rootstock and scion circumferences [21]. Cumulative yield varied from 377,6 to 570,3 kg in 2001 for HSVd (CVd-IIc isolate) and the control, respectively. This equated to 34% cumulative yield lost [21]. The negative impact of HSVd infection on cumulative yield has been reported in another study carried out on Orange Maltaise demi sanguine grafted on Alemow (*C. macrophylla*). HSVd-infected trees have been periodically monitored for a period of 12 years (from 2005 to 2017) for growth and fruit yield. HSVd-infected trees showed a significant reduction of yield of about 76% compared to healthy control [46]. As for CEVd, the effect of HSVd infection on the quality of fruit from Washington Navel orange trees grafted on Carrizo citrange rootstock has been evaluated from 2004 to 2007. The results of this experiment showed that the quality of the fruit was not affected by HSVd infection. However, a reduction occurred in the diameter of the harvested fruits [26].

3.3 Transmission and epidemiology

As pointed out before, for CEVd, propagation of infected budwoods and mechanical inoculations with contaminating tools were reported as the principal causes for the omnipresence of multiple viroid species, including HSVd, among citrus orchards [34]. Mechanical transmission of HSVd has been already reported. Indeed, the results of a transmission assay carried out under greenhouse conditions revealed that all HSVd strains are mechanically transmitted from citron to healthy citron by a single slash with a knife blade [32]. As for CEVd, the potential involvement of gots in HSVd spread has been shown under controlled conditions [34]. Top working, a common practice in Mediterranean countries, seems to have largely contributed to HSVd spread in Mediterranean citrus orchards [9]. HSVd is not known to be seed-borne [47] in citrus or to have natural vectors [11, 48].

4. Signaling pathways in citrus exocortis and cachexia pathogenesis

It is usually accepted that although the mechanisms through which viroids interact with their hosts are beginning to be dissected, the key triggering events and molecular mechanisms underlying viroid pathogenesis remain unclear [49, 50], and CEVd and HSVd are no exception. As demonstrated by various types of citrus pathogens [51, 52], further investigation of the molecular basis of viroid-host interactions is crucial to better understand the pathogenesis of viroids, and thus help to develop effective strategies to combat viroid diseases [50, 53]. Important changes occur in the chloroplast, cell wall, peroxidase, and symporter activities upon infection of Etrog citron with CEVd [54]. The CEVd-infected citron system has been subsequently used for studying the feedback regulation mechanism using transcriptomic analysis. The analysis of the woody host response to CEVd revealed the activation of basic defense and RNA-silencing mechanisms following CEVd infection. In other words, a large number of genes (about 1530) encoding key proteins involved in the RNA silencing pathway, and proteins related to basic defense

responses are expressed following CEVd infection [53]. Furthermore, a recent study elucidates the role of phytohormone pathways, particularly those linked to ethylene, in disease development and ribosomal stress caused by CEVd infection by using tomato as an experimental host [55]. For HSVd, a small RNA-mediated gene silencing response has been highlighted upon the infection of lemon by HSVd. The large amounts of HSVd-small interfering RNA (siRNA) from both central and variant domains have been suggested to be involved in interference with host gene and symptom development [56].

5. Detection methods

5.1 Biological indexing and cross protection

Generally, biological assays based on indicator host plants expressing typical symptoms of infection and able to withstand higher levels of viroid replication played an essential role in both viroid detection and characterization [57]. CEVd and HSVd are viroid diseases present in citrus orchards around the world [11]. The biological diagnosis through indexing method is considered as an efficient tool to test the health status of a plant, regarding a disease by inoculation with the grafting of the budwood or any other infected tissue in indicator plants that allow viroid replication, symptoms expression [11, 58], and the enhancement of viroid concentration [59]. However, bioassays for CEVd and HSVd detection and identification may require a panel of indicator host plants [60]. Certain considerations need to be respected for the proper indexing of citrus viroids. These include the use of excellent plants, the work under warm temperatures, and the use of citron index plants grown one per container. As mentioned previously, citrus viroids are highly mechanically transmissible and tools must be disinfected to avoid their spread [9].

The citron test is a very sensitive and diagnostic index for determining the presence of CEVd [9]. However, indexing, in vivo for CEVd diagnosis is time-consuming, labor-intensive, and requires technician greenhouses [59]. It can take 90 days after inoculation or grafting onto indicator plants [61-63]. Symptoms of slight to severe epinasty leaves wrinkled and twisted to the reverse with light to dark brown cracks in petiole and branches, blisters in the petiole, corking of the midrib, and reduced growth are the main symptoms observed on Etrog citron Arizona 861-S indicator plants graft-inoculated with CEVd-infected budwoods [28, 29, 59, 63]. An in vitro indexing procedure has been developed to minimize the risks of epidemics caused by viroids including CEVd. It has been proved that the in vitro indexing of CEVd is efficient as well as the *in vivo* diagnosis, and requires between 20 and 40 days less to reach the maximum incidence after inoculation. Epinasty, growth reduction, and rugged leaves with dry tips, and reduced size are the main symptoms observed on the sprouts planted *in vitro* and grafted with CEVd-infected callus [59]. The same symptoms have been reported for sprouts grafted with CEVd infected cortex [62]. Cuban Shaddock (Citrus maxima (Burm.) Merr.) has been proved to be the best rootstock, compared to rough lemon (Citrus jambhiri Lush.) or Volkamer lemon (Citrus volkameriana Ten.), for symptom expression on Arizona 861 S-1 citron indicator plants for indexing exocortis [64]. Gynura is also considered as an excellent indicator for CEVd. This latter reacts strongly in this host plant [9].

As to cachexia, Parson's special mandarin budded on vigorous root-stock such as rough lemon or Volkamer lemon is reported as an excellent indicator for the disease [9, 65]. The biological indexing may take up to one year before symptoms are seen. The reaction of Parson's Special mandarin may differ depending on HSVd isolates. In other words, some isolates are very mild reacting, whereas others are quite severe in their reaction to the indicator plant. Indeed, a mild strain reaction consists of just a slight browning at the bud union or cut back region of the Parson's Special mandarin while a severe reaction consists of the appearance of gum in the wood that may extend via the entire plant [9]. Cuban Shaddock has been proved to be the best rootstock, compared to rough lemon or Volkamer lemon, for symptom expression on Clemeline 11–20 indicator plants for indexing cachexia. Furthermore, the application of 0,5% foliar urea sprays, alone or in combination with 20 ppm gibberellic acid showed to produce more intense expression of cachexia symptoms in the indicator Clemeline 11–20 than the unsprayed control [64].

Cross protection is a biological assay, in which the infection of a plant with a viroid strain ensures protection from infection with another strain of the same viroid. This bioassay can be used for indirect viroid biological indexing. It has been applied in the diagnosis of several viroids including CEVd and HSVd. Typically, the principle of this method is based on the infection of the plant with a mild strain of a viroid, followed by its inoculation with inoculum from a plant suspected to be infected with a severe strain of the same viroid. Positive indexing of the viroid is revealed by the non-expression of symptoms in the tested plants [60]. It has been shown in a cross-protection assay, performed with CEVd-129 as a "protecting" strain against the severe type strain of CEVd that a mild strain of CEVd could lead to apparent "protection" against challenge inoculation with the severe strain. However, it is important to highlight that variability has been shown in the induced protection effect. The latter varied from only a brief delay to almost total impairment of symptom expression. The level of protection depends on the length of the interval between the inoculations with the mild and severe strains [66].

5.2 Nucleic acid-based methods

Since viroids lack a protein capsid, serological techniques used routinely in plant viruses' detection are not applicable [67]. Nucleic acid-based methods, including polyacrylamide gel electrophoresis (PAGE), hybridization (dot- and northern-blots and micro-/macroarrays), amplification (reverse transcription-polymerase chain reaction (RT-PCR) and reverse transcription loop-mediated isothermal amplification (RT-LAMP)) and sequencing (next-generation sequencing and Sanger sequencing), offer rapid cost-effective, and reliable diagnosis of viroids [60].

PAGE is considered as the first molecular technique used for the rapid (2–3 day period) identification of viroid infected plants. This technique continues to play a crucial role in the identification of new viroids since it is the only diagnostic method that is sequence-independent. PAGE analysis under denaturing conditions showed that many Citrus species may harbor numerous viroids including CEVd and HSVd [57]. PAGE and ethidium bromide or silver staining is considered as the first molecular technique applied for CEVd detection [22, 68]. However, it seems that the sensitivity of this technique requires an adequate viroid accumulation level [22]. In other terms, the PAGE procedure was used successfully to directly detect CEVd from field-grown sweet orange and grapefruit trees. The key was reported to be the use of large (50 g) samples of succulent, expanding-flush tissue collected during the summer season. However, samples collected from field-grown trees in January and February did not give consistent detection in trees known to be CEVd-infected, presumably because lack of new growth and low temperatures do not favor CEVd replication [69]. PAGE analysis can routinely resolve as many as four different viroids in the same sample. For instance, it has been shown that this technique can resolve two HSVd variants differing in length by only four nucleotides i.e. 303 nts vs. 299 nts [57].

Since the beginning of their use in the 1980s, dot blot hybridization and hybridization of tissue imprints began to replace PAGE for routine viroid detection. This is mainly because these methods allow the processing of a large number of samples [57]. A northern hybridization protocol, which relied on the analysis of preparations from bark tissues, was proved to be more sensitive than PAGE to detect CEVd and HSVd from field-grown plants of different citrus species and cultivars [70]. A citrus viroids-multiprobe composed of full-length clones of HSVd, CEVd, and two other citrus viroids has been constructed for the simultaneous detection of viroids associated with citrus trees. All the tested viroids were effectively detected with this multiprobe when tested by both northern hybridization and dot blot methods. It is important to highlight that this multiprobe does not allow the identification of the viroid type species resulting in a positive signal [71].

Due to the small size of viroids, numerous RT-PCR approaches can be applied for both their detection and subsequent characterization. In the case of CEVd and HSVd, numerous RT-PCR and real-time RT-PCR approaches have been developed and proved to allow the detection of CEVd and HSVd in both singleplex or multiplex assays [63, 72–76]. A list of some RT-PCR and related tests developed to detect CEVd and HSVd are presented in **Table 2**. Some of these tests allow also the discrimination between mild and severe CEVd strains and the identification of HSVd isolates associated with cachexia symptoms [77]. The multiplex one-step RT-PCR assay developed by Wang et al. [75] is considered a good tool streamlining the simultaneous detection of up to five citrus viroids, including CEVd and HSVd. This enables to reduce time and labor without affecting sensitivity and specificity. Indeed, serial dilution experiments showed that the singleplex RT-PCR sensitivity was similar to that of multiplex RT-PCR for all the tested viroids [75]. This type of assay could be used in high throughput screenings of viroids associated with citrus in field surveys, germplasm banks, nurseries, as well as in other viroid disease management programs [74]. Similarly, the multiplex RT-TaqMan PCR assay developed by Papayiannis [76] enables accurate discrimination between CEVd and HSVd with a diagnostic sensitivity and specificity of 100%. It is important to emphasize that in conventional RT-PCR tests, the overall sensitivity and specificity were lower and varied between 97 and 98% for HSVd, and 94 and 95% for CEVd. Therefore, this essay presented 1000-fold more analytical sensitivity [76]. The specificity of the tests described previously was confirmed by including healthy controls and/or plant tissue infected with other citrus graft transmissible virus and bacteria pathogens and non-targeted citrus viroids. Both singleplex and multiplex assays did not cross-react with any non-inoculated negative controls or other citrus pathogens [63, 74, 76]. To date, PCR-based approaches have been proven efficacy on viroid direct detection. However, false positives and negatives due to amplicon contamination and failure to generate cDNA of suitable size during reverse transcription, respectively, are not uncommon and therefore preclude the application of RT-PCR for large scale indexing [70].

Next-generation sequencing (NGS) technologies are currently becoming routinely applied in different fields of virus and viroids studies. These advanced technologies have therefore contributed to a revolution in both the detection and discovery of plant viruses and viroids [78–81]. NGS has also provided an alternative method to identify viroids in the citrus cultivars. In other words, transcriptome sequencing has shown efficacy in citrus viroid diagnostics. Indeed, this method enabled the simultaneous identification of numerous viroids from various citrus samples, including CEVd and HSVd [82]. A deep sequencing approach, combined with bioinformatics analysis, is already being implemented for HSVd detection in *C. lemon* in China. This finding suggests that HSVd could infect this host and potentially be a pathogen that causes disease on *C. lemon* trees [56].

Name of RT-PCR test	Sequence	Tm (°C)	Genomic coordinates	Size of the expected product	References
Primer/Probe Name	5′-3′				
RT-PCR					
Singleplex					
CEVd-R	GGGGATCCCTGAAGGACTT	60	80-98 ^a	371 bp	[72, 85]
CEVd-F	GGAAACCTGGAGGAAGTCG		99-117 ^a		
HSVd-F 27-mer VP-20	CGCCCGGGGCAACTCTTCTCAGAATCC	60	78–102	251 bp	[37, 73]
HSVd-R 26-mer VP-19	GCCCGGGGGCTCCTTTCTCAGGTAAG		60–85		
86-97 by the second sec	CTCCAGAGCACCGCGGGCCCTC	DN	120–140	140 bp	[37]
66-97 by the second sec	CTGGGGAATTCTCGAGTTGCCGC		1–23		
Multiplex					
CEVd-F194	TTTCGCTGCTGGCTCCACA	58	194–212	196 bp	[63]
CEVd-R18	ACCTCAAGAAAGATCCCGA		371–18		
HSVd-F1	GGGGCAACTCTTCTCAGAATCC		81–102	302 bp	
HSVd-R1	GGGGCTCCTTTCTCAGGTAAGTC		58-80		
CEV-R	CCGGGGATCCCTGAAGGACTT	58	78-98 ^a	371 bp	[75]
CEV-F	GGAAACCTGGAGGAAGTCGAG		99-119 ^a		
HSVd-R	CCGGGGCTCCTTTCTCAGGTAAGT		59-82 ^b	302 bp	
HSVd-F	GGCAACTCTTCTCAGAATCCAGC		83-105 ^b		

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Name of RT-PCR test	Sequence	Tm (°C)	Genomic coordinates	Size of the expected product	References
Primer/Probe Name	5'-3'				
Real time-RT-PCR					
Singleplex					
CEVd – 161 F	GTCCAGCGGAGAAACAGGAG	60	181-200 ^c	105 bp	[74]*
CEVd -258 R	AGAGAAGCTCCGGGCGA		270-286 ^c		
CEVd – 187 P FAM	TCCTTCCTTTCGCTGCT		212-228 ^c		
HSVd-208 F	GAGACGCGACCGGTGG	60	216-231 ^d	88 bp	
HSVd-295 R	GCTCAAGAGGGATCCGCG		286-304 ^d		
HSVd-226 P TET	TCACCTCTCGGTTCGTC		234-250 ^d		
Multiplex					
CEVd-RTR_F	GTCGCCGCGGGGATCACT	60	142–159	64 bp	[63]
CEVd-RTR_R	CCAGCAGCGAAAGGAAGGA		187–205		
HSVd-RTR_F	GGAATTCTCGAGTTGCCGCA		5–24	127 bp	
HSVd-RTR_R	CCGCGGCCCTCTCT		118–131		
CEVd-RTR_P	CCAGCGGAGAAACAG		163–177	Ι	
HSVd-RTR_P	CAACTCTTCTCAGAATCC		85–102		

Name of RT-PCR test	Sequence	Tm (°C)	Genomic coordinates	Size of the expected product	References
Primer/Probe Name	5'-3'				
CEVdF	GCGTCCAGCGGAGAAACA	60	158–175 ^e	68 bp	[76]
CEVdR	CAGCGACGATCGGATGTG	I	226–208 ^e		
CEVdTAQ	{FAM}-TCGTCTTCCTTTCGCTGCTGG-{BHQ1}	I	$181-204^{e}$		
HSVdF	GCCTTCGAAACACCATCGA	l	159–177 ^f	71 bp	
HSVdR	CACCGGTCGCGTCTCATC	l	230–213 ^f		
HSVdTAQ	{HEX}-CGTCCCTTCTTTTTACCTTCTCCTGGCTC-{BHQ2}		179–208 ^f		
The same primers have been ^a GenBank Accession no. NC- ^b GenBank Accession no. NC- ^c GenBank Accession no. CEV ^d GenBank Accession no. U211 ^c GenBank Accession no. U211 ^f GenBank Accession no. GO2.	also tested in multiplex Real time-RT-PCR test. 001464. 01351. d. HQ284019. d. KJ810553. 26.				

Table 2.

Primer sequences and their annealing temperature (Tm), primer/probe location, and expected size of PCR products for each primer pair when used to amplify CEVd and HSVd by RT-PCR and related tests (this is not a full or exclusive list).

6. Control strategies

In vitro somatic embryogenesis, from both style and stigma cultures, has been proved to be a highly effective sanitation method leading to the complete elimination of the main virus and virus-like diseases associated with citrus. Furthermore, it has been shown efficacy to eliminate diseases induced by viroids, and the production of healthy citrus plants [83-85]. This method was applied to eliminate CEVd and HSVd from some Citrus species [83]. For example, somatic embryogenesis has been tested on 13 genotypes, belonging to the Algerian germplasm collection of two different Citrus species, lemon and sweet orange, infected by at least one grafttransmissible agent, including CEVd and HSVd. This method has shown efficacy to eliminate CEVd from 12/13 tested genotypes. However, HSVd was proved to be the most infectious viroid since it has been eradicated only from 5/13 tested genotypes. It is important to emphasize that no somaclonal variability has been highlighted in lemon regenerated plants. However, a genetic instability has been observed in some of the regenerated orange plants Washington navel 251 [83]. Sanitation by in vitro shoot-tip grafting has also been proved to be a very effective method for citrus graft-transmissible diseases eradication including citrus viroids (success rate of about 100%) [86–88]. For instance, it has been reported that CEVd and HSVd can be routinely eliminated from citrus by shoot-tip grafting. Since citrus viroids are extremely tolerant of heat, the use of thermotherapy as a sanitary method is not effective in eliminating viroids from citrus budwoods [9].

No naturally occurring durable resistance has been observed in most species, despite non-hosts for viroids exist. Therefore, the effective control methods for viroid diseases consist mainly of detection and eradication, and cultural controls [50].

7. Viroids situation in the mediterranean region: focus on Morocco

Exocortis and cachexia are widespread diseases in the Mediterranean region. CEVd and HSVd have been reported in most Mediterranean countries and are among the most prevalent citrus viroids in the region [9]. The development of reliable diagnostic methods facilitated extensive surveys for CEVd and HSVd in different parts of the region. Both viroids were successively identified in many countries, including Morocco [89–92], Cyprus [33], Spain [26], Egypt [43, 93], Italy [61, 94], Tunisia [46, 95], France [21], Syria [96], and Turkey [42].

In Morocco, exocortis and cachexia are among the major citrus viroid diseases [90, 91]. These diseases are prevalent in citrus orchards and can be found in all *Citrus* species and varieties [89–92, 97]. Mechanical transmission of citrus viroids, including CEVd and HSVd, via working tools seems to be behind the widespread of these phytopathogens and their detection in both old and young plantings in all surveyed citrus orchards [92]. Research, recently completed from 2008 to 2018, to monitor CEVd and HSVd prevalence, in the main citrus-growing areas of Morocco (Gharb, Haouz, Loukkos, Moulouya, Souss, and Tadla), showed that CEVd and HSVd are omnipresent in almost all citrus-growing areas of the country with relatively high prevalence. That is the case for example for the Gharb area where CEVd and HSVd were detected at a prevalence of 85% [89] and 21% [92], respectively. Concerning genetic analysis, a first sequence comparison among six Moroccan HSVd isolates collected in the six main citrus-growing areas of Morocco has been recently reported by Afechtal et al. [92]. Phylogenetic analysis showed that the six HSVd isolates are clustered into one group within the "citrus-type". Furthermore, it seems that sequence variability is neither a function of host plant nor a function of the symptoms [92].

8. Conclusions

Citrus viroids, including CEVd and HSVd, are distributed mainly by the introduction and propagation of infected budwoods, by top working, and by mechanical transmission [9, 11]. Both viroids are known for their ability to infect a large number of host plants [36]. CEVd and HSVd are destructive to certain citrus varieties and, can cause yield losses that may be as high as 34 to 76 percent depending on the combination viroid-rootstock-scion [21, 46]. The mechanisms through which CEVd and HSVd interact with their hosts and induce pathogenesis are beginning to be deciphered. In other words, the involvement of RNA-silencing and basic defense mechanisms following CEVd and HSVd infection has been highlighted [54, 56].

Once introduced and established in a country, both viroids can spread relatively rapidly because of their ability to be transmitted via mechanical means [9, 11]. Since CEVd and HSVd have a high resistance to heat, the chemical treatment appears to be the best method to disinfect CEVd- and HSVd-contaminated tools. For instance, a 0,25 to 0,5 and a 1 percent solution of sodium hypochlorite appears to be the best option to eliminate CEVd and HSVd, respectively, from contaminated hedging and budwood cutting tools [11, 98]. Like all citrus viroids, CEVd and HSVd seem to be successively eliminated from propagative material by shoot-tip grafting or by the deployment of nucellar budlines. Being extremely tolerant of heat, CEVd and HSVd have not been successfully eliminated from budwood by applying thermotherapy [9]. Certification programs must include measures to control viroid spread in nurseries [32]. The majority of rootstocks that are tolerant to the citrus tristeza virus [*Closteroviridae*; *Closterovirus*; CTV] are susceptible to citrus viroids. Therefore, in the absence of a certification program, exocortis disease usually follows upon the replantation of these rootstocks [9]. Since no useful sources of natural resistance to viroid disease are known, diagnostic tests continue to play a key role in efforts to control viroid diseases [67]. Nowadays, several nucleic acid-based methods for detecting CEVd and HSVd exist, including PAGE, hybridization, amplification, and sequencing [60]. Although biological assay has several disadvantages, it will always play a pivotal role in viroid research. Indeed, Cuban Shaddock has been proved to be the best rootstock for symptom expression on Arizona 861 S-1 citron and Clemeline 11–20 indicator plants for indexing exocortis and cachexia, respectively [64]. Besides, gynura seems to be an excellent indicator for CEVd [9]. A combination of both molecular and biological assays should lead to the most effective means for viroid identification and characterization [60].

Complicated interactions, including antagonism and synergy, occur between viroids coinfecting the same citrus host. These interactions may lead to different symptoms, canopy volumes, fruit yields, and commercial performance. Although no obvious physiological changes in citrus hosts have been described in mixed infections of CEVd and HSVd and both viroids do not induce severe symptoms in citrus [24, 99], their interaction was intriguing because they are commonly found simultaneously infecting different citrus cultivars and they have identical biological properties within the same host. The relationship between the two viroids has been investigated over 3 years (from 2011 to 2013). Results showed a positive correlation between CEVd and HSVd in specific tissues of two citrus cultivars (blood orange and Murcott mandarin). This result has been supported by three findings: titer enhancement, localization similarity, and lack of symptom aggravation under mixed-infection conditions. Compared to their concentrations under single-infection conditions, a significant increase in the CEVd and HSVd population has been observed under mixed-infection during 6 and only 1 season of the 12 monitored seasons, respectively. This result is somewhat surprising because no competition phenomenon for host resources occurs between the two viroids although they have
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the same biological functions and share identical cellular and subcellular spaces [27]. This issue merits consideration in future research.

Regarding the current situation of CEVd and HSVd in Morocco, this chapter provides a general overview of their spread in the Moroccan citrus-growing areas. Preventing the introduction and the establishment of exocortis and cachexia diseases in the Moroccan citrus orchards can be set up through the use of viroid-free (certified) planting material, disinfection of pruning tools, regular monitoring of citrus orchards to ensure early detection of both diseases, and by avoiding top working practice. This review pointers to new research avenues in exocortis and cachexia diseases in Morocco or elsewhere. These research fields could include for instance the characterization of CEVd and HSVd isolates, searching for secondary hosts, and developing sustainable control strategies. Investigating the prevalence of CEVd and HSVd infection in numerous natural host plants, and the characterization of the viroid sequence variants is valuable especially that a cross-transmission phenomenon between different hosts seems to be possible for HSVd [100].

Studying functional genomics through transcriptomic analysis and/or proteomic approaches in citrus-CEVd/HSVd interaction would be an interesting approach to shed more light on the full mechanisms underlying the complex and varied events associated with such interactome, and thus contribute to the development of novel diagnostic methods and plant protection strategies. This further advanced research will expand our understanding of CEVd and HSVd epidemiology and the mechanisms behind their spread across the world in general and Morocco in particular, and could potentially help in devising innovative management strategies of both viroids.

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Conflict of interest

The authors declare no conflict of interest.

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Chapter 6

Indexing Virus and Virus-Like Diseases of Citrus

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Abstract

Citrus is a highly nutritive and prized fruit crop around the world. It contributes a substantial share in local consumption and exports of a nation to earn a handsome foreign exchange. The production of citrus is under the threat of citrus decline. Different factors are responsible for the citrus decline but virus and virus-like diseases have the major role in this decline. Virus and virus-like diseases alone or in association with other biotic and abiotic factors exist in the citrus orchards. Therefore, indexing of diseases caused by virus and virus-like pathogens is the key factor to manage these citrus diseases. Proper facilities and skilled personnel are the pre-requisite for the diseases indexing procedures. Biological, serological and molecular indexing is sensitive, reliable and durable strategy for managing different citrus virus and virus-like diseases under different conditions. Moreover, indexing of viruses and virus-like pathogens are very important for the production of disease free citrus nurseries. This chapter gives a brief review for the commonly used biological, serological and molecular assays for the detection of citrus virus and virus-like pathogens.

Keywords: citrus, detection, diseases indexing, viruses and virus-like pathogens, graft-transmissible diseases, viroids, RT-PCR, ELISA

1. Introduction

Citrus belongs to family Rutaceae and holds an important position among fruits all around the globe. It is the most cultivated fruit in the world after grapes. Citrus is believed to be originated from southeastern Asian region [1]. Northern hemisphere accounts for about 70% of the total citrus production and approximately 80 citrus species are native to India and other tropical and sub-tropical areas of Asia [2]. Citrus being a perennial fruit tree is usually produced through vegetative propagation of scion on rootstock. Combination and compatibility of scion and rootstock can result in high yielding citrus plants. The United States, China, Brazil and the Mediterranean countries contribute two third of global citrus production and are regarded as major citrus producing countries [3]. Citrus products and by-products provide the basis for local agricultural industries, which generate employment and raise income, and in many cases, this industry constitutes an important source of foreign revenue for developed and developing countries such as Pakistan.

A number of factors and certain conditions are collectively responsible for fluctuations in citrus production. Selection of rootstock, agronomic practices and management in citrus nurseries and orchards, propagation methods and biotic and abiotic factors contribute their share to some extent in reduced citrus production. Like other commercial crops, number of diseases, insect pests and genetic problems affect the citrus production. Diseases are one of the major limiting factors for the low citrus production and gives a serious threat to citrus industry. These diseases are caused by fungi, prokaryotes, nematodes, viroids, viruses and virus-like pathogens. Among these, viruses and virus-like pathogens play a major role in citrus decline. These pathogens incur varying degree of damages to citrus plants and make their life span shorter, causing low yield and deterioration of quality and ultimately loss of economy which leads towards the citrus decline [4].

Citrus decline is the matrix of all above mentioned factors and conditions. The common diseases, playing an important role in citrus decline are citrus gummosis caused by Phytophthora sp. and Fusarium sp., citrus canker caused by Xanthomonas sp., Huánglóngbìng caused by Candidatus laberibactor sp., citrus stubborn caused by Spiroplasma citri and one of the most devastating citrus viruses *i.e.* citrus tristeza virus. Citrus viruses play a vital role in its decline by using the prevailing conditions and many other factors as these are bud/graft-transmissible and have systemic infections. A variety of symptoms has been observed regarding the infection of citrus viruses resulting in systemic infection. No viral diseases on citrus was under discussion or the hot issue before 1940 but during and after 60 years, thirty economically important viruses and virus-like diseases of citrus were recognized as a cause of citrus decline in different parts of the world [5-7]. Unfortunately, citrus orchards are short lived and decline within 15 years as against their potential of 50 years or more. This is mainly attributed, among other factors, to the prevalence of graft-transmissible virus and virus-like diseases, faulty nursery operations and poor orchard management. However, most of the problems originate from nurseries.

Therefore, it is the time when citrus nurseries should operate on highly technical and scientific lines and start providing disease-free and certified plants to the growers. In the first instance, nurseries should be registered and indiscriminate multiplication and sale of uncertified citrus plants must end. For this purpose, the most imperative points such as the prevalence and detection of citrus viral diseases, selection of material, production of disease-free material and streamlined screening procedures are highlighted in this bulletin. If the guidelines are properly followed and certified bud-wood becomes available for producing disease-free citrus plants, the problem of citrus decline can be minimized.

1.1 Citrus pathology

Citrus pathology is the study of citrus diseases caused by biotic (pathogens) and abiotic factors. It is now being considered as a major part in the field of plant pathology. Being a major fruit crop in the world, citrus production always remains important for the citrus industry. Physiology, morphology, biochemistry and behavior of the citrus tree towards the prevailing climatic conditions are the key areas to be kept in mind while investigating the citrus diseases. Etiology of citrus diseases and their detection methods help to manage these diseases. A plenty of information regarding the diseases of citrus and their control has been published around the world.

2. Virus and virus-like diseases of citrus

Virus, viroids and virus-like diseases, however, infecting different citrus species could not receive due attention because of the lack of laboratories with proper facilities for their proper identification. These diseases are also known as

'graft-transmissible diseases' (GTDs) and the term used for the casual agents is 'citrus graft-transmissible pathogens' (CGTPS) [8]. These are an emerging threat for citrus industry. Major viruses and virus-like pathogens include citrus tristeza virus (CTV), citrus yellow vein clearing virus (CYVCV), citrus variegation virus (CVV), concave gum, psorosis, cristacortis, ringspot, exocortis, *Cachexia-xyloprosis*, *Candidatus liberibacter asiaticus* and *Spiroplasma citri* [9, 10]. A brief description of these virus and virus-like pathogens is summarized below (**Table 1**).

Although plant pathologists have put their efforts for the identification and management of virus and virus-like diseases of citrus but there are some areas need to be investigated. A comprehensive book has been written by Roistacher in 1991 regarding the detection of virus and virus-like diseases of citrus. These diseases reduce the citrus yield and ultimately result in the loss of low foreign exchange. Diseases caused by viruses and virus-like pathogens are infectious, contagious and devastating due to their systemic nature. They are transmitted through different means in nature; through vegetative propagation, by insect vectors and horticultural tools used for the routine activities in citrus orchards and nurseries. These diseases have a considerable economic importance because of their involvement in

Sr. No.	Citrus disease	Pathogen	Transmission	Incidence %	Host	Importance
1.	Citrus greening disease (CGD) Huánglóngbìng	Bacterium- like organism	Psyllid: Diaphorina citri	20–90	Sweet orange, grapefruit, orange jessamine	Associated with citrus decline
2.	Citrus tristeza virus (CTV)	Closterovirus	Aphid species (Aphis gossypii, Toxoptera citricida)	7–18 Up to 48	Sweet orange, lime, mandarin	Economically important
3.	Gummy bark (GB)	Virus probable	Grafting, mechanical	20–30	Mandarin, sweet orange	-do-
4.	Bud union crease (BUC)	Virus probable	Grafting, mechanical	20–30	Mandarin, sweet orange	-do-
5.	Cristacortis	Virus probable	Not known	10	All citrus species	-do-
6.	Exocortis	Viroid	Mechanical	7–10 Up to 16	Sweet lime	-do-
7.	Cachexia- xyloporosis	Viroid	Mechanical	4–10	Mandarin	-do-
8.	Citrus stubborn disease (CSD)	Prokaryote	Leaf hopper (<i>Neoaliturus</i> <i>haemocops</i>)	2–7	Sweet orange, grapefruit, periwinkle	-do-
9.	Yellow vein clearing (YVC)	Virus	Grafting, vector not known	2	Lemon, sour orange	Minor importance
10.	Ring spot/ Variegation	Virus	Not known	2–3	Sweet orange	Minor importance

Note: All diseases are graft-transmissible. No adequate information on vector transmission is available except their identity; viroids problems are favored by warm conditions.

*The above summarized information is extracted from the work of [10–12].

Table 1.

Major virus and virus-like diseases of citrus in Pakistan, their transmission and hosts^{*}.

Citrus species	Virus				Viroid		l Prokaryote Virus-like symptoms				oms	PS	DE		
	CTV	IVV	RS	YUC	Ex	сх	GR	ST	BS	GB	BU	BP	Misc.		
Sweet orange (Mosambi)	+		+		+	+	+	+		+	+	+	+	+	+
 Sweet orange (Mosambi)	+		+		+	+	+	+		+	+	+	+	+	+
Mandarin	+					+	+		+	+	+	+	+	+	+
Sweet lime					+						+	+	+	+	+
Grapefruit								+		+	+	+	+	+	
Lemon	+	+	+	+											
Acid lime	+								+				+		+
Rough lemon		+	+				+							+	
Sour orange	+	+		+									+		
 Orange jessamine					+		+			+					

Note: "+" is the indication of presence of infection on the citrus varieties.

CTV = Citrus Tristeza Virus, IVV = Infection variegation, RS = Ring spot; Ex = Exocortis, CX = Cachexia xyloporsis, GR = Greening disease, ST = Stubburn Disease, BS = Bark Scaling, GB = Gummy Bark, BU = Bud Union Disease, PS = Psorosis, DE = Decline.

[10, 12].

Table 2.

Citrus species and presence of viruses and virus-like diseases.

the citrus decline [4]. Millions of citrus trees have been died due to CTV. The CGTPS usually have two types of effects either quick decline or long term losses. These diseases are very difficult to control or manage unless or until by the application of integrated management practices. The appropriate diagnosis or indexing method plays an important role for the management of CGTPS [8].

The major symptoms due to virus and virus-like pathogens are vein clearing, bark cracking, yellowing of leaves, leaf dropping, gummosis, mosaic, rugosity, bark scaling, stem pitting, dwarfing, chlorosis and mottling [10, 13]. The virus and viruslike diseases, infecting different citrus species in Pakistan, have been neglected for a long time due to lack of proper facilitations in the research laboratories and skilled personnel for their detection and characterization. A brief description is presented in **Table 2** regarding the citrus species and viruses and virus-like diseases in Pakistan. Indexing facilities are very important for the diagnosis of plant pathogens. Similarly, unlike other pathogens viruses and virus-like pathogens are very sensitive to their indexing through different techniques. Pathogen detection system always played an important role in management of virus and virus-like pathogens. Proper indexing facilities help in the characterization and differentiation of different viruses and their isolates. Management of viruses and virus-like pathogens is only possible when appropriate indexing procedures and facilities are available.

3. Insects as vectors of virus and virus-like pathogens

Insect pests have always been key role players in the direct or indirect transmission of plant pathogens in agricultural and horticultural crops [14–16]. Citrus tristeza, cachexia-xyloporosis, greening or Huánglóngbìng, infectious variegation, vein

enation, yellow vein clearing, exocortis and stubborn are the most conspicuous viral diseases of citrus all over the world including Pakistan [11, 17]. These diseases are usually graft-transmissible and phloem-restricted. Although these diseases along with other fungal, bacterial or mycoplasmic infections of citrus are usually spread through unhealthy mechanical intrusions and by the use of infected uncertified bud, scion or rootstock in plant propagation, many type of sap-feeding insect pests play important role in the transmission of these diseases such as leafhoppers, aphids, psyllids, whiteflies and thrips [17–20].

Among the vector borne viral diseases of citrus, citrus tristeza (CTV) which is caused by a *Closterovirus* is the most dominant and widely studied viral diseases of citrus. It is transmitted by different aphid species primarily by black citrus aphid (*Toxoptera citricida* Kirk.) and cotton-melon aphid (*Aphis gossypii* Glov.) [17, 21]. Another emerging viral disease of citrus is the yellow vein clearing (CYVCV) caused by a *Mandarivirus*. It was first observed in Pakistan in 1988 in the orchards of sour orange (*C. aurantium* L.) and lemon (*C. limon* L.) [22], and later on it was reported in China, India, Iran and Turkey [23–26]. This CYVCV is reported to be vectored by e transmitted by whiteflies and aphids (*Aphis craccivora* and *A. spiraecola*) [25, 27]. Although not virus borne, citrus stubborn is a destructive disease being caused by a bacterium *Spiroplasma citri*. It is usually transmitted by many species of leafhoppers, primarily by *Scaphytopius nitridus* and Circulifer tenellus in citrus-growing suburbs of California and Arizona and by *Circulifer haematoceps* in the Mediterranean zones [17].

4. Indexing strategies

Indexing is an indispensable procedure to produce and diagnose disease-free plants. Different techniques or combination of techniques have been applied in this regard and the effectiveness of each depends upon the facilities available. Generally indexing can be divided into two types.

- A. Field indexing; also known as biological indexing including the mechanical inoculation through direct contact or vegetative propagation and/or through insect transmission.
- B. Laboratory indexing; also known as quick indexing including serological, molecular and chemical assays.

Commonly used indexing methods are tissue grafting, budding, insect transmission for biological indexing and enzyme-linked immunosorbent assay (ELISA) and polymerase chain reaction (PCR) for quick indexing strategies. Although all viruses and virus-like pathogens can be detected through PCR and its derivatives, polyacrylamide gel electrophoresis (PAGE) is commonly used for the detection of viroids.

4.1 Biological indexing

Biological indexing is the inoculation or introduction of virus source (infected sample) into the indicator plants for detection and purification. It involves one of the common indexing methods such as vegetative propagation of infected scion (grafting/budding) to indicator plants, mechanical inoculation of indicator plants or transmission of virus through the insect vector (*e.g.* aphids for CTV, psyllids for greening and leaf hoppers for prokaryotic diseases). Biological indexing is usually

Sr. No.	Disease	Appropriate number of test plants	Indicator plants (for inoculation)	Favorable temperature	Symptoms on the indicator (indexed) plant
1.	Citrus tristeza	5	Mexican lime (C. aurantifolia), sweet orange on sour orange root stock, Duncan grapefruit	65–68° F Cool-warm	Vein clearing, stem pitting, leaf cupping, decline on sour orange stem pitting on grapefruit.
2.	Yellow vein clearing	5	Lemon (C. limon), sour orange (<i>C. aurantium</i>)	Cool	Yellowing and clearing of veins
3.	Ringspot/ Infections variegation	4	Citron (C. medica), cowpea (Vigna unguiculata)	Cool	Ringspot, necrotic local lesions, distorted leaves.
4.	Psorosis	4	Sweet orange (C. sinensis)	Cool	Flecking on leaves
5.	Cachexia- xyloporosis	5	Mandarin (C. reticulata) Parsons special	Warm >95°F	Gum in bark, scion and at bud union
6.	Exocortis	5	Citron (C. medica)	Warm	Tip browning, leaf epinasty
7.	Cristacortis	4	Grapefruit, sweet orange	Cool- warm	Flecking
8.	Concave gum	4	Sweet orange	Cool	Oak leaf pattern, narrowing of leaves
9.	Citrus greening disease	5	Mandarin (<i>C. reticulata</i>), Grapefruit (<i>C. paradisi</i>), Murraya sp.	Cool- warm	Leaf blotches, chlorosis
10.	Citrus stubborn disease	4	Sweet orange, grapefruit, periwinkle (Catharanthus roseus)	Warm	Stunted shoot, smelling of leaves, Zn deficiency like signs.
11.	Yellows (probably Aster)	5	Grapefruit, Periwinkle (C. roseus)	Warm	Chlorosis
12.	Bud union crease	5	Sweet orange	Warm	Brown line at bud union
13.	Gummy bark	5	Mandarin	Warm	Gum in the bark

Note: The above information is extracted from the work of [10, 12] and personal communication of Dr. S.M. Mughal. (Dr. Mughal was also in the team of Dr. Catara during the surveys of citrus growing areas of Pakistan in early 1980's).

Table 3.

Biological indexing of citrus graft-transmissible pathogens.

time consuming, require glasshouse facilities and takes about 6–12 months for results. At least 3–4 plants are required per treatment. Biological indexing of graft-transmissible pathogens, indicator plants and symptoms in the indicator plants are summarized in **Table 3**.

Detailed methodology for biological indexing has been described much in literature [28–31]. Followings are the generalized and simplified steps to be kept in mind during the biological indexing on the basis of available literature.

i. Sow the seeds of test plants (usually Mexican lime or acid lime) in the sand in germinating tray. Transplant the seedlings in pots having potting media (sand, soil and moss @ 1:1:1 ratio) after 17–28 days of germination, depending on the germinating conditions.

- ii. Inoculate the seedlings at 4–6 leaf stage.
- iii. Keep the indicator plants in insect-free chambers before and after inoculation.
- iv. **Mechanical inoculation:** grind the virus infected samples in phosphate buffer (pH 7.2). Dust the carborundum powder on to the indicator plants, but not too much, before inoculation (excessive dusting of carborundum will cause the necrosis and misled results).
- v. Pass the crude virus extract from double layer muslin cloth and then apply to the indicator plants with the help of forefinger and leave for 3–4 min. Remove the excess sap from indicator plants under tap water.
- vi. **Insect transmission:** Do rearing of vector insects in the laboratory or collect the vector insects directly from the field and keep them in laboratory for few days to be acclimatized with the rearing conditions. Provide them with the artificial or natural diet.
- vii. Observe the fasting period according to the nature of transmission (nonpersistent, persistent or semi-persistent) before allowing them to be fed on virus source to acquire the virus.
- viii. For non-persistent transmission pre-acquisition fasting time is 30–90 min. Long fasting period enhances the chances of quick acquisition of virus from the infected source.
- ix. Transfer the insects on to the young leaves with virus symptoms/virus infected plants with the help of camel brush and allow them to feed for few min.
- x. After few min, immediately transfer the viruliferous insects on to the indicator plants and keep the plants in insect-free chamber to avoid the contamination from other insects.
- xi. Maintain the insect population on indicator plants for at least 24 hr. and eradicate them after that through insecticides.
- xii. For persistent transmission and semi-persistent transmission preacquisition fast has no effect. In both cases long acquisition feeding period enhances the chance of transmission.
- xiii. Maintain the insect population for a week and eradicate them with insecticide in case of semi-persistent transmission while maintain the insect population through transferring them on new indicator plants till they are alive.
- xiv. **Vegetative propagation:** Take the infected bud-wood from the virus source and graft on to the rootstock/indicator plants and keep the indicator plants in insect-free zones.

xv. Temperature range between 65 and 95°F helps the appearance of symptoms on indicator plants for viruses and virus-like pathogens. Observation time also varies from 3 to 16 months for different viruses, virus-like pathogens and viroids.

Laboratory indexing/advanced detection methods

There are rapid methods, highly specific, routinely applicable and some of which test large number of samples. These methods are summarized in **Table 4**. ELISA is the main laboratory indexing method used for the detection of CTV, PAGE for viroids and PCR for all diseases. Mother plants (plants recovered by nucellar embroyony *in-vivo* or *in-vitro*, by thermotherapy or micro-grafted plants or by micro-budding may be indexed by any of the above methods. Although 'chromatography' is a useful in chemical indexing of certain virus and virus-like pathogens but it is less reliable than vegetative propagation indexing. Electron microscopy is also helpful for the detection of greening and stubborn diseases other than the viruses. Moreover, *S. citri* can be cultured on a specific medium. Now-a-days, commercial kits are available for the ELISA, PCR and other detection methods along with the instructions.

4.2 Serological assays

Serology involves the quick indexing of plant viruses, based on the antibody– antigen reaction. Enzyme-linked immunosorbent assay (ELISA) is one of the widely

Sr. No.	Methods	Tested for	Advantages and limitations	References
1.	Immunofluorescence, tissue staining Azure A, Light microscopic observations	Citrus tristeza virus, yellow vein virus, greening diseases	Simple, economical, limited number of samples, time consuming, non-specific	[29, 32]
2.	Gel immunodiffusion	Citrus tristeza virus	Economical, time consuming, require quality antiserum, where ELISA facilities are not available.	[6]
3.	Enzyme-linked Immunosorbant Assay (ELISA and its variants)	Citrus tristeza virus and some prokaryotes	Rapid, economical, specific, routinely applied for large number of samples, quantitative, sensitive	[33]
4.	Electron microscopy (EM)	Citrus tristeza viru and other viruses	Quick for elongated viruses (CTV, CYVV), requires proper facilities	[6]
5.	Immunosorbant electron microscopy (ISEM)- Decoraton Technique	Citrus tristeza virus	Quick, specific, require antiserum and proper facilities, limited sampling	[6]
6.	Polyacrylamide Gel Electrophoresis (PAGE)	Citrus viroids (Exocortis, Cachexia)	Excellent for viroid detection and characterization, requires purification of viroids and proper conditions and facilities	[34]
7.	Molecular hybridization (RNA/DNA Probes), Polymerase Chain Reaction (PCR)	Virus and virus like diseases, CTV.	Highly sensitive, routinely applicable, time consuming, require primers and equipment facilities	[6]

Note: Large scale screening of material is possible with any of the method(s) mentioned above. However, there are several limitations including time and availability of proper facilities and trained manpower.

Table 4.

Advanced methods for the detection of viral diseases of citrus.

used in detection of plant viruses. It is relatively cheap and can test large number of samples.

ELISA with its derivatives, direct (DAS-ELISA) and indirect (DAC-ELISA), is the main serological indexing tool used for most of the citrus viruses at large scale samples.

Followings are some general steps followed during the ELISA based detection or indexing [35].

- i. **For DAS-ELISA;** Coat the antibodies in the ELISA plate and keep at 4°C for overnight or 37°C for 4 hr. for incubation.
- ii. Wash the plate with washing buffer for 3 times with the interval of 5 min.
- iii. Add the antigen (virus sap extracted from infected samples) into the wells of ELISA plate and incubate as above.
- iv. Repeat the washing and coat the ELISA plate with enzyme-labeled antibodies and incubate as above.
- v. Repeat the washing step and add the substrate followed by incubation for 30 to 90 min for visual observation of color change and read the micro-plate through ELISA reader/spectrophotometer for quantitative data.
- vi. For DAC-ELISA: Add the antigen and incubate the plate as above.
- vii. Wash the ELISA plate as in DAS-ELISA.
- viii. Add the primary antibody and incubate.
- ix. Add the secondary antibody and incubate.
- x. Add the enzyme-labeled antibodies and incubate.
- xi. Add the substrate and then observe the color change after incubation and read the plate through ELISA reader for quantitative data.

Note: Repeat the washing step after every step before adding the substrate. Stop the reaction in both types of ELISA with the help of 1 N NaOH.

4.3 Molecular assays

Molecular detection of citrus viruses and virus-like diseases has revolutionized the subject and provided the platform to detect the early stages of infection to reduce the economic losses. The molecular hybridization techniques supplemented with nucleic acid amplification methods based on PCR, in which high-throughput sequencing approaches can be adopted to identify the strains in relation to evolutionary history or phylogenetic assemblages [36, 37]. Although, nucleic acid based methods are highly sensitive and discriminatory allowing specific strain typing, but it bears the problems in reproducibility [38, 39]. Progressive efforts have been made to decrease the troubleshoots and hurdles to improve the amplification systems by improving the sensitivity and specificity of detection by limiting the high contents of plant related enzyme inhibitors. In contest, nested and multiplex PCR provides high sensitivity and make the possible to detect several targets in single assay [40]. Moreover, highly sensitive technologies by conducting the amplification of nucleic acids in an isothermal reaction, nucleic acid sequencebased amplification (NASBA) and reverse transcription loop-mediated isothermal amplification (RT-LAMP) provides specific detection of viruses and virus-like diseases.

The addition of real-time PCR for high-throughput testing allows the automation of PCR by combing the fluorimeteric approaches to detect and quantify the targets simultaneously [41, 42]. The combination of different protocols including the serological techniques and molecular approaches will increase the accuracy and reliability of virus diagnostic. Furthermore, in future prospects, nucleic acid arrays and biosensors assisted by nanotechnology will open new corridors to revolutionize the detection of plant viruses and virus-like diseases.

Citrus tristeza virus (CTV) is the most dangerous citrus disease all over the world and is also known as quick decline disease reducing the population of citrus trees significantly [43-45]. However, the utilization of advanced diagnostic methods, such as, biological indexing, electron microscopy (EM), ELISA and PCR or reverse transcriptase PCR (RT-PCR) is providing promising detection of the virus particles and leading towards the management strategies of CTV [46]. The application of conventional PCR is sensitive and specific under optimized and controlled conditions. However, sometimes, it is not possible to judge the amount of pathogens in the samples. Therefore, researchers have to employ other subsequent techniques for complete detection and quantification. Meanwhile, with real-time PCR approach, users can monitor the reaction and also the quantification of the specific pathogen in the sample. While setting up the real-time reaction for virus detection, it is the basic requirement to adapt the specific conditions of the detection system and instrument, and the characteristics of the reaction reagents and cycling procedures in which the most important are primer design, reaction components and conditions. The real-time PCR works well with small amplicons (5–200 bp), while standard PCR allows amplification of several hundred bases without sensitivity and specificity. Moreover, concentrations of MgCl₂, primers, and dNTPs are usually higher than conventional PCR [47].

The new developing chemistries are setting up the protocols with different characteristics depending upon the target and assay requirements. In addition to the most widely working chemistries (SYBRGreen, TaqMan, Scorpion, Molecular Beacons), there are more novel chemicals or technologies such as Amplifluor; Locked Nucleic Acid (LNA) Probes, Sigma Proligo; Cycling Probe Technology (CPT), Takara; Light Upon eXtension (Lux) Fluorogenic Primers, Invitrogen Corporation; Plexor Technology, Promega [48, 49]. Real-time technology is being used also in multiplex formatting for the specific detection and strain identification for several viruses [50–55]. Furthermore, real-time reaction in multiplex system is difficult to optimize due to different ratio between the targets and the reaction. The replacement of conventional PCR with real-time PCR is providing new horizons towards the multiple detection system of plant viruses especially of the citrus viruses and virus-like diseases.

5. Detection of citrus viroids

After the discovery of viroid group of pathogens as an infectious agent to the plants, new aspects in virology were come in front of researchers to be addressed. Viroids are the smallest pathogens which consist of 246 to 401 nucleotides. They are

low molecular weight, circular and single stranded RNAs. Viroids exist as free RNA because they lack protein coat [56]. Since viroids do not code for protein and enzyme, they rely on host enzyme for protein synthesis system and replication. To date, 38 viroids have been identified and they are classified into 2 families *i.e. Pospiviroidae* and *Avsunviroidae* [57].

The major economic important viroids in different plants are coconut viroids (CCCVd), citrus viroids (Exocortis and cachexia and variants), Hop stunt viroid and Potato spindle tuber viroids [57]. The origin of viroids is still questionable as they do not have natural host relationship [58, 59].

Citrus production is also affected by viroids. These are the emerging threat to citrus industry. To date, seven citrus viroids have been detected so far in citrus viz. Citrus Exocortis viroid (CEVd), Citrus Bent Leaf viroid (CBLVd), Hop Stunt viroidcitrus (HPSVd-cit), Citrus Dwarfing viroid (CDVd), Citrus Bark Cracking viroid (CBCVd), Citrus viroid V (CVd V) and Citrus viroid VI (CVd VI-OS). These have been distributed in different geographical areas as shown in **Table 5** [70]. Diseases caused by citrus viroids are citrus exocortis disease (CED), citrus cachexia disease (CCD), citrus leaf bending disease (CLBD), citrus bark cracking disease (CBCD) and citrus dwarfing disease (CDD). Among these, citrus exocortis and citrus cachexia-xyloporosis are the most devastating and widely distributed [57]. These diseases cause a reduction in yield, size of fruit and quality of production [8]. These are transmitted directly and through propagation [71]. Stunting, dwarfing, bark cracking, yellowing of leaves, backward leaf bent, pin holing, yield loss and ultimately tree decline are the common symptoms of citrus viroid diseases [63, 71, 72]. Citrus viroids alone or with other viruses or prokaryotes in the host contribute considerably in tree decline [73]. Exocortis and cachexia are the major viroids which are widely distributed in citrus orchards. Other citrus viroids have also been detected from citrus orchards in different parts of the world [73]. Unlike viruses, viroids do not have protein coat, therefore, these are very difficult to detect through serological methods. For this purpose, molecular techniques such as PCR, PAGE are available for the detection of citrus viroids. These are sensitive, sophisticated and rapid detection techniques. Molecular techniques not only help in the detection but also in the characterization of viroids.

Viroid	Geographical distribution	References
Citrus excocortis viroid (CEVd)	Australia, Argentina, Brazil, Japan, Taiwan, Corsica, China, India, Israel, Spain, Pakistan, South Africa USA, Uruguay, Iran	[8, 57, 60–62]
Citrus bent leaf viroid (CBLVd)	Israel, Japan, Australia, China, Uruguay, Pakistan, UAE, Iran, Spain	[8, 60, 61, 63–65]
Hopstunt viroid (CVd- II),	Israel, Brazil, Uruguay	[8, 60, 62]
Citrus dwarfing viroid (CDVd)	USA, Uruguay, Pakistan	[8, 60, 65, 66]
Citrus bark cracking viroid (CBCVd)	USA, Uruguay	[60, 67]
Citrus viroid V (CVd-V)	Spain, Iran	[61, 68]
Citrus viroid VI (CVd- VI)	Japan	[69]

Table 5.

Geographical distribution of citrus viroids.

Genus	Citrus viroid	Length (nucleotides)	Diseases
Pospiviroid	Citrus excocortis viroid (CEVd)	371	Citrus exocortis disease
Hostuviroid	Hop stunt viroid-citrus (CVd-IIa) Citrus cachexia viroid (CVd-IIb)	299–302	Citrus cachexia disease.
Cocadviroid	Citrus viroid IV	284–286	Citrus bark cracking disease
Apscaviroid	Citrus bent leaf viroid (CVd-I),	318	Citrus leaf bending disease
	Citrus dwarfing viroid (CVd-III)	294–297	Citrus dwarfing disease
	Citrus viroid V	295	
	Citrus viroid VI	330	

Note: The above information is extracted from the work of Hammond & Owens (2006) and King et al. (2011).

Table 6.

Classification of citrus viroids (king et al., 2011; Hammond & Owens, 2006).

5.1 Pathogen description and characterization

All citrus viroids are classified in different genus under *Pospiviroidae* as mentioned in **Table 6**.

5.2 Diagnostic methods for citrus viroids

Biological indexing is done through graft inoculation in indicator plants. It is very suitable to check the symptoms produced by citrus viroids and their severity. The most important host for indexing CEVd is Etrog citron (*Citrus medica*, Arizona 861) because of its great sensibility and rapid symptom expression [74]. According to Nakahara et al. [75], bioassay on Etrog citron is the most sensitive technique in detection of viroids although it takes more time compared to other methods. **Molecular tools** are now widely being used in the detection of citrus viroids. Combinations of several molecular techniques are very useful for reducing the time and to allow large numbers of samples to be examined and to identify each citrus viroid species [75, 76]. **PAGE** is also used to separate variation based on molecular weight. PAGE is not suitable for indexing large number of samples because it is not cost-effective. It is used to test the circularity of viroid RNA by two-dimensional denaturing PAGE (2D-PAGE) [77–79]. Sequential-PAGE is also commonly used and capable of detecting all citrus viroids [71].

Reverse transcription- polymerase chain reaction (RT-PCR) is the most commonly used method to detect citrus viroids. It is also a reliable method for quick screening and detection of citrus viroids [80]. It is known for its high specificity and ability to detect unknown viroids or variants [57]. **Multiplex RT-PCR** is another approach to detect simultaneously more than a viroid by using several set of primers. For instance, CEVd, CBLVd, CVd 1-LSS, CVd-II, CVd-III, CVd-IV and CVd-VI were successfully detected simultaneously via multiplex RT-PCR [69]. **Real-time RT-PCR** is also used to detect citrus viroid. It is a quantitative PCR technology basically same as RT-PCR but it measures and quantifies products generated during each cycle of PCR [81]. **Molecular hybridization** is based on the specific interaction between complementary purine and pyrimidine bases forming A-U and G-C base pairs. According to Targon et al. [82], imprint hybridization technique is fast, sensitive and economic methods to be used as a routine for citrus viroid indexing in the certification programs. However, dot-blot technique is required an appropriate amount of extracted nucleic acids [75], and it is not suitable

for detection of new or unknown viroids. Another molecular approach to detect viroids is Northern blot hybridization. CEVd, CBLVd, CVd-II, CVd-III and CVd-VI were successfully detected by Northern blot hybridization using specific probes in inoculated Etrog citron [83].

5.2.1 RT-PCR for detection of citrus viroids

5.2.1.1 Samples collection

Collect the leave samples based on virus and viroids-like symptoms in the field. Bring the leaves samples to laboratory for processing and preservation until use as follows;

- a. Collect the leave samples in sterile plastic bags and place in ice box.
- b. In the laboratory, wash the samples first in 10% bleach followed by distilled water.
- c. Dry the samples and put in the plastic bags.
- d. Label the plastic bags and store them at -80° C until further use.

5.2.1.2 Nucleic acid extraction

Extract the nucleic acids from leave samples using the TESLP buffer [84] as follows;

- 1. Grind the 2-3 g of leaves (about 10–12 leaves) using mortar and pestle with liquid nitrogen. The slurry needs to be transferred to 50 ml screw cap tubes.
- 2. Add 10 ml of TESLP buffer [0.13 M Tris–HCl (pH 8.9), 0.017 M EDTA (pH 7.0), 1 M LiCl, 0.83%SDS, 5%PVP] into the tube.
- 3. Add 16 μ l of 2-mercapthoethanol into the mixture.
- 4. Incubate the mixture for 30 min at room temperature in the rotary shaker.
- 5. Centrifuge the mixture at 11000 rpm for 15 min.
- 6. The supernatant needs to be transferred to a new 50 ml screw tube.
- 7. Add phenol:chloroform:iso-amyl (PCA, 25:24:1) @ 3:2 and mix well using vortex followed by centrifugation for 15 min, 11,000 rpm at room temperature.
- 8. Transfer the supernatant into a new 15 ml screw tube and add CA (24:1) @4:3. The mixture needs to be mixed well using vortex and repeat the step 7.
- 9. The supernatant is obtained into a new 15 ml screw tube @ 1 volume of supernatant with 0.9 volumes of 90% isopropanol.
- 10. The tube is inverted 3–4 times to mix the components. Do not vortex or centrifuge.

- 11. The mixture is incubated at -80°C for 30-40 min (or -20°C for 3-4 hr. or overnight).
- 12. The mixture is centrifuged for 15 min, 11,000 rpm at room temperature.
- 13. The isopropanol is discarded and the pellet obtained is transferred into 1.5 ml micro centrifuge tube.
- 14. The pellet is washed with 1 ml of 70% ethanol followed by washing with 1 ml absolute ethanol until the clean pallet is obtained.
- 15. The pellet is suspended in 50 μ l of sterile double distilled water.
- 16. The pellet is immediately used for RT-PCR or stored in -20°C until use.

Reverse Transcription Polymerase Chain Reaction [69]:

5.2.1.3 Synthesis of cDNA

The extracted RNA is used to run RT-PCR. Reverse Transcription process is carried out in two steps to synthesis cDNA as follows;

Step 1: (1X)

Experimental RNA = 5 μ l Reverse primer = 1 μ l Double distilled water = 2.5 μ l Total Volume = 8.5 μ l The reaction is incubated at 80°C for 12 min then immediately transferred to ice for 5 min.

Step 2: (1X)

```
AMV-RT = 1 \mul
dNTPs = 2 \mul
RNAse Inhibitor = 0.5 \mul
MgCL<sub>2</sub> = 4 \mul
RT buffer = 4 \mul
Total volume = 11.5 \mul
```

The reaction is incubated at 55°C for 30 min. After 30 min, the process is stopped when it reaches to 10°C. The cDNA obtained is stored in -80°C freezer until use (or it can be used immediately).

5.2.1.4 PCR protocol

The final volume of PCR should be 25 μ l which consists of 12.5 μ l of PCR master mix, 5 μ l of cDNA, 5.5 μ l of sterile double distilled water, 1 μ l of forward primer and 1 μ l of reverse primer.

The conditions for PCR amplification (35 cycles) are as follows: a. Denaturation:

1.94°C for 10 min

- 2.94°C for 30 seconds
- 3.60°C for 1 min

b. Annealing at 60°C for 10 seconds.

c. Extension at 72°C for 10 seconds and then 5 min.

The list of specific primers used is given in **Table 7**.

5.2.1.5 Agarose gel electrophoresis

The amplified RT-PCR product is separated using 2% agarose gel as follows [85];

- 2% agarose gel is prepared with 1x TBE buffer
- Samples are loaded in the gel and electricity is provided at 100 volts for 50 min.
- The gel is stained with Ethidium bromide for 10 min and washed with distilled water for 5 min.
- The gel is visualized under Trans UV and captured with Gel Doc XR system.

5.2.1.6 PCR product purification

Positive PCR products with expected size are purified using MinElute® Gel Extraction Kit according to the standard protocol provided with Kit.

1. The expected size of band is excised from the agarose gel with a sterile, sharp scalpel.

Viroid	Туре	Sequence	Target (Product size)
CEVd	RT-Reverse	5' -CCGGGGATCCCTGAAGGACTT-3'	371 bp
	PCR-Forward	5' -GGAAACCTGGAGGAAGTCGAG-3'	
CVd-I	RT-Reverse	5' -TCGACGACGACCAGTCAGCT-3'	233 bp
	PCR-Forward	5' -TCCCCTTCACCCGAGCGCTGC-3'	
CVd-I- LSS	RT-Reverse	5' -ACGACCGCTCAGTCTCCTCT-3'	247 bp
	PCR-Forward	5' -CTGTAACCGGACCGGTCTCCTTC-3'	
CVd-II	RT-Reverse	5' -CCGGGGCTCCTTTCTCAGGTAAGT-3'	302 bp
	PCR-Forward	5' -GGCAACTCTTCTCAGAATCCAGC-3'	
CVd-III	RT-Reverse	5'-TCACCAACTTAGCTGCCTTCGTC-3'	271 bp
	PCR-Forward	5' -CTCCGCTAGTCGGAAAGACTCCGC-3'	
CVd-IV	RT-Reverse	5' -TCTATCTCAGGTCGCGAAGGAAGAAGC-3'	209 bp
	PCR-Forward	5′ -TCTGGGGAATTTCTCTGCGGGACC-3'	
CVd-VI	RT-Reverse	5' -GTCCGCTCGACTAGCGGCAGAGAGC-3'	166 bp
	PCR-Forward	5' -CGTCGACGAAGGCATGTGAGCTT-3'	

Table 7.

List of specific primer for citrus viroids [69].

- 2. The gel slice is put in a sterile 1.5 ml micro-centrifuge tube and weighed.
- 3. QC buffer, provided with the kit, is added @ 3:1 volume of gel.
- 4. The gel slice is incubated at 50°C for 10 min until the gel slice has completely dissolved.
- 5. The mixture is vortexed every 2–3 min to facilitate the dissolution of gel slices.
- 6. Then, 1 gel volume of isopropanol is added and mixed by inverting with pipette.
- 7. The MinElute spin column is placed into 2 ml collection tube.
- 8. The sample is transferred into the MinElute column and centrifuged at 13000 rpm for 1 min.
- 9. The flow-through is discarded and put back the column into the same collection tube. 750 μl of Buffer PE is added to MinElute column and let it stand for 1–2 min.
- 10. Centrifuged at 13000 rpm for 1 min and the flow-through is discarded.
- 11. The process is repeated to remove Buffer PE completely.
- 12. The ethanol residual left at the bottom of the column is discarded and MinElute column is placed into a sterile 1.5 ml micro-centrifuge tube.
- 13.30 μ l of EB Buffer is added to the center of MinElute membrane to elute DNA. The mixture is let to stand for 1 min, and then is centrifuged at 13000 rpm for 1 min.
- 14. MinElute column is discarded and the tube is stored in 20°C.

5.2.1.7 Molecular Cloning (TOPO TA cloning kit, Invitrogen)

Positive PCR samples will be cloned using the TOPO TA cloning kit according to the standard protocol provided along with the Kit as follows;

Ligation

- 4 µl purified PCR products.
- 1 µl vector (pCR2.1-TOPO).
- 1 µl Salt solution.
- Incubate in PCR machine/ heat block at 25°C for 30 min.
- Add 2 µl of ligation mixture into competent cell *E. coli* do not pipette up and down, just thaw a bit/ swirl.
- Put 30 min in ice.

Transformation

- Put 30 sec in 42°C water bath (heat shock).
- Put in ice for 5 min (immediately after heat shock).
- Add 250 µl SOC medium to mixture-seal competent cell tube with parafilm.
- Put at 200 rpm in 37°C incubator shaker for 1 h 30 min.
- Warm the petri dish in incubator for 20-30 min.
- Spread 40 µl X-gal on petri dish (LBA media).
- After spread the X-gal, put the petri dish in incubator for 20–30 min.
- Finally, spread the sample mix on petri dish and incubate overnight at 37°C.

Note: Strictly follow the incubation time and temperature in the protocol during cloning.

5.2.1.8 Two Dimensional poly acrylamide gel electrophoresis (2D PAGE)

2D PAGE is carried out to for the detection and to check the circularity of Viroid RNA. Following is the recipe and protocol for PAGE.

Gel Ingredients

- Acrylamide (A)
- Bisacrylamide (B)
- 40% AB in 50 ml distilled water @ 19:1 ratio

Non Denaturing Gel:

Ingredients	8% GEL	5% GEL
40% AB	6 ml	6.25 ml
10X TBE	3 ml	5 ml
dH ₂ O	20.25 ml	37.7 8 ml
10% APS	750 µl	937.5 μl
TEMED	40 µl	43.75 μl
Total Volume	30 ml	50 ml

- Mix the gel with magnetic bar.
- Wash the glass with KOH and dH₂O.
- KOH washing buffer includes 10 g KOH + 10 ml dH₂O + 90 ml and 99% ETOH.
- Rinse the glass with dH₂O and let it dry.

- Prepare the gel and cast into electrophoresis set.
- Let the gel to polymerase for 30 min.
- Pre-run empty gel for 20 min at 10 mA.
- Pre-run sample for 10 min at 10 mA and then run sample for 1 hr. plus at 20 mA.

5% Non-denaturing Gel

- 40% AB: 8.7 ml
- Urea: 25.2 g
- 10XTBE: 5.25 ml
- 10% APS: 750 µl
- TEMED: 45 µl
- dH2O: 17.25 ml
- Total volume: 52.5 ml

Fixer 1:	
ETOH (10%, V/V)	10 ml
Acetic Acid (5%, V/V)	5 ml
dH ₂ O	85 ml
Fixer 2:	
ETOH (10%, V/V)	10 ml
Acetic Acid (5%, V/V)	0.5 ml
dH ₂ O	89.5 ml
Silver Solution:	
Silver Nitrate	0.3 g
dH ₂ O	150 ml
Developer Solution:	
3 mM NaBH ₄	0.023 g
Formaldehyde	0.75 ml
0.375 M NaOH	3 g
dH ₂ O	200 ml

Silver Staining:

- Fix the gel in fixer 1 for 10 min at room temperature in shaker.
- Fix the gel in fixer 2 for 10 min at room temperature in shaker.

- Dip the gel in silver stain solution for 1 hr. under dark condition.
- Rinse with dH_2O twice with the interval of 1 min.
- Add developer solution in the end.
- Stop developing by adding 5% acetic acid.

5.3 Citrus tristeza virus (Ctv): a case study

5.3.1 Introduction

CTV belongs to the genus Closterovirus of the family Clostervoiridae. Virus particle is a monopartite, positive sense, comprising of ssRNA genome of approximately 20Kb in size. It is the largest known form of a plant virus and its genome is encapsulated in a flexuous rod 2000 nm long particles composed of coat protein subunits of 25KDA [86–89]. ssRNA genome comprised of 19,296 nucleotides that encode for 12 open reading frames [90]. CTV probably originated in Asia and has been spread to all citrus growing areas by infected plant material movement and now is widely distributed to all major citrus growing areas as summarized in Table 8. Over the two decades *i.e.* 1930–1950, millions of citrus trees were destroyed due to CTV infection and citrus orchards were almost wiped out in Brazil, Spain, and Argentina. This virus was the killer of three million citrus trees grafted on sour orange rootstock alone in south California [91–94]. The tristeza disease was first reported in Florida in 1959 and by 1980s became the serious threat to citrus industry [95]. By 1991, an estimation of total world loss of 100 million trees was recorded due to CTV in Argentina, Brazil, Spain, California, Venezuela and other areas [96, 97]. Several strains of CTV have been identified primarily on the basis of their biological reaction in several citrus species and indicator plant. The major groups of strains are mild that cause barely detectable clearing of leaf veins in Mexican lime; decline-inducing strains cause death of trees when propagated on sour orange rootstock. Stem pitting strains cause mild to severe pitting of stems and branches of grapefruit and orange resulting in low yield [95, 98]. Almost all the citrus varieties and hybrids have been infected with CTV [91]. Symptom expression of CTV in citrus hosts is highly variable and depends upon host species (rootstock and scion combination), virulence of CTV isolates and soil or environmental conditions. Characteristics symptoms of CTV are vein clearing, decline, stem pitting, seedling yellows, stunting and leaf corking on different citrus hosts like sweet orange, grapefruit, grafted on sour range root stock. Severity of infection and symptoms expression on cultivars vary from mild to severe isolates [99–101]. CTV is transmitted in nature by different species of aphids in a semi-persistent manner and through grafting [102, 103]. The most efficient vector involved in semipersistent manner is T. citricida Kirkaldy (brown or black citrus aphid) when compared with other aphids.

5.3.2 Indexing

Serological and biological indexing: Indexing includes biological, serological and molecular methods, which are the common procedures according to their reliability, sensitivity and duration to detect the CTV. During a survey in Spain, 22 CTV isolates were collected on the basis of geographical information, source tree and symptomology and then were characterized by biological indexing. Diversified

Regions	Countries	Status
EPPO	Israel, Spain, Turkey	Present
	France	Found but not established
	Algeria, Cyprus, Egypt, Italy, Morocco, Tunisia	Scattered infection
Asia	Brunei, China, Georgia	Present
	India	Widespread
	Indonesia. Iran, Japan	Present
	Jordan	Unconfirmed
	Korea, Malaysia, Nepal	Present
	Pakistan	Present (Scarce Information available)
	Philippines, KSA, Sri Lanka, Taiwan, Thailand, Viet Nam, Yemen	Present
Africa	Cameroon, Chad, Ethopia, Gabon, Ghana, Kenya, Mauritious, Mozambique, Nigeria, South Africa, Tanzania, Zaire, Zambia, Zimbabwe	Present
	Libya	Unconfirmed
North America	Bermuda, Mexico, USA	Present
Central America and Caribbean	Antigua, Barbuda, Bahamas, Belize, Costa Rica, El Salvador, Guatemala, Honduras, Jamaica, Netherlands Antilles, Nicaragua, Puerto Rico, St. Lucia, Trinidad, Tobago	Present
	Dominica	Unconfirmed
South America	Argentina, Bolivia, Brazil	Present (wide spread)
	Chile	Found, not established
	Colombia, Ecuador, Guyana, Peru, Paraguay, Suriname, Uruguay	Present
Oceania	American Samoa, Australia, Fiji, New Zealand	Present
Source: Anonymous, 2	2004.	

Table 8.

Geographical Distribution of Citrus tristeza closterovirus.

symptoms were produced on 9 indicator species. Mexican lime was found to be a good indicator host [104].

In Morocco, 14 diverse isolates were selected from samples during survey and then characterized on the basis of reaction pattern. Among these 14 isolates, four were severe and two were mild isolates. Isolates were also indexed against a series of monoclonal antibodies [105]. DAS-ELISA was used to detect the CTV from the samples collected during a survey in Western and Midwestern development regions of Nepal [106]. One hundred and eighty-eight samples were analyzed through biological indexing and DAS- ELISA to detect tristeza, psorosis and similar diseases like-symptoms including viroids in orange varieties in all the regions and the cachexia was detected as the most important and widespread disease [107]. Biological indexing is still considered as an important tool using for the characterization of CTV isolates. Different strains were identified through symptoms expression on differential hosts, including Mexican lime and sweet orange. Moreover, they

observed visual symptoms of different strains on Mexican lime and sweet orange through biological indexing followed by ELISA [108]. Detection of CTV in Spain was compared by indexing using monoclonal and polyclonal antibodies [109].

Molecular indexing: Different nucleic acid based indexing methods have been developed for the quick detection of CTV. The adaptability of these methods depends upon the reliability, time duration and sensitivity. Alteration in protein patterns in rootstock bark from CTV infected tree were analyzed through PAGE [110]. There was a clear modification in protein pattern but not in CTV free trees. Similarly, Northern blot technique was used to compare dsDNAs extracted from CTV infected and CTV free plants. Two out of the three CTV isolates were detected by this method [111]. CTV was also detected in the three aphid species through RT-PCR. IC-RT-PCR was used to amplify the coat protein gene [112]. Sensitivity of cDNA probe was slightly better than hybridization with ³²P-labeled probe. Similarly, hybridization with tissue print with DIG-probe could differentiate CTV isolates grown under green house or field conditions [113]. In Taiwan, RT-PCR was found to be a rapid and sensitive assay than other serological methods but one step RT-PCR, which is the combination of reverse transcriptase and polymerase chain reaction in one tube. It is more sensitive and detects the CTV when virus concentration is very low. Comparison between ELISA and RT-PCR revealed that ELISA was better than RT-PCR at detecting mild CTV strains as the virus was detected in all parishes, while RT-PCR detected CTV in only 8 parishes. It would appear that the primers used for RT-PCR are more specific for severe CTV isolates [114]. Some modifications were introduced in PCR-ELISA to increase its sensitivity and reduced the costs of detection. PCR-ELISA is the immune-detection of PCR products and effective for detection and differentiation of plant viral nucleic acids. PCR-ELISA being a laborious and expensive method was modified and simplified by using asymmetric PCR. It made PCR-ELISA more sensitive than TaqMAN[™], a fluorescence-based detection method.

Three microscopy procedures for detecting CTV were compared which provided additional alternatives for very rapid CTV indexing, including the use of EM, SSEM and light microscopy. In light microscopy, inclusions were found in young phloem tissues of all CTV-infected hosts examined. Similarly, in SSEM virus particles were found on grids prepared with antiserum and extracts from infected tissue. CTV particles could be detected in pooled samples representing one in 100. Similarly, virus particle fragments were observed infrequently in samples representing one infected plant in 1,000 samples [32].

6. Conclusion

Citrus is an important fruit crop of the world and has a great potential for local consumption, export purposes and industrial uses. Unfortunately, citrus orchards are facing the problem of low productivity due to citrus decline. This is mainly attributed, among other factors to the prevalence of graft-transmissible virus and virus-like diseases, unhygienic nursery operations and poor orchard management. However, most of the problems arise from nurseries. It is the time that the nurseries should operate on highly technical and scientific lines and should work on providing disease-free and certified plants to the citrus growers. To establish the disease free nurseries, indexing of virus and virus-like diseases are the major area that needs to be focused. Implication of traditional and modern high-throughput biological, serological and molecular indexing techniques, such as ELISA, RT-PCR, PAGE, should be put in practice for the detection and indexing of virus and virus-like diseases of citrus plants. Moreover, citrus nurseries should be registered and indiscriminate multiplication and sale of uncertified citrus plants should be prohibited.

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Chapter 7

Xanthomonas citri ssp. *citri* Pathogenicity, a Review

Juan Carlos Caicedo and Sonia Villamizar

Abstract

The infectious process of plant by bacteria is not a simple, isolated and fortuitous event. Instead, it requires a vast collection of molecular and cell singularities present in bacteria in order to reach target tissues and ensure successful cell thriving. The bacterium *Xanthomonas citri* ssp. *citri* is the etiological agent of citrus canker, this disease affects almost all types of commercial citrus crops. In this chapter we review the main structural and functional bacterial features at phenotypical and genotypical level that are responsible for the symptomatology and disease spread in a susceptible host. Biological features such as: bacterial attachment, antagonism, effector production, quorum sensing regulation and genetic plasticity are the main topics of this review.

Keywords: Biofilm, Secondary Metabolites, Antibiotic, Xanthomonadine, Quorum sensing

1. Introduction

The surface of the plants is one of the most hostile environments, prevailing factors at the phyllosphere such as: the low availability of nutrients, the high incidence of UV rays, the fluctuating periods of temperature and humidity, mechanical disruption by winds, antibacterial compounds produced by the host plant or by microorganisms member of leaf microbiome, among others, make the bacterial persistence and survival itself a pathogenicity strategy. Due the symptoms development ceases when one pathway involved in the bacterial epiphytic survival is seriously threatened [1]. In phytopathogenic bacteria whose infection route is the phyllosphere, it is important to understand how phenotypic traits upset to ensure survival and surface fitness, and how these traits interact with the phyllosphere microbiome in order to secure the onset of infection (**Figure 1**). Besides, over the time, on a large-scale, plant leaves will age and fall, thus, the phyllosphere bacteria must have to anticipate living outside of the leaf, for example in the air, soil or reach to young leaves [2].

Bacterium Xanthomonas citri ssp. citri (Xcc) is the etiological agent of bacterial citrus canker. This bacterium is equipped with a huge arsenal of cellular structures that allow its survival in the phyllosphere before it reaches the target mesophyll tissue. Xcc secretes toxins that directly affect the survival of its competitors. Once in the mesophilic tissue Xcc produces effectors that are responsible by the appearance of spongy and corky pathognomonic lesion of citrus canker. In this chapter we will review the both bacterial life style outside and inside of the host.



Figure 1.

Phytopathogenic bacteria plant infection. A. Surface leaf survivor and biofilm formation. B. Bacterial movement to natural opening on leaf. C. Phytotoxins secretion to modulate stomatal closure. D. Effector secretion that affect the cell host behavior. E. Degrading cell wall plant protein secretion.

2. Xanthomonas citri ssp. citri taxonomy

The bacterium Xantomonas citri ssp. citri is a gram negative rod shape bacteria with a single polar flagellum. Xcc belongs to the Xanthomonas genus from the gamma proteobacteria group. This genus is constituted by 28 species and more than 150 pathovars [3]. In the early 1900s, due to pathogenicity experiments, the bacterium was classified as *Pseudomonas citri* [4]. Subsequently, the bacterium was classified as *Pseudomonas citri* [4]. Subsequently, the bacterium was classified as *Xanthomonas citri* [5]. The bacterium continued in X. citri until 1978, when it was classified in X. campestris pv. citri in order to reserve citri at the specific level [6]. In 1989 Gabriel suggest the replaced of bacterium as X. citri [7]. Using DNA–DNA hybridization approach and based on renaturation rates, the bacterium was classified as X. axonopodis pv. citri by Vauterin [8]. Lately, It was suggested major changes to Xanthomond taxonomy, it which were based on multilocus sequence analysis (MLSA) and digital DNA–DNA hybridization of whole genome nucleotide, the author has been recommend the names Xanthomonas citri ssp citri for the etiological agent of citrus cancer type A [9].

3. Microbe-host interaction

3.1 Microbe -host interaction outside the susceptible host "epiphytic life style"

Bacterial citrus canker disease cycle begins with the deposit of inoculum of *XCC* at the leaf surface by rain splash. Subsequently, the bacteria move toward the natural opening of leaves, the stomata, then, the bacteria reach the apoplastic space and start the infection process inside the host "endophytic lifestyle". In this section we are going to focus on the structures, toxins, molecules and extracellular substances that favor and promote the epiphytic interaction between XCC and susceptible citrus host.

Xanthomonas citri ssp. citri Pathogenicity, a Review DOI: http://dx.doi.org/10.5772/intechopen.97776

3.1.1 Type IV pili

Several bacterial genera are endowed with filamentous appendages called pili. These filamentous organelles include the chaperon- Usher pili, type IV pili (T4P) and gram-positive pili. All types of pili are homopolymers ensembled of thousands of units of pilin protein. The outstanding function of pili is the attachment to surfaces, besides, in *Xcc* pili type IV is also responsible for the twitching motility and biofilm formation [10]. Type 4 pili is unique in its dynamism, since, it polymerizes and depolymerizes in very fast cycles, which leads to instantaneous extension and retraction cycles producing considerable mechanical force [11], as a consequence, this organelle could attract several substrates like DNA or bacteriophages in order to internalize to periplasmic space, as well as to secrete protein across the membrane [12]. Twitching motility is a bacterial displacement that able to cell to move over humid on organic and inorganic surfaces on a fashion independent of flagella [13]. In the process of biofilm development, the T4P contributes in the initial steps exactly in the reversible attachment phase and subsequently, in the formation of mushroom microcolonies. Contribution of T4P in the pathogenicity in XCC is not completely demonstrated, however, the mutation of *pilM* gene responsible to encode a membrane protein that participate in the T4P pili ensemble reduce drastically the bacterial virulence [10].

3.1.2 Type V secretion system (non fimbrial adhesins)

Xanthomonads encode type V secretion system (T5SS), it which has a function as non fimbrial adhesins [14]. Compared with the other bacterial secretion systems, the secretion system 5 is one of the simplest complexities from the structural point of view; it is smaller and has only presence at the outer membrane of gram negative bacteria [15]. This T5SS do not have a direct energy source, there is no ATP accessible in the periplasm space neither proton gradient. Consequently, the name of autotransporter has been coined for the this T5SS [16]. The T5SS is comprised of two domains: the β barrel that is located at the out membrane and a secreted passenger. There are five subtypes of T5SS from Va-Ve and recently a new subtype the Vf has been discovered [17]. The bacterium *Xcc* is endowed with three subclasses of T5SS: Va, Vb and Vc. (Figure 2). Va is a classical auto-transporter, it which transport proteases, lipases and adhesins. The type Vb is a secretion system knowing as Two-Partner Secretion System (TPS), which is composed by a translocator protein and a cognate passenger protein. Translocation from the cytoplasm to the periplasm space occurs by Sec translocase pathway once the perception of amino terminal from signaling peptide is done. The passenger protein has effector function and is termed TpsA. It is transported by TpsB, which forms a pore in the outer membrane in order to enable the TpsA translocation. TpsB also comprise two periplasmic domains. TpsB typically contains a 16-stranded beta-barrel domain that forms the outer membrane pore and two periplasmic POTRA (Polypeptide transport associated). Its function is the recognition of the cognate partner via binding to a TPS domain in TpsA.

The T5SS subclass Vc have a trimeric transporter adhesin conformation, this surface exposed adhesin assembles as homotrimeric structure at the outer membrane [18]. Proteomic and functional studies involving T5SS have revealed roles in pathogenicity to host primarily implicated in the adhesion, especially in the initial steps of pathogenicity process [19, 20].

3.1.3 Xanthomonadin pigment

Xanthomonads bacteria produce a yellow pigment membrane bound known as Xanthomonadins. Several studies have shown that Xanthomonadin has a pivotal



Figure 2.

Schematic representation of T5SS present in Xcc. β barrel domain and POTRA are characterized with blue, linker, passengers transported are represented in green and two partner secretion system domain are characterized with red.

role in a epiphytic survival and in plant-pathogen interaction [21, 22]. In the early years this yellow pigment was associated with the carotenoids. However, it was only until its full characterization was achieved that this pigment represents a unique group of aryl-polyene, water insoluble new type of pigment [23]. Genomic analysis shows that a region near to 25.4 kb contains seven transcriptional units (*pigA*, *pigB*, pigC, pigD, pigE, pigf and pigG). This gene cluster encodes necessary elements for Xanthomonadin biosynthesis [24]. Biological roles of xanthomonadin in a pathogenicity context are: (i). favor the bacterial epiphytic survival, since, Xanthomonadin avoid the photodamage produced by UV light irradiation that results in ROS production. Similar as structural related carotenoids, Xanthomonadin absorbs wavelengths between UV-C to red light. This pigment gives the bacteria additional advantages against the other phyllosphere colonizer bacteria as it is to deal with stress related factors such as UV irradiation and consequently the photo oxidative damage. Xanthomonadin also offers protection against visible light in the presence of exogenous photosensitizers. Cellular location of Xanthomadin (outer membrane) strongly suggests that this pigment stabilizes cell membrane in the epiphytic phase of this phytopathogenic bacterium. Previous studies in which Xcc deletion mutants of the *pig* genes were used and which were inoculated using the needleless syringe pressure technique did not show a significant reduction in virulence compared to the wild type phenotype inoculated using the same technique. Instead, when using the spray

infection method, that resembles the natural infection method, it which involve the epiphytic fitness stage, the *Xcc* pig mutant strains display great reduction in the virulence compare with the wild type phenotypes [25]. (ii) *Antioxidant activity*, the oxidative stressors as ROS and H_2O_2 injury the membranes, DNA and proteins, the carotenoids pigments could efficiently quench the ROS.

3.1.4 EPS xanthan and LPS

The EPS in Xanthomonas is named as xanthan, this polysaccharide surrounds the outer membrane through non-covalent ligations [26]. Pathogenicity roles in Xanthomonas genus differ greatly depending on specie, e.g. in Xanthomonas campestris, xanthan suppresses induced innate immunity by calcium chelation [27]. In addition xanthan increases the plant susceptibility to X. campestris due to avoiding the callose deposition [28]. In Xac there is controversy regarding the direct participation of xanthan in the pathogenicity process, while some authors find just a discrete participation in the epiphytic survival [29], another study shows that xanthan deletion mutants reduce the surface leave colonization ability and consequently the severity of citrus canker disease was deeply reduced [30]. Xanthan is a key component in the biofilm formation. The gene cluster *gum* is responsible for the xanthan production and exportation. This gene cluster comprise 12 successive genes with one operon-like identical direction of transcription i.e. *gumB* to *gumM*. The first two genes of cluster *gumB* and *gumC* encode components of channel than spans the outmembrane and the periplasmic space and enable the xanthan secretion [31].

The LPS is the major component of the outer leaflet of the outer membrane. The LPS in *Xcc* have a classic conformation being a tripartite glycoconjugate forming by: lipid A that carries a core oligosaccharide and polysaccharide the O- antigen. LPS that lack the O-antigen are named as lipooligosaccharide (LOS) or rough-type LPS. LPS has an essential role in bacterial growth acting as a barrier for antibacterial compounds and delivering protection against stress as well contributing to the structural proprieties of outer membrane. Lipid A is fairly conserved in most gram-negative bacteria, however, in Xanthomonas genus there is variation in the core oligosaccharide and O antigen structures, there may even be variation between the different species of Xanthomonas [32]. Nowadays is has been established that LPS has a double role in plant-microbial interaction; (i) elicitor of immunity plant response and (ii) It has a role in the promotion of virulence, because it acts as a barrier against antimicrobial activity compounds produced by root hair. *Xcc* is able to overwhelming plant defense responses induced by LPS.

3.1.5 Quorum sensing and biofilm formation

One discovery in microbiology that completely changed the conception of microbial ecology in the last two decades was the establishment of cooperative behavior in bacterial populations. This social behavior allows members of the bacterial community to adapt to new ecological niches, colonize new habitats, gain a competitive advantage against potential competitors and resist or avoid the host defense [33]. This cooperative behavior is based on a cell to cell communication system known as Quorum Sensing. Quorum sensing (QS) is a system of bacterial cell–cell communication that enables the microorganism to sense a minimum number of cells (quorum) in order to respond to external stimuli in a concerted fashion [34]. The process of QS relies upon the production, release and detection of small signaling molecules called auto-inducers. Each bacterial cell produces a basal quantity of auto-inducers, which are exported to the extracellular environment and

reflect bacterial population density. At high cell densities, the auto-inducers reach a critical concentration, at which point they are recognized by their cognate receptor, triggering a cascade of biological functions [35].

The autoinducer in *Xcc* is a short chain fatty acid molecule known as DSF (<u>D</u>iffusible <u>S</u>ignal <u>Factor</u>). Once this DSF accumulates at the extracellular space up to a critical level, it is sensed by its cognate receptor and triggers a cascade of biological function via the internal second messenger cyclic di-GMP, which is involved in virulence, resistance and biofilm formation. The encoding genes for quorum sensing components in *Xcc* form a cluster termed as *rpf* (Regulation of Pathogenicity Factors). For detailed revision of DSF quorum sensing circuit in *Xcc* [36].

Once Xcc reaches a leaf surface, it begins the initial adhesion process that was mention above. This attachment is followed by the formation of biofilm-like structures. Biofilm classical definition is an aggregated composed by several bacterial communities, which are embedded in a self-produced matrix of EPS, these bacterial cells are attached to each other or/and to a surface [37]. Biofilm is composed by polysaccharides, nucleic acids (eDNA), proteins, and have a pivotal role in attachment and protection against biotic and abiotic factors. In *Xcc* the biofilm formation in leaf and fruit surfaces is a main virulence factor in the early stage of development of citrus canker disease. In Xcc biofilm formation and dispersion is modulated by the quorum sensing autoinducer molecule DSF. How it was mention before DSF autoinducer promotes the biofilm formation because it stimulates the EPS production and pilus ensemble. On the other hand, DSF negatively regulates the biofilm formation because; it upregulates β 1–4 mannanase, ManA, leading to EPS dispersion and disassembly of biofilm [38]. Our previous study shown that quorum sensing signaling plays an essential role in the epiphytic stage survival, which is crucial at the early phase of pathogenicity development. Since, quorum quenchers bacteria belonging to genus Pseudomonas and Bacillus, it which were isolated from leaves of susceptible citrus host, which displayed the ability to disrupt the DSF pathway in *Xcc* and reduce citrus canker severity in a high susceptible citrus host [34].

3.1.6 T4SS and T6SS potentiates the Xcc antagonism with bacteria inhabiting the phylloplane and the soil amoeba

Nutrient limitation in the phyllosphere additional to environmental changes conditions, make the surface of the leaves one of the most hostile, restrictive and competitive habitats [38]. The type IV protein secretion system is used by bacteria to inject proteins and/or DNA into the prokaryotes and eukaryotes targets. Xanthomonas are endowed with genes that encode components of T4SS, the encoding genes VirB7, VirB8 and VirB9 responsible for the outer membrane pore formation. Genes that encode for VirB3, VirB4, VirB6, VirB8, Vir11, VirD4 and VirB10, responsible for the pore formation at the inner membrane. Finally, the gene that encodes for the subunits VirB2 and VirB5 that form the extracellular pilus structure. Besides, the encoding gene for VirB1 subunit predicted as a periplasmic lytic transglycosylase that plays a role in peptidoglycan alteration throughout T4SS biogenesis [39].

A recent study shows that in *Xcc* there are near to 12 proteins that interact with inner membrane associated ATPase VirD4, that is responsible for the recognition of substrates to be secreted [40]. These proteins share a C terminal domain termed XVIPCDs (Xanthomonas VirD4-interacting proteins conserved domains). These proteins are translocated into the target bacteria cell resulting in the dead of the receptor cells [41]. This bactericidal T4SS is knowing as X-T4SS and the effectors secreted by this nanomachine are termed X-Tfes (Xanthomonadales likeceeae t4SS effectors). Finally, a recent study reported that T6SS protect *Xcc* against the predatory amoeba Dyctiostelium [42].

3.2 Microbe -host interaction inside the susceptible host

Once the bacterium *Xcc* reaches the mesophilic tissue, after of epiphytic fitness and survival events mention before, must have to face the host defense response and parallel to express the pathogenicity factors;

3.2.1 T3SS the main pathogenicity determinant

The type 3 Secretion systems T3SS is the main protein secretion system widely studied in relationship to the pathogenicity. This secretion system is shared with several pathogenic bacteria ranging from animal to plants. This system is known as the "needle" and it works by delivering effector proteins directly to the target cells and modifying their behavior. Effectors from *Xcc* strains determine the host range. i.e. avirulence factors limit the specificity at the pathogen race/cultivar level by triggering immunity reactions in hosts with a related specific resistance gene. [43]. The effector delivered by the T3SS in *Xcc* belongs to the AvrBs3/PthA family. *Xcc* contains four PthA genes that encode transcription activator-like effector (TALE); of these four genes, pthA4 is responsible for the formation of citrus canker lesions. In citrus host the gene known as CsLOB is targeted by the TALE encoded by the *Xcc* gene pthA4; this gene was assessed in two susceptible host to *Xcc* infection, i.e., grape fruit and sweet orange [44]. CsLOB1-specific function still remains unclear; some previous studies suggest that CsLOB1 is involved in the regulation of development of lateral organ and metabolism of nitrogen and anthocyanin. Some plant hormones such as auxin, gibberellin, and cytokines also have proven to exert an effect on CsLOB1 gene [45]. Therefore, TALE have been shown to promote host cell transcriptional reprogramming as a virulence strategy [46].

4. Conclusions

The bacterium *Xcc* uses various adaptation and colonization strategies, it which are mainly aimed at guaranteeing its epiphytic survival, either by overcoming stress factors of biotic origin (predators, competitors, nutrient limitation) and abiotic origin (UV radiation, humidity and temperature variability). Because, this phase of epiphytic adaptation is crucial for the subsequent development of citric cancer symptoms in the susceptible host. Despite, these mechanism not having a direct effect on the health of the host, they become virulence factors, since its abolition avoid the subsequent development of the characteristic symptoms of citric cancer. Already inside the host, the bacterium uses as the main direct pathogenicity factor, the inoculation of effector proteins TALE, this effector is responsible for inducing cell hyperplasia, leading to rupture of the leaf epidermis and resulting in raised corky and spongy lesions surrounded by a water-soaked margin, the pathogno-monic lesson of bacterial citrus canker.

Conflict of interest

The authors declare no conflict of interest.

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Chapter 8

Climate Change and Citrus

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Abstract

Climate change is the change in the statistical distribution of weather patterns that lasts for an extended period. Climate change and agriculture are interrelated processes and affect in many ways. Citrus fruits are one of the largest fruit crops in the world. Yield loss at a drastic level due to abiotic stress annually in which temperature and water stress are the main environmental factors. These factors cause biochemical, anatomical, physiological, and genetic changes in plant structure and lead to defective growth, development, and reproduction, which ultimately cause a reduction in the economic yield of the crop. An increase in temperature and water stress at critical phenological stages of citrus results in reduced tree fruit set, decrease in fruit growth and size, increase in fruit acidity, low tree yield, reduced fruit peel thickness, and pre-harvest fruit drop. Stomatal conductance and net carbon dioxide assimilation in citrus leaves can be reduced by super optimal leaf temperature. Water deficit reduces the transpiration rate, stomatal conductance by stomatal closure associated with ABA content and causes an abrupt decrease in photosynthesis and CO₂ assimilation in citrus which reduce trees overall growth and production. Interventions in agronomic practices, breeding strategies, and biotechnological approaches can mitigate climate change effects on citrus. The groundwork against climate change is compulsory for better global livelihood and food security.

Keywords: Citrus fruits, environment, global warming, abiotic stress, genetic improvement, climatic adaptation

1. Introduction

Citrus and its related genera i.e., Poncirus, Eremocitrus, Fortunella, and Microcitrus belong to the family Rutaceae [1, 2]. Citrus is a prominent fruit tree of tropical and sub-tropical regions that require a suitable climate for quality production. Citrus fruit quality and quantity are inclined by multiple factors including climatic conditions [3]. Change in optimum climate elements like low temperature/freezing, heat stress/heatwaves, CO₂ assimilation, drought/water scarcity, intensive rainfall, and relative humidity, may affect directly and indirectly citrus production [4].

Citrus tree (rootstock and scion) growth, development, fruit production, and fruit quality is reduced under the biotic and abiotic stresses [5]. Citrus with tolerant rootstocks against biotic and abiotic factors improve the growth and productivity of the trees [6]. The potential citrus yield is 18–20 tones ha⁻¹, which goes up to

25 tones ha⁻¹ in the developed world; however, the citrus average yield in Pakistan is 10–12 tones ha⁻¹ and is affected by abiotic and biotic stresses [7]. The yield gap is due to biotic, abiotic, and general factors, like agronomic practices in countries of climate risk [8].

The productivity and growth of plants are affected by climate change especially drought and high temperatures collectively [9]. Reactive oxygen species accumulate superoxide and hydrogen peroxide [10] is due to water stress and high-temperature stress which reduce the biochemical, physiological, and molecular regulation. Reduction in carbohydrate accumulation affects the flowering, fruit set, and fruit yield. However, to reduce the negative plant physiological stresses, there should be good management practices in citrus orchards. Choice of better scion enhances citrus trees to produce higher yield with good fruit quality [11].

Citrus has a phenological life cycle of the whole year, starting from February to next year January. Flowering starts during February–March in subtropical regions and is generally considered a critical period for citrus production. An increase in temperature and water stress after pollination inhibits ovule fertilization [12], which in return reduces tree fruit set, increases June fruit drop, and reduces tree yield [13–15]. Fruit growth phases from button size to mature fruit are more sensitive to heat stress and deficit irrigation. Citrus under water deficit conditions faces reduced fruit growth and ripening, which is associated with a decrease in fruit size, an increase in fruit acidity [16], and low tree yield [13]. Water stress at the pre-harvest stage in oranges develops fruit peel wrinkles [17]. An increase in optimum temperature at fruit ripening causes pre-harvest fruit drop and reduced yield (**Figure 1**).

To deal with heat and water deficit stress, there is a need to improve agronomic management practices and adopt breeding and molecular approaches. Agronomic management practices encompass factors like irrigation, nutrition, pruning, pests, diseases, and other injuries which have a key role in citrus fruits quantity and quality [19]. Breeding approaches need to search out/develop rootstocks that are tolerant/resistant against abiotic and biotic stresses. Based on breeding techniques, better rootstocks can be developed that can mitigate climate risk and other major biotic factors [20]. Molecular approaches are very helpful to deal with heat and



Figure 1. Key phenological stages and management activities [18].

water deficit conditions. Modulation in genes and gene expression related to stress help plants to cope and mitigate stress adversity. Henceforth, somatic hybridization, mutation, somaclonal variation, and genetic transformation techniques help to improve trees for thermotolerance [21].

Thus, in this chapter, we present an overview of climate change i.e., heat and drought stress impact on citrus and its management through agronomic, breeding, and molecular approaches.

2. Climate change

The main reason behind climate change is greenhouse gasses; especially carbon dioxide accumulation in the environment. Fossil fuel burning is the primary source of greenhouse gasses emission. The use of pesticides in agriculture and cutting of forests are also contributing to the proliferation of such gasses that cause climate change. An optimum amount of these gasses is necessary for controlling the earth's temperature, but now their concentration is increasing dramatically. From the expected beginning of human civilization about thousand years ago to 1900, the carbon dioxide concentration in the environment was 0.03%, but now due to climate change, it has been reached to 0.04%, the highest in history [22].

2.1 What is the effect of climate change?

The earth's mean temperature has risen for the past hundred years [23]. The increase in temperature of the earth due to climate change can affect the environment adversely. Today, the average temperature is 4°F more in comparison to the last Ice Age [24]. Global warming is causing the melting of polar caps and warming the ocean's water, which is leading to greater storms and frequent floods along with heavy winds and rains. A heat rise is also enhancing the incidences of wildfires, which damage natural habitat and creatures [25]. Climate change threatens the world's population. The world is severely facing the issue of climate change, especially the third world countries. American and European countries are prepared well against climate change [26]; however, the countries of the







Figure 3.

Global surface temperature anatomy 1880–2018 [28].



Figure 4. Top 17 countries facing the risk of extremely high water stress [29].

Middle East, Asia, and Africa are more exposed to environmental changes due to less preparedness and technology to tackle these issues [27]. Norway would be the country likely to survive climate change due to its low vulnerability against climate change. The neighboring countries: Finland (third), Sweden (fourth), Denmark (sixth), and Iceland (eighth) are well prepared. The countries least likely to survive global warming change include the Central African Republic, South Africa, Eritrea, Chad, Somalia, and the Democratic Republic of the Congo. These countries have poor infrastructure, unstable governance, poor health, and food and water scarcity (**Figure 2**).

2.2 Rise in temperature

NASA center graph associated with climate (**Figure 3**) indicates the average global surface temperature during the era of 1880–2018. After 1940, an abrupt increase in temperature was noted for a duration of two years and then continuous high temperature was witnessed after 1980–2016 [28]. The researchers believe

that global temperature will rise continuously over the next few decades, mainly due to humans generated greenhouse gases. The IPCC (The Intergovernmental Panel on Climate Change) predicts a rise of 2.5–10°F over the next century [29].

2.3 Water scarcity

Water is highly important for plants and its global importance is not difficult to understand. There is a frequent rise in water scarcity due to changes in climatic patterns. It is expected that the world will face a decrease of 66% in water availability up till 2050. The water cycle is adversely affected by climate change. Due to the changing climate, several areas are getting dry. There are 17 nations under the extremely high risk of water scarcity; out of which 12 are in North Africa and Middle East [29]. India and Pakistan, two Asian countries, fell in the list of 17 countries having a risk of water scarcity (**Figure 4**).

3. Effect of heat and water deficit on tree health

The yield of any crop begins to decrease when the temperature exceeds the ideal temperature range and the water level falls below the ideal water demand of the crop. Temperature and precipitation variables of climate are described as diagrammatic sketch alternatively for intensity and duration of drought, which show a small portion of the climate space presently exceeding tree mortality threshold (**Figure 5**). It is predicted that there will be high temperature and drought due to extreme climate change, which can cause severe damage to agriculture and could become a risk for tree populations [31].

3.1 Tree physiology

Plant physiology includes all the dynamic processes of growth, metabolism, reproduction, defense, and communication responsible for plant survival [32, 33]. Heatwaves affect the plant's physiological processes and responses, their ability to tolerate heat, as well as the effectiveness of strategies used for thermotolerance



Figure 5. Climatic variables, temperature, and precipitation, with a range of variability [30].

improvement [34]. In citrus fruits, the temperature above than optimum causes a big difference in the leaf to air vapor ratio as well as high leaf temperature, but the shade conditions can relieve the water pressure and lower the temperature of the leaves [35]. Stomatal conductance and carbon dioxide uptake are reduced by superoptimal leaf temperature and water tension [36].

The gas exchange activity is reduced badly in citrus trees under deficient irrigation [37, 38]. CO_2 assimilation, conductivity, and transpiration rate decrease under water stress, so these gas exchange parameters are a water stress indicator [39–41]. Citrus trees under water deficit conditions reduce the conductivity of stomata and increase photorespiration [42], which reduces the yield, size, and quality of fruits [43]. Under groundwater deficiency, citrus trees lead to stomata closure associated with high ABA content and result in a sudden decrease in photosynthesis [16, 44, 45] and production losses [14, 35]. Chlorophyll *a* is easily damaged compared to chlorophyll *b* due to lack of water. Genotypes/species that maintain stomata conduction under dry conditions also maintain chlorophyll fluorescence and high growth levels [46].

3.2 Tree morphology

Altered temperatures and water deficit conditions affect citrus leaf-to-air vapor pressure during the day in the early morning and midday [8], indicating a vapor pressure of 4.3 kPa at 37–40°C and 6.2 kPa at \geq 40°C, respectively. Citrus exposed to the temperature above than optimum (37°C) and vapor pressure deficit (3.6 kPa) at $330 \,\mu mols^{-1} CO_2$ concentration during midday, shows depression in carbon dioxide exchange rate [47]. Swingle citrumelo, a citrus species increased its total biomass when kept under slightly high temperature [8]. Drought stress affects both plant vegetative and reproductive growth parameters [48, 49]. Citrus under water deficit lessens vegetative growth, fruit size and quality and orchards face a major economic loss [14]. Orange trees exposed to prolonged or excessive water deficit can lead to leaf drop, gradual drying of the tips of the branches, and a drastic decline in fruit production due to severe flower and fruit drop [50]. Young lemons exposed to water stress showed a decrease in daily stem diameter and water flow [51]. The growth of plant roots is dependent on the soil water availability. The roots of irrigated soils are well distributed and widespread compared to roots with less irrigation. Valencia orange roots on the Swingle citrumelo rootstock and a significant difference in root distribution between irrigated and non-irrigated trees were observed [52]. Water stress decreases the growth and metabolism of citrus fruits [43, 53] and increases the cost of extracting juice [21]. Dryness also reduces the thickness of fruit peel, making citrus fruits more sensitive to damage during handling and transportation [54]. Irrigation stoppage during initial and final growth phases of Lane Late orange (Citrus sinensis Osbeck) significantly reduced yield [55]. General studies of water stress in citrus show that the extent and duration of water stress at critical development stages are more vulnerable to the production of citrus. On the other hand, the cultivar and properties of orchards *i.e.*, soil, climate, and cultivation also play important role in success under deficit irrigation [56].

3.3 Tree water status

The transport of water is determined inside the plant by the availability of soil water and relative humidity. Plant physiological adjustments under changing environmental conditions maintain the turgor pressure of the plant cell. In perennial species, seasonal variations in environmental conditions can affect water relationship. In citrus fruits, the large crown and low hydraulic conductivity of the

trunk and roots contribute to severe water scarcity [57]. Transient water deficit in citrus at midday [58] reduces photosynthetic rates [47]. Root hydraulic conductivity decreases under drought stress to prevent the plants from mortality. High temperatures can also increase the loss of root moisture to a harmful level [59]. In plants, heat stress appears as the supply of water is insufficient to meet the evaporation requirement. Heat stress is linked to drought as the plant and soil quickly lose water at high temperatures. It is known that heat stress and drought reduce nutrient uptake and photosynthetic efficiency in plants [60].

3.4 Tree biochemistry

Citrus leaf water potential and leaf abscisic acid (ABA) are the indicator of water stress. Citrus rootstock Rangpur lime grafted with scion Pera orange resulted in decreased leaf water potential and decreased leaf ABA concentration when subjected to water stress [61]. Citrus trees produce endogenous hormones and their regulation by promoting synthesis and accumulation under severe water stress [43]. Plant phytohormones are found in a minor quantity but drought stress accumulates jasmonic acid [62]. Drought synthesizes roots ABA and leaves by transpiration stream [43, 63]. The amount of sugar, like non-reducing sugar (sucrose) and reducing sugar (fructose and glucose), in contrast to the sorbitol content, decreases dramatically over the drought period. Water stress accumulates proline contents, an important osmoprotective agent, and its concentration increases with increasing water deficit conditions in citrus orchard [64]. Proline levels in leaf were recorded in Gada dahi citrus rootstock on day 24 of the water stress in comparison to tolerant rootstocks, which indicated that the accumulation of proline was greater in susceptible genotypes than in tolerant genotypes due to higher stress. Lower accumulation of proline was due to its protective function, removing radicals, maintaining the redox balance, and reducing cell damage [65]. The total phenol content also increases in plants under drought stress, compared to normal irrigated plants [66, 67]. Proteins are involved in several processes that change the plant metabolism under stress conditions and activate the plant defense signal [19, 68]. The protein content of drought-tolerant genotypes is generally higher than that of drought-sensitive genotypes. Carrizo citrange, a tolerant genotype, shows notable soluble proteins in leaves and roots [69, 70]. Higher MDA and H_2O_2 contents observed in plants under water stress indicate greater oxidative damage, which determines the severity of the plant and indicates low efficacy of antioxidant machinery of Carrizo citrange drought-tolerant rootstock [43]. Plants produce several antioxidant enzymes, such as CAT, SOD, and POD to treat the cell damage caused by stress at the oxidative level. SOD is the main enzyme that is expressed under stress, especially under water stress conditions. Carrizo citrange has shown an excellent defense mechanism under water stress with high activity of CAT, SOD, and POD in roots and leaves [19, 68].

3.5 Tree anatomy

Alteration in anatomy by applying heat stress is established. Stress treatment at 40–45°C was given to similar size plants and anatomical changes (size of the epidermis, size of pith, cortex, leaf thickness, epidermal cells, parenchyma tissue) in root, stem, and rhizomes were studied. The thickness of mesophyll, epidermis, and cortex was increased in stressed plants [71]. Some common anatomical changes include increased densities of stomata and trachomatous, cell size reduction, stomata enclosure, and higher xylem vessels in roots and shoots [72]. It has been demonstrated that grafted plant size is reduced on dwarfing rootstock, and such plants are unable to maintain drought or water-deficient conditions [73]. The researchers explain that the vessel density of root and stem are decreased with tree height [74]. The rootstock growth ability is dramatically affected by the number of xylem traits, xylem phloem ratio, vessel size, and vessel density [73, 75]. Maintaining hydraulic conductance of stem, root [74, 76], vessel size and number is the basic factor in hydraulic conductance maintenance [77]. Fewer small vessels may decrease hydraulic conductance, as a result, growth decrease in fruit trees [78]. Water stressed leaf spongy cells have a dense arrangement and reduce the conductivity of leaf diffusion. These results give an idea to understand the direct relationship between mesophilic conductivity and the porosity of the soils [79].

3.6 Tree genetics

Stress-related genes are activated through high temperature and drought, [80], and sugars, different functional proteins, amino acids, and amines are synthesized through these genes [81]. HSPs are the heat shock proteins consisting of a group of genes relevant to heat stress in plants and animals [82, 83]. Heat shock proteins play an important role in maintaining/removing ROS, cell membrane integrity, producing antioxidants, and osmolytes [84, 85]. Heat shock proteins protect plant cells/tissues from drought and heat stress [84]. Citrus HSP70 expression has been examined against water scarcity and high-temperature stress in the Poncirus trifoliata rootstock. In P. trifoliata HSP70 and HSP90 genes against abiotic stress are upregulated [86]. HSP90s play a vital role in signal transduction, cell cycle regulation, protein breakdown, genomic mutation, and protein trade [81, 87]. Aquaporins are transmembrane channel proteins found in tonoplasts, plasma membranes, and other intracellular membranes and are abundantly expressed in plant roots [88, 89]. Major intrinsic proteins (MIPs) are a superfamily of aquaporins that regulate intracellular water passage [90]. The plasma membrane proteins (PIPs) are the most important group of natural proteins that respond to water transport. Overexpression of PIP under abiotic stress conditions confirms the importance of PIP for heat and water stress tolerance [91] as the combination of heat stress and the scarcity of groundwater generally limits the physiology, growth, and productivity of plants [92].

3.7 Tree productivity

The citrus phenological cycle starts from February to next year January in subtropical regions. Flowering starts during February-March and is generally considered a critical period for fruit production. An increase in temperature and water stress after pollination inhibits ovule fertilization [12] which reduces tree fruit set, increases fruit drop, and reduces tree yield [13]. Phases of fruit growth from button size to mature fruit are more sensitive to deficit irrigation and heat stress. Hence, reduced fruit growth, and delayed ripening occur which are associated with a decrease in fruit size, increase in fruit acidity, and low tree yield [13]. Drought at a pre-harvest stage in oranges develops wrinkle on fruit peel [17]. An increase in optimum temperature at fruit pre-harvest causes fruits drop and reduced yield. Citrus under different phenological stages respond to deficit irrigation or water stress and contribute negatively to yield/production and fruit quality. In an experiment, eleven-year-old sweet orange scion grafted on Carrizo citrange were evaluated against water stress and revealed 10–12% relative yield decline. Gonzalez [37] compared Clementina (Citrus clemen*tina*) tree performance under 25–50% deficit irrigation during initial fruit enlargement and pre-maturation phases and recorded a significant negative effect on fruit yield [13]. Navelina sweet orange (Citrus sinensis Osbeck) yield reduced significantly

when irrigation was reduced at 55% with respect to crop water requirement during flowering and fruit set.

4. Management of citrus under climate change

4.1 Agronomic management

Implementation of proper orchard management practices decreases the adverse effects of heat and drought stress. The management includes trees requirement based nutrition and irrigation techniques, organic and synthetic mulches, as well as selecting the most suitable cultivars/rootstocks that are resistant to various stresses.

The selection and development of new rootstocks tolerant to biotic and abiotic stress is inevitable for the stable production of citrus under the scenario of climate change. New and known diseases and environmental conditions also help to force developing new citrus rootstocks according to the demand [93]. Citrus rootstocks like Volkamer lemon (*C. volkameriana*), Rangpur lime (*C. limonia*), and Rough lemon (*C. jambhiri* Lush.) resist water stress and increase the production of cultivars grafted on these.

Fertilizer application can also be helpful to manage plants against abiotic stresses [94]. Application of Ca and K macronutrients and B and Mn micronutrients modify the function of stomata under heat/high-temperature stress [95]. K, Ca, B and Mn activates physiological and metabolic processes that help maintain a high water potential in tissues, which increases tolerance to heat stress [96]. The use of N, K, Ca and Mg also reduces the toxicity of ROS, thereby increase the levels of antioxidant enzymes in plant cells [96].

Plant growth regulators (PGRs) in managing water and heat stress also play an important role. PGRs like cytokinins, abscisic acid (ABA), and salicylic acid play role in resistance to heat and drought. The application of PGRs increases the water potential and chlorophyll content in citrus trees [97]. The exogenous use of ABA increases productivity in the absence of water [98]. ABA formulations are available with commercial manufacturers to improve the drought tolerance of trees [99, 100].

Mulching underneath the trees is often used as a technique for water conservation [101, 102]. Mulches are used to maintain moisture levels high in the soil, control soil temperature, and evaporation [93, 103]; thereby reduce the need for irrigation during growing seasons [104]. The need for water in the soil is decreased and the ability to withstand drought and heat is increased by using mulches [105, 106]. Plastic films are more effective than organic compost for groundwater protection [107].

4.2 Breeding strategies

Citrus rootstock breeding programs are aimed to combine biotic and abiotic tolerance/resistance in new rootstocks. However, conventional plant breeding (**Figure 6**) in mitigating the abiotic stresses has limited success against plant productivity [108]. Similarly, developing better rootstocks through breeding by the conventional method is a long-term approach due to many difficulties, particularly the complexity of citrus biology (high heterozygosity, long juvenility, polyembry-ony) [109, 110]. Typically, from a breeding program, it takes at least 15 years for a new standard variety to emerge in the citrus industry. Moreover, a sexual hybrid is difficult to identify at an early stage. In this case, trifoliate leaf (a morphological marker) is used as a male parent and unifoliate as a female for identification of sexual hybrid at the seedling stage [111]. The trifoliate trait is dominant, and the



Figure 6.

Traditional cross-hybridization in citrus. (A) a large unopened bud, (B) emasculation, (C) pollination of the emasculated flower, (D) bagging of the pollinated flower, (E) bagged twig, (F) general view of the seed parent after crossing.

seedlings showing the parental trifoliate pollen phenotype are considered hybrids. In absence of a trifoliate pollen parent, the hybrids can be identified by using SSR and RAPD molecular markers [112].

Besides citrus biological constraints, valuable traits like resistance to low/high temperature, root rot, viruses, nematodes, salinity, and drought are important to incorporate in rootstocks. A list of some important biotic and abiotic traits of citrus rootstocks are presented in **Table 1**, which can help in the development of new tolerant/resistant rootstocks.

Several rootstock hybrids have been released to the citrus industry worldwide. Swingle citrumelo rootstock is a hybrid of Duncan grapefruit and Trifoliate orange, crossed by Swingle in 1907 and released in 1974. Since then, it has been used successfully as the standard rootstock for better traits *i.e.*, moderate drought

Rootstocks	Alkalinity	Chloride	Drought	Tristeza	Nematodes	Phytophthora	Fruit quality
Cleopatra mandarin	Т	ТТ	рТ	Т	S	рТ	G
Chios mandarin	Т	S	рТ	Т	S	SS	_
Sweet orange	Т	рТ	Т	Т	S	SS	G
Macrophylla	TT	TT	TT	S	S	Т	L
Pomeroy Poncirus	SS	SS	S	R	Т	Т	Н
C-35 citrange	S	S	Т	R	Т	Т	Н
4475 citrumelo	S	рТ	рТ	R	Т	Т	Н
Citrandarin	рТ	рТ	_	R	рТ	Т	G

Abbreviations in the table: T; tolerant, TT; very tolerant, pT; poorly tolerant, S; susceptible, SS; very susceptible, G; good, L; low, H; high.

Table 1.

Some biotic and abiotic traits of selective citrus rootstocks [113].

tolerance and high fruit quality. Citranges are hybrids of Washington Navel orange and *Poncirus trifoliata*, out of which Carrizo and Troyer are the two main citranges. These can tolerate water shortages and produce excellent quality fruits. Benton citrange is a cross of Ruby blood orange and trifoliate orange developed in late 1940 and is more tolerant to heat and water scarcity [114]. Brazilian sour orange also shows tolerance against heat, drought, and their combined stress [85].

4.3 Biotechnological interventions

Hybridization by somatic approaches is a protoplast fusion process that has become an important tool for plant production, combine (partially or totally) desired cultivars somatic cells, species, or genera, resulting in the development of new genetic combination. In addition to intergenerational mixtures in somatic hybridization, more emphasis has been placed on interspecific mixtures between *C. reticulata* and *C. maxima* [109, 110] to meet the specific needs of the citrus industry. Poncirus is drought-prone, while Citrange C-35 is more drought-tolerant. Among these rootstocks, 4475 citrumelo have the best ability to adapt to the environment. Cleopatra mandarin + *Poncirus trifoliata* and Cleopatra mandarin + C-35 Citrange somatic hybrids have resistance to CTV, tolerance to nematodes, and phytophthora. The Sweet orange + *Poncirus* and the Sweet orange + C-35, as well as the Sweet orange + Citrumelo 4475, can adapt to low moisture soils and tolerate biotic stresses. Macrophylla is a productive rootstock and adapts well to saltwater, limestone, and water stress [115].

In vitro mutagenesis and somaclonal variation are important tissue culture techniques being used in citrus improvement. Somaclonal variation, genetic and phenotypic variation between plants, can be used to improve citrus cultivars under conditions of water and heat stresses. Genetic improvement by *in vitro* selection of Satsuma mandarins (*Citrus unshiu* Marc.) has been made successfully; however, the frequency of somaclonal variation by factors, including genotypes, explant culture length, sources, and environmental composition [116, 117]. Cell lines success stories of some salt-tolerant cultivars are *C. sinensis* cv. "Shamouti" [118], *C. limonium* [119] and "Troyer citrange" [120].

Genetic transformation is an alternate technique for citrus genetic improvement. PEG-mediated genetic transformation of citrus fruits is a direct DNA transfer method [120] that seeks to express an aminoglycoside phosphotransferase II gene in isolated protoplasts from sweet orange (Citrus sinensis Osbeck) culture for suspension. The genetic transformation of citrus fruits has mainly been carried out from young materials such as embryogenic cells from the epicotyl segment of *in vitro* germinated seedlings. Excess protein for late embryogenesis (OHL), heat shock proteins, and certain transcription factors that affect the expression of various stress-related target genes have also been used to improve drought tolerance in transgenic plants. Drought-induced genes with different functions have been identified through molecular and genomic analyses in a variety of plant species such as the C/CBF family (Shinozaki) [121]. By regulating stress gene expression and signal transformation, plants indirectly become more stressresistant [97, 122]. The TDF genes have been identified as drought-induced and the proteins encoded include fructose aldols bisphosphate, a cold-like protein found in WCOR413. The PIP2 protein, an aquaporin specializing in a water channel to transport water across the plasma membrane, and the tonoplast have been observed in sweet orange. TDF21, TDF38 and TDF80 are involved in the regulation of signal transduction and expression of genes. These are sensitive to stress and also regulate the expression of stress-induced genes, possibly induced by drought.

5. Future research strategies

Soil microbiome research offers the opportunity to improve abiotic stress in plants. The mechanisms by which plants recover from drought and/or heat stress can be mediated by microbes surrounding the plant, particularly the roots, and these are involved in various stages of plant growth. Advances in the application of new molecular and genomic tools and technologies have paved the way for the study of plant microbiota, and these promising advances enable the study of the biological functions of various microorganisms both inside and outside the host tissue.

Significant advances in the characterization of the plant genome and the optimization of techniques for manipulating the plant genome have contributed and will further improve our knowledge and ability to develop stress-resistant plants. Ultimately, genetic engineering or transgenic methods must be combined with conventional breeding activities and supported by markers in order to obtain the desired improved varieties.

Plants have developed complex adaptive mechanisms to withstand diverse and complex abiotic stresses. With the advent of new technologies such as genomics and genetic transformation, significant advances have been made in understanding these complex traits in higher plants. However, the commercial application of successful research results requires additional validation of the products or prototypes in the field.

These efforts will lead to tangible practical outcomes that may help mitigate the effects of climate change, especially concerning drought and heat stresses, and will contribute to improved crop productivity and food security.

6. Conclusion

Plants adaptation is considered a striking strategy to manage the impacts of climate change. In climate change, the most important factors are fluctuating patterns of temperature and drought which have an adverse effect on plants physiology, morphology, water status, biochemistry, anatomy, genetics, and productivity. Hence the emphasis should be on the development of production systems for improved water-use efficiency and to adapt to the hot and dry conditions through agronomic practices. Development of climate-resilient citrus rootstocks and scion through genomics and biotechnology are essentially required.

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Conflict of interest

The authors have no conflict of interest with any person or institution.

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Section 3

Citrus Nutritional and Nutraceutical Importance

Chapter 9

Citrus Fruits: Nutritive Value and Value-Added Products

Abu Saeid and Maruf Ahmed

Abstract

Citrus fruits are essential sources of food and energy and play a critical role in supplementing healthy diets. Citrus fruits contain mostly carbohydrates such as sucrose, glucose, and fructose and are good dietary fiber sources, which help prevent gastrointestinal disease and promote high circulating cholesterol. Besides, citrus fruits are also significant sources of vitamin C and various bioactive compounds. It is suggested that these components are of vital importance in improving human health due to their antioxidant properties and being converted to vitamin A. However, citrus fruit is still being used for different purposes like juice, jam, jelly, squash, pies, cake, candies, marmalades, etc. Most citrus waste materials are currently used as animal feed. Innovations are occurring in the conversion of citrus by-products into valuable commodities with the development of innovative technologies. This chapter has put up primary and secondary research findings of citrus fruits, especially lemon and pomelo, their chemical properties, composition, and their use in health and cosmetic needs.

Keywords: citrus, lemon, pomelo, nutritional properties, value-added products

1. Introduction

Citrus is an evergreen shrub that belongs to the Rutaceae family from South Asia, China, India and the Malay Archipelago, which is native to the subtropical and tropical regions of Asian regions [1]. The genus of citrus includes sweet orange (C. sinensis: 61.1 % of world citrus production), tangerine (C. reticulata: 19.9 %), limon and lime (*C. limon* and *C. aurantifolia*: 12.1 %) and grapefruit (*C. paradisi*: 5%). Minor types of citrus, which constitute much of the remaining 2.0%, include sour orange (C. quarantium), shaddocks (C. grandis), citrus (C. medica), which seem to be promising sources for many beneficial human nuts [2]. Citrus fruit is divided into two sections like peel and flesh (**Figure 1**). Peel is made from epicarp or flavedo (colored peripheral surface) and mesocarp or albedo (white soft middle layer). The peel (60-65%), internal tissues (30-35%), and seeds (0-10%) comprise citrus fruits [3]. Citrus fruits provide carbohydrates, such as sucrose, glucose, and fructose mostly. Fresh citrus fruits are also an immeasurable source of dietary fiber associated with gastrointestinal disease prevention and lowered circulating cholesterol. Citrus fruits also have a distinct aroma and delicious taste along with low protein and fat content.

Citrus fruits also provide the most potent source of vitamins C and B (thiamines, pyridoxines, niacins, riboflavin, pantothenic acids, and folate). The fruit also leads to the use of phytochemicals, such as carotenoids, flavonoids, and limonoids [1].



Figure 1. *Structure of the citrus fruit* [2].

Citrus phytochemicals contain antibacterial, antiviral, antifungal, anti-carcinogenic, anti-thrombotic, or anti-inflammatory agents [4]. Several studies have proposed citrus fruit evaluation as a healthy and delicious diet [5]. Prior research suggested that citrus and citrus products are rich sources of vitamins, minerals, and dietary fibers [6]. However, the bioactive and non-nutrient compounds in citrus are appreciated to reduce the risk of various chronic diseases [7].

Citrus fruits are eaten as fresh goods and juice throughout the world. Peel is discarded as waste containing many secondary components with significant antioxidant activity related to other fruit portions [8]. In recent years, flavonoids such as polymethoxy flavones (PMFs), which are present in citrus fruits, have been attracted growing attention by their antioxidants [9] and anti-cancer properties [10]. Various bioactive compounds in citrus peel extract and powder may reduce overall cholesterol, triglycerides, LDL, and glucose levels [11]. Citrus by-products produce a range of value-added products, including essential oils, pectin, enzymes, single-cell collagen, natural antioxidants, ethanol, organic acids, and prebiotics. Orange, lemon, mandarin, and grapefruit contained essential oils show antifungal activity upon the fungi A. niger, A. flavus, P. chrysogenum, and P. verrucosum. The essential oil may be regarded as acceptable for the food industry as alternatives to chemicals [12]. Pectin extracted from *Citrus* peel is used in various industrial food processes as gelling agents, including jam, jellies, and as thickener, texturizer, emulsifier, and stabilizer in dairy products. Pectin is also used to jellify properties in the pharmaceutical, dental, and cosmetic industries [13]. Therefore, this chapter highlighted the nutritional values of major essential nutrients such as Vitamin C, carotenoids and vitamin A, Folate, Dietary fiber, flavonoids, and limonoids, as well as value-added products such as food ingredients, pectin, essential oil, enzymes, a natural antioxidant, and packaging film retrieved from citrus especially lemon and pomelo fruit.

2. Characteristics of citrus fruits (lemon and pomelo) and their chemical compositions

2.1 Lemon

Lemon (*Citrus limon L.* from Rutaceae) is one of the most common globally and ranks third among the Citrus species globally by 4,200,000 metric tons after orange and mandarin [14]. Lemon fruits typically consist of three parts: pulp, skins (albedo and flavedo), and seeds. It offers an extensive supply of natural compound

products such as citric acid, ascorbic acid, minerals, flavonoids and essential oils [15]. Lemon bioactive compounds like flavonoids, vitamins, minerals, dietary fiber (**Table 1**), and essential oils are used in the food, cosmetic, and pharmaceutical industries. Most by-products of the lemon juice industry can provide functional foods with nutritional substances such as non-digestible carbohydrates, dietary fiber and bioactive (flavonoids and ascorbic acid). Lemon fruits can function against photo-oxidamage because carotenoids exist. Lemon fruit, rich in flavonoids, has a significant role in the healthy diet, particularly in preventing diseases such as obesity, diabetes, lowering blood lipids, cardiovascular disease, and some forms of cancer [18]. The citrus fruits used for direct consumption or converted into juices, jam, jelly, molasses, lemoncello beverage and more in addition to the lemon skin are added value products such as pectins, essential oil and functional ingredients [12, 18].

2.2 Pomelo

Pomelo is one of the most commonly grown and eaten citrus fruits and orange, mandarin, lemon, and grapefruit [19]. Pomelo (**Table 1**) is a promising source of carbohydrates, proteins, fiber, vitamins and minerals originating in warm tropical climates in south-eastern Asia [20]. The presence of bioactive (carotenoids, lycopene,

Component	Lemon (Citrus limon)	Pomelo (Citrus maxima)
Moisture (g/100g)	84.2	87.0
Fiber (g/100g)	1.6	1.60
Carbohydrate (g/100g)	10.8	11.5
Protein (g/100g)	0.9	0.6
Fat (g/100g)	0.8	0.20
Vitamin (mg/100g)		
β-carotene	50.0	120
Thiamine	0.02	0.03
Riboflavin	0.01	0.03
Vitamin C	37	20.0
Mineral (mg/100g)		
Ca	70.0	10.0
Mg	12.0	21.6
Na	1.50	2.70
Р	10.0	20
K	148	106
Fe	0.23	0.40
Zinc	0.12	0.15
Cu	0.20	0.19
Total phenol (mg GAE/g)	204.40	70.56
Total flavonoids (mg QUE/g)	27.50	13.06
Carotenoid (mg/100 mL)	0.31-0.35	0.72-0.73

Table 1.

Chemical composition of Citrus fruits as [15–17].

polyphenols, flavonoids, limonoids, fiber and vitamin C) contributes to their protection against oxidative stress, hyperglycemia, and high blood pressure. Due to its essential health promotion properties, pomelo segments in food products are growing in importance in producing functional foods [21]. Pomelo is eaten fresh or made into juice [19], or pomelo fortified noodles help the diabetic population [21]. On the other hand, researchers have investigated alternative ways of restoring pomelo peels to the advantage of value-added products such as pectin, essential oils, polysaccharides, phytochemicals [19]. Production of juice and consumption of fresh fruit create large quantities of agricultural waste. The main components of wet Pomelo Peel waste, like other citrus fruits, include water, cellulose and hemicellulose, soluble sugars, lipids (mainly D-limonene), and bioactive compounds (i.e., polyphenols, mostly flavonoids).

3. Nutritional values of citrus fruits

Citrus has many natural plant compounds such as vitamin C, carotenoids (some can convert to vitamin A), folic acid, flavonoids, and fiber. **Table 2** shows the amount of vitamin and mineral consumption in lemon and pomelo fruits.

3.1 Vitamin C (ascorbic acid)

Citrus is a valuable source of vitamin C. By consuming a moderate amount of citrus fruits each day, an individual can achieve 100 percent Vitamin C level. Vitamin C is an essential water-soluble vitamin essential for the body's defense [22]. It is transmitted through muscle fibers, carnitine biosynthesis, neurotransmitters, collagen, and bones because these particles connect the fibers. The immune system

	Vitamin C	Vitamin A	Folate	Fiber
Oranges	53-88 mg	17 µg	30 µg	2.4 g
Children under 9 y (%)	213-589	3-6	15-20	10-13
Persons 9+ y	59-195	2-4	8-10	6-11
Pregnant/Lactating women	44-110	2-3	5-6	8-9
Grapes	31-61 mg	58 µg	13 µg	1.6 g
Children under 9 y	125-244	12-15	7-9	6-8
Persons 9+	35-135	6-10	3-4	4-8
Pregnant/lactating women	26-76	4-8	2-3	6
Tangerines	27-72 mg	46-144 μg	16 µg	1.8 g
Children under 9	107-480	9-36	8-11	7-9
Persons 9+	30-160	5-24	4-5	5-9
Pregnant/lactating women	21-90	4-19	3-4	6
Lemons/limes	29-61 mg	2-22 µg	11-16 µg	1.8-2.8 g
Children under 9	116-407	0.4-6	4-7	11-15
Persons 9+	32-135	0.2-4 2-4		9-13
Pregnant/lactating women	24-76	0.2-3	1-2	10

Table 2.

The number of nutrients and the percent of the recommended daily allowance or adequate intake met from the consumption of 100 g of selected citrus fruit [22].

can be effectively stimulated by consuming vitamin C, which boosts white blood cells [23]. When Vitamin C is taken for pregnancy, it can decrease pre-eclampsia risk [24]. Some studies indicate that vitamin C supplementation can reduce the severity of colds symptoms or duration [23]. Anti-oxidants such as Vitamin C could reduce the risk of artery stiffening and cardiovascular diseases [25]. Above 200 mg of vitamin C daily is a healthy intake, and citrus fruits are a huge source of this vitamin. Lemon provides 37 mg of ascorbic acid per 100 g of fruit [16]. Pomelos have 52.3 mg of ascorbic acid in 100 g of the flesh [26].

3.2 Carotenoids and Vitamin A

There are many types of carotenoids, including terpenes responsible for pigments commonly found in plants, and there are about 600 carotenoids in foods and 50 in human bodies [27, 28]. The highest carotenoid levels, such as lutein, zeaxanthin, lycopene, and vitamin A, are found in fruits and vegetables, including orange and carotene. Benefits of carotenoids in foods include improving immune function, promoting bone formation, promoting eye health, and maintaining visual quality [22]. There is a large amount of data supporting that carotenoids reduce the risk of cancer, macular degeneration, cataracts, skin damage to the sun, and cardiovascular diseases [29]. Higher consumption of β -carotene is linked to a lower breast cancer risk [30]. Beta carotene, lycopene, or lutein may decrease the rate of UV-induced lipid peroxidation in human skin fibroblast cells [30]. Lutein is inversely related to colorectal cancer in both men and women [31]. The levels of lutein, zeaxanthin, β -cryptoxanthin, and β -carotene in the lemon and pomelo, were around 2.95, 0.81, 0.81 and 10.3 (µg/g, db), respectively [32]. The content of carotenoids in pummelos' peel was 0.012-0.015 mg/gdb [33].

3.3 Folate (folic acid)

Folic acid, which is a water-soluble vitamin, and its derivatives are collectively called folate or folacin. The most notable folate compounds in Citrus are the reduced 5-methyl tetrahydrofolate (monoglutamate) and polyglutamate compounds [34]. Folate plays a vital role in DNA, which is involved in homocysteine regulation and protein production primarily through the methylation transfer reactions [22]. Because there is a high DNA production during pregnancy, a folate deficiency is significantly linked to birth defects such as neural tube defects [35]. Lack of folic acid caused higher levels of homocysteine, raising heart disease and atherosclerosis [22]. Previous studies show that citrus fruits' daily consumption can help improve folate levels, which will subsequently decrease blood homocysteine (tHcy), thus reducing cardiovascular disorder and neural tube defects [36]. Citrus is a parallel source of dietary folate that can help to cover up to 10% to 20% of the recommended daily allowance of adults, children, and infants with a consumption of 100 g of citrus fruits. The consumption of citrus fruit is an easy way to obtain vitamin C and dietary folate, which is vital for absorption in the body. Lemon, a citrus fruits representative, has eleven to sixteen micrograms of Folate in 100 grams [22]. According to El-Otmani and Ait-Oubahou [37] Citrus limon contained 11mg of folic acid per 100 g of citrus.

3.4 Dietary fibre

The fiber is found in vegetables and fruits cannot be digested and absorb in the small intestine. There are two kinds of dietary fiber; soluble and insoluble fiber. Insoluble fibers are highly fermentable and connected with carbohydrate and lipid

metabolism, while soluble fibers contribute to fecal bulk and reduce transit time [34]. Although pectin, cellulose, and hemicellulose comprise the most abundant dietary fiber on the plants, they also contain only trace amounts of lignin. Pectin is citrus' primary fiber, which occurs primarily in citrus peels and rinds. Consumption of citrus fruit can contribute significant quantities of pectin in a diet. Dietary incorporation of pectin appears to affect several metabolic and digestive processes; principal interest affects glucose absorption and cholesterol level [38, 39]. There is a significant benefit in consuming citrus fruit because of its pectin content. Dietary incorporation of pectin appears to have many implications for metabolic, digestive, and health affairs. One way fiber can reduce colon cancer is by diluting and trapping the harmful chemicals in the colon from bile-absorption and bile-excretion [34]. Scientific studies have proven that fiber can help promote laxation and satiety, the uptake and reabsorption of glucose, fat, cholesterol, and bile acids, thereby lessening heart disease risk and possibly enhancing healthy intestinal microbial fermentation [40, 41]. Citrus fruits significantly reduce cholesterol levels depending on the esterification degree of fiber consumption, viscosity, and molecular mass [22]. A fiber-rich diet has a low risk of deadly chronic diseases such as diabetes, heart disease, weight, and cancer and lowers cholesterol levels and blood sugar [42]. Several epidemiological studies reported that citrus peel support reducing plasma liver cholesterol, total serum cholesterol, serum triglyceride levels, and total liver lipids [43].

3.5 Flavonoids and limonoids

Citrus pulp, peel are rich sources of flavonoids. Toh et al. [44] found that pomelo peeled (356.95 mg/QE) had higher total flavonoid content than pomelo pulp (13.06 mg/QE). Makni et al. [15] found the amount of quercetin in lemon flesh (56.16 mg Eq Quercetin/g dry weight) was higher than in peel (27.50 mg Eq Quercetin/g dry weight). Citrus fruits are also rich in flavonoids such as hesperidin, hesperetin, naringin, naringenin, diosmin, quercetin, rutin, nobiletin, tangeretin, and others [45]. Citrus flavonoids have both antioxidant and anti-inflammatory properties, and because of that, it can increase the antioxidant capacity and effect reducing cholesterol and triglycerides levels and provide more excellent bone health [22]. Preclinical studies and clinical trials demonstrated that flavonoids' effects in the forms of hesperidin and its aglycone hesperetin prevent various types of diseases, including neurological, psychiatric, and cardiovascular disorders [46]. Over the years, naringin and hesperidin are gaining attention for their great antioxidant capacity, contributing sweet flavor to foods and beverages [47]. Some known naturally occurring flavonoids have potency in defending against certain types of RNA and DNA viruses [48].

Limonoids are also known as flavonoids, which are compounds found in citrus fruits. In citrus fruits, there are two groups of limonoids: aglycones and their corresponding glucosides. Bitter taste in citrus results from limonoids present. Limoninin's most important constituents are glycosides called limonin and nomilin [22]. In animal and human cell lines, limonoids slow down the development of aggressive cancers like the pancreas, colon, stomach, and breasts. On the other hand, limonoids are also reported to reduce skin cancer in animal models. Limonoids are known for their medicinal or health beneficial effects like anti-cancer, anti-microbial and antimalarial activities [49]. Limonoids have antibacterial and antiviral effects. Some limonoids are known to stimulate the in vivo production of the detoxifying enzyme glutathione S-transferase in the liver and inhibit the formation of chemically induced tumor cells in the oral cavity, forestomach, small intestine, colon, lung, and skin of



Figure 2.

Proposed model for signaling pathways leading to growth inhibition by obacunone in estrogen-responsive breast cancer (MCF-7) cells [53].

animals [50, 51]. Limonoids found in citrus fruits decreased the spread of cancer cells in animal studies [52]. Studies showed that lemon-lime oil in the form of obacunone appeared to prevent breast cancer by inhibiting aromatase enzyme and antiinflammatory pathways [53] (**Figure 2**). Several studies have revealed that limonin and nomilin are found in fruits, pulp, and seeds. Pummelo juice contained 18 ppm of limonin and 29 ppm of total limonoid glucosides [54]. Wattanasiritham et al. [55] reported limonin content of 18 ppm in the juice of pomelo cultivars. Limonin levels in extracted juice from seven pummelo cultivars from Florida ranged from 10.07 to 29.62 ppm. As shown by analysis, mature lemon seeds contain 1300 µg/g of different limonoid glucosides on a fresh weight basis [56]. The mature seeds contain much higher amounts of glucosides than commercialized juice, therefore. Fong et al. [57] documented that the commercial lemon juice had only 82 µg/g of glucosides.

4. Value-added products

Citrus fruits are known for being highly fragrant, with a tart taste and higher vitamin C content. The world has a wider variety of citrus fruit because of the continued type changes, such as sour oranges, oranges, pummelos, lemons, and limes, among others [22]. Nowadays, citrus pulp/pomace, seed, and peel are used for various commercially valuable products such as food ingredients, pectin, essential oils, enzyme production, a natural antioxidant, and packaging film formation.

4.1 Food ingredient or food products

The nutritional supplement of pomelo fruit segments has been added to products to be developed like noodles prepared with 30% new segments and 5% dry. These noodles can satisfy those with regular diabetes and the general public [21]. A high dietary fiber food was created by reducing the dietary fiber-rich pomelo peel to a powder that contained nearly 50% of dietary fiber. Lemon fruit is usually eaten fresh, but it is also processed to make juices, jams, jellies, molasses, candies and much more [18]. Lemon juice has been used as a coagulant during the manufacture of wara cheese [58]. Another innovation implemented in the beef burger is to use lemons for "enhancing the cooking properties of the burger" [59]. Lario et al. [60] had reported that the high-fiber lemon powder extract from lemon peel debris by-products is an ideal additive for food products (as meat, dairy, and bakery products).

4.2 Pectin

Pectin is an agent of gelling, emulsifying, stabilizing, texturizing, which appears as a white to light brown powder broadly accepted as a functional ingredient [61]. Fruit peels are a highly desirable pectin source because they cover up to 20% of the fruit's total Pectin [62]. Pomelo is a highly valued source of natural Pectin. About 20.75% pectin is derived from lemon peel for jams. In the study, the high extraction (36.71%) of Pectin from lemon peel has something to contribute to this industry [63]. Moneim et al. [64] recommended utilizing 20.75% of the lemon peels' total weight in making pumpkin jam. The researchers were added 16.740% of Pectin from pomelo peel to the pressed carrots before storage [65]. Studies by Methacanon et al. [66] have shown that pectin yield was 23.19% for pomelo peel. On the other hand, Roy et al. [65] were found that pomelo peels are a good source of Pectin, and then carrot jam made by extracting Pectin from pomelo peel.

4.3 Essential oil

Essential oils (Eos) are volatile, complex, natural mixture of aromatic oils obtained from plants [67]. Citrus essential oil is commonly known to produce a good fragrance and has been officially approved for healthy public consumption. All over the world, EOs are used in cosmetics, perfumery, toiletries, flavoring, beverage, pharmaceuticals and other personal hygiene products [68–70]. Lemon oil is often used on the skin because of its antimicrobial and antifungal properties. Pomelo peel (PP) has approximately 299 recognized volatile compounds. A significant number of these volatile compounds is considered as terpenoids (189 volatiles, 63.2%). Various kinds of chemicals released are monoterpenoids, monocyclic monoterpenoids, bicyclic monoterpenoids, diterpenoids, acyclic sesquiterpenoids, monocyclic sesquiterpenoids, bicyclic sesquiterpenoids, and tricyclic sesquiterpenoids. Another major volatile present in PP EO is nonterpenoid alcohols (4.7%), nonterpenoid aldehydes (6.0%), nonterpenoid hydrocarbons (5.7%), and esters (8.7%). The unknown volatiles covered 11.7% (35 volatiles) of total volatile compounds. The structures of widely known terpenoids in Polypropylene EO has shown in **Figure 3a,b**. The most critical monoterpenoids (1 to 16) and sesquiterpenoids (17 to 29) are present in PP EO [19]. According to studies conducted, lemon essential oils retain aroma in foods because of their natural preservative and flavoring properties [71]. The effect of lemon essential oils on the cheesemaking process dramatically reduces microorganisms' population, especially those of the Enterobacteriaceae family [58]. In the food and pharmaceutical industries, citrus



Figure 3.

(a) Major monoterpenoids in pomelo peel essential oils (1-16); (b) Major sesquiterpenoids in pomelo peel essential oils (17-29).

EOs can be employed to inhibit mold and fungal growth. Lemon EO (Citrus limon) was used as a possible fungicide to manage the pathogenic fungi attacking grapevines, namely Eutypa sp., Botryospaeria dothidea, and Fomitiporia mediterranea. The antifungal activity was observed for EO against all the three fungi with the highest action against strain Eutypa sp. (82% inhibition) and the lowest tolerance (33.1%) towards F. Mediterranean [72]. These essential oils repress the growth of mold and yeast. The raspberries coated with alginate and lemon EO (0.2 percent) or orange EO (0.1 percent) halted bacteria, yeast, and mold and also reduced the quality deterioration right after harvest [73]. The lemon EOs mixed in the chitosan films can be used to control L. Multicellular pathogens in refrigerated foods Researchers Rahmawati et al. [74] observed that an edible coating applied with lemon essential oil only delayed aging of tofu and fresh strawberry.

4.4 Enzymes

The most basic usage of citrus peels is to produce the pectinolytic enzyme for beneficial purposes. Larios et al. [75] studied endo-polygalacturonase production by Aspergillus sp. CH-Y-1043 using untreated lemon peel and citrus pectin as carbon sources. Lemon peel being employed as a substrate in submerged cultures to obtain high pectinase titers by A. flavipes FP-500 and A. terreus FP-370 [76]. A. niger produces approximately 2,181.23 U/L pectinases from lemon, peel pomace in a solid-state reactor [77]. Seyis and Aksoz [78] have shown that lemon pomace and peel are suitable substrates for heterotrophic xylanases enzyme production using fungus Trichoderma harzianum. Aspergillus niger LFP-1 was studied in solidstate fermentation (SSF) using pomelo (Citrus grandis) peels as a substrate [79]. Maller, et al. [80] determined that lemon peels are extremely capable of triggering the production of Polygalacturonase in the aspergillus niveus. Pectin lyase yield increased through fungal strain Aspergillus oryzae process derived from lemons peel and used in solid-state fermentations [81]. Studies said that Polemo pericarp powder utilized as a substrate for Aspergillus oryzae JMU316 has Naringinase enzyme [82]. Pectinase enzyme produced from pomelo peels by By Aspergillus niger through Solid State Fermentation [83]. Lemon peels could be a good source of naringin, which could be used as a carbon source in submerged fermentation for naringinase production using Aspergillus niger [84]. Naringinase is essential for the production of sweetener precursors, preparation of prunin, aroma enhancement in winemaking, biotransformation of antibiotics, and rhamnose manufacturing [82].

4.5 Natural antioxidant

Antioxidants are chemical substances that can reduce or prevent the damage caused by free radicals in the body, thus reducing the risks of cardiovascular disease and cancer [85]. Results of studies showed that lemon peel contained almost 75.9% of antioxidant content. The unique ability of Paneer was derived from compounds found in the Peel of orange, lemon and pomegranate [85]. Peel taken from Tambun White pomelo type contains higher levels of antioxidants and is also a rich source of natural antioxidants [44]. Lemon peel and flesh had the highest antioxidant capacity, and they had a significant impact on the prevention of cardiovascular diseases and other diseases [86].

4.6 Packaging film formation

More sustainable, biodegradable plastic has gained popularity among environmental scientists. The researcher created plastic films from citrus peels. By applying the peels of citrus, biodegradable packaging material could be made [87]. Wu et al. [88] prepared fruit peel as the edible packaging film with high content biopolymer to form film for packaging. The film was designed to incorporate tea polyphenols, which causes interacting molecules to become more closely crowded. Soy protein with essential oil of lemon peel was used to create a degradable film and create cheese curdorants for preservation [89]. Dias et al. [90] reported that the use of citrus essential oil and its aroma significantly improved consumers' health and significantly increased the acceptance of biscuits' packaging. Das et al. [91] demonstrated that chicken feather keratin combined with pomelo peel pectin to form biodegradable composite film and wrapping of fried fish fillets resulted in less weight loss, hardness value, and reduction in the surface microbial count.

5. Conclusion

Citrus has positive effects on human health, and it could be an essential raw material to the biotechnological industry. Citrus is a mighty source of vitamins, minerals, and dietary fibers. Bioactive and non-nutrient compounds in citrus are valuable for controlling chronic diseases such as diabetes, cholesterol, obesity, cardiovascular disease, and some forms of cancer. Besides citrus, vitamin C also

has other benefits, including fighting diseases such as cardiovascular disease, boosting white blood cells, immune function, and symptoms or duration of colds. Peel, flesh/pomace, and seed from the citrus fruit are employed in making different novel foods like noodles, extract pectin, enzyme extracts, and essential oil. Therefore this information might be necessary for the readers because it gives facts about the popular citrus fruit. Also, choosing the best citrus for an edible ingredient can be beneficial for citrus processors.

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Chapter 10

Citrus Essential Oils: A Suite of Insecticidal Compounds

Bulbuli Khanikor, Kamal Adhikari and Bikash Rabha

Abstract

Citrus essential oils (CEOs) and their constituent compounds are being reported to have multifarious activities. In this chapter an attempt is made to discuss the insecticidal activities, as well as CEO profile of different vegetative part of Citrus species and biocidal potentiality of their constituent compounds against diverse insect pests. It is observed that in most of the CEO constituent profile, limonene is the major constituent compound. Other important constituents present in different percentages in different CEOs are β -citronellal, linalool, pinene, β - caryophyllene, β -myrcene, terpinene, citral etc. These plant EO constituents are reported to have insecticidal effects against diverse insect species. Taking the four peel EOs of Citrus limon, Citrus paradisi, Citrus medica, Citrus maxima commonly grown in North Eastern part of India, study on their insecticidal effects against Dolichoderus affinis (Hymenoptera: Formicidae) was made and result is presented showing higher fumigant toxicity of C. medica and C. limon oil against the ant sp. With the increasing awareness for using safe insecticidal products among consumers, the citrus EOs with their attracting terpene compounds having good insecticidal potency bear all attributes to be used as commercial green pesticides in coming days both in indoor and outdoor management of insect pests.

Keywords: essential oils, limonene, Dolichoderus, Citrus medica

1. Introduction

The genus Citrus has tremendous industrial value all over the globe not only for its nutritive juicy high valued fruits but also for the essential oils present in its different vegetative parts. Thus, both the Citrus fruits and citrus essential oils bear potential to generate livelihood & to boost the country's economy. Citrus essential oils (CEOs) with diverse biologically active compounds of terpene groups with pleasant aroma have already achieved significant positions in flavor, food, cosmetic industries. At the same time, because of their antimicrobial activities as well as anticancer, antioxidant, anti-inflammatory, metabolic disorder alleviating activities etc. these oils and their compounds have been getting importance in pharmaceutical and medical sectors for the last few decades [1]. A good number of studies also reported insecticidal potential of citrus EOs extracted from different citrus sp. and their constituents at different times, a few of which are commercialized to be used by the consumers against insect pests. There are 33 recorded species of citrus worldwide (ThePlantList.org) with many recorded and unexplored varieties present in different parts of the world. The essential oil profile of different citrus species varies although some of the constituent compounds are common but present in different amounts in the total bulk oil. Even the oil profile of different vegetative parts of a single citrus species are not identical. Understanding of essential oil profile of diverse citrus species grown in wild, semi wild and cultivated state across the globe at different seasons is the much-needed task as the quality of the oil, oil yield percentage, consistency of the constituents even varies with seasonal changes, geographical location, harvesting time of the plant parts, soil type etc. however from the existing GC–MS profile of different *Citrus* sp. reported at different times and from different places, it is apparent that two -three dominant compounds are mostly present in most of the Citrus species. The literature revealed that the Citrus EO comprises more than 200 compounds of which 85–99% are volatile and 1–15% nonvolatile compounds. The volatile compounds comprise mostly monoterpenes (predominant limonene), some sesquiterpenes and their oxygenated derivatives [2].

Pest control sector is dominated by synthetic pesticidal products for many decades. At recent times with increasing concern to ecofriendly product, plant essential oils are getting renewed interest as they are not only effective but also comparatively safe and environment friendly in comparison to synthetic counterparts. Essential oils are part of natural plant defense system and many of them are proved effective and some are exploited for integrated management practices of pest and pathogens. As some citrus species are naturally resistant to certain group of pests and or pathogens, it is assumed that certain bioactive compounds may present in the essential oil part of those citrus species. It is already established that citrus essential oils of different citrus species are effective against wide range of pest and pathogens. It is also important to have an insight about the interaction of citrus constituents against its own insect pest and pathogen complex to be used as insecticidal, repellent and bactericidal etc. A few papers highlighted beneficial effects of using citrus essential oil against its own pest and pathogen complex. The added advantage of considering CEO as insecticidal and insect repellent is that the plant is edible therefore safe for residual contamination or toxicity to consumer. At the same time the pleasant aroma offers consumer acceptance.

2. Citrus EO against insect sp

CEO and extracts have been tried against a wide range of insect pests for assessing their insecticidal as well as repellent properties. In some parts of the world citrus plants have been traditionally used to ward off a insect pests. Some recent reports especially of the last two decades of the insecticidal and repellent effects of different citrus sp. are presented below. Most of the works were carried out on dipteran, lepidopteran, hemipteran and coleopteran insect pests.

Topical toxicity of the essential oil of *Citrus hystrix* with LD50 of 26.748 μ L/g and antifeedant properties leading to severe growth inhibition has been reported against tobacco armyworm *Spodoptera litura* [3]. The fumigant toxicity and repellent effect of the n-hexane extract of the plant leaf was documented against stored grain pest *Lasioderma serricorne* [4]. Fumigant toxicity of peel oils of lime, orange, mandarin, tangerine, grapefruit and lemon were reported against three store grain pest species *Callosobruchus maculatus*, *Sitophilus zeamais and Dermestes maculatus* [5].

The peel essential oil of the plant is reported to possess repellent effect against *Callosobruchus maculatus* [6], *Aedes aegypti* and *Anopheles minimus* [7]. Similarly, the insecticidal and repellent activity of *Citrus reticulata*, *Citrus limon* and *Citrus aurantium* peel oils was demonstrated against *Callosobruchus maculatus* [8]. Insecticidal activity of *Citrus limon* and *Citrus sinensis* against vine mealybug, *Planococcus ficus* [9]. The larvicidal and adulticidal effects of *Citrus limon* and

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Citrus sinensis are mentioned against *Attagenus fasciatus* and *Lasioderma serricorne* [10]. The seed and peel extracts of *Citrus limon* L. was reported to have the highest larvicidal toxicity (LC50 values of 395.59 ppm for seed; 468.69 ppm for peel) after 24 hours over EOs of *Citrus grandis, Citrus sinensis, Citrus paradise, Citrus reticulate* [11]. Essential oil of *Citrus reticulata* and *Citrus sinensis* was reported effective against the fourth instar larvae and adults of *Tribolium castaneum* with higher potency of *Citrus reticulata* [12].

The seed EOs of *Citrus reticulate* var. kinnow, *Citrus reticulate* var. freutrall, *Citrus sinensis* and *Citrus jambhiri* was tested against *Tribolium castaneum* with promising efficacy in terms of LC50 for *Citrus jambhiri* followed by Citrus reticulate and *Citrus sinensis* [13]. Similarly Oboh et al. [14] recorded insecticidal efficacy of *C. sinensis* peel essential oil against Callosobruchus mamulatus, *Tribolium confusum*, *Sitophilus oryzae* with LC50 value of 21.8, 38.9, 60 µl/l.

Comparative evaluation of toxicity of EOs of C. limon, C. aurantifolia, C.sinensis in filter paper impregnation method showed highest toxicity of C. limon (95% mortality) followed by C. aurantifolia (92.5%) and C. sinensis (82.5%) against carpenter ant Camponotus nearcticus [15]. But, Guerra et al. [16] comparatively lower topical toxicity (15% mortality) of C. limon EO against Camponotus pennsylvanicus among the eleven different EOs tested. Essential oils and or extracts of C. maxima or C.grandis have been reported effective against different mosquito species. In our earlier studies we recorded differential biocidal activities of essential oil extracted from peel and leaf part of Citrus grandis grown in Assam against different developmental stages of Aedes aegypti and Culex quinquefasciatus [17, 18]. EO extracted from leaves was found more effective against egg stage while oil from peel was recorded more effective against larval and adult stages of A. aegypti [18]. The leaf and peel oil of the plant was recorded highly effective against egg and larval stage with LC50 value of below 50 ppm but did not found much effective against adult stage of *Culex* quinquefasciatus although having repellent properties with good protection time [19]. In a recent study we observed synergistic larvicidal response of Citrus grandis leaf oil with Allium sativum bulb oil against C. quinquefasciatus [19]. Manorenjitha et al. [20] tested hexane, ethyl acetate, methanol, water and essential oil extract of C. grandis peel extract for evaluating oviposition deterrent and repellent properties on *Aedes aegypti*. They observed promising oviposition deterrent activity of ethyl acetate fraction (10 ppm concentration) in breeding plates kept within mosquito cage and effective repellency (94.7%) of 20% essential oil fraction of the peel while offering animal bait in modified tunnel test. A study for toxicity assessment on worker termites Odontotermes feae, essential oils of Citrus grandis with LC50 value of 273.36 ppm was found to show maximum toxicity out of Citrus paradisi, Cassia fistula, Citrus grandis EOs [21].

The peel essential oils of *Citrus aurantifolia* has been reported as insecticidal, repellent, and larvicidal against *Aedes aegypti* [22]. In our previous study, we observed the ovicidal, larvicidal and adulticidal effects of leaf and peel essential oil of *Citrus aurantifolia* against *Aedes aegypti* [23].

Promising fumigant toxicity of the peel EO of *Citrus aurantium* and *Citrus sinesis* from the north east Brazil with LC50 value of 5.80 μ L/L of air and 3.80 μ L/L of air respectively and oviposition deterrent activity at 3.5 and 7.0 μ L/L of air against the whitefly *Bemisia tabaci* [24]. The same oil was reported effective against the larval and adult stages of tomato leafminer *Tuta asoluta* (Lepidoptera: Gelechiidae) [25]. The insecticidal activity of *Citrus aurantium* EO against adult housefly *Musca domestica* was reported next to the activity of EO of *Citrus sinensis* [26]. *Citrus aurantium* leaf EO was found as effective fumigant against sawtoothed grain beetle *Oryzaephilus surinamensis*, cigarette beetle *Lasioderma serricorne* and rice weevil *Sitophilus oryzae* with LC50 value of 64.94, 202.49 and 364.25 μ L/L of air

respectively [27]. Similarly, Bnina et al. [28] noted fumigant toxicity of peel, leaves and flowers essential oil of Citrus aurantium against four stored grain pests namely Tribolium castaneum, Liposcelis bostrychophila, Sitophilus granarium, Cryptolestes ferrugineus with LC50 value of 64.78%, 23.11%, 101.50% and 20.62% respectively. They also noted repellent property of the oil against these pests. Yazdgerdian et al. [29] tested contact and residual toxicity of eleven essential oils including Citrus aurantium, Citrus sinensis, Citrus limon against wooly beech aphid, Phyllaphis fagi (Hemiptera: Aphididae) and rice weevil, Sitophilus oryzae (Coleoptera: Curculionidae) and recorded highest residual toxicity (40%) of C. aurantium among the citrus sp. tested against the targeted species. Similarly, without affecting the seed viability of stored cowpea and consumer acceptability, peel EOs of *Citrus* nobilis and Citrus medica was reported to show significant reduction of egg laying, egg hatching, and adult emergence percentage of pulse beetle *Callosobruchus maculatus*. Both the oils showed dose dependent repellency with higher effect for EO of *C. nobilis* [30]. Like that of EOs, the nonpolar petroleum ether extract of ripe fresh fruit of C. aurantium was reported effective against adults of olive fruit fly Bactrocera oleae (Diptera: Tephritidae) in petri dish residual exposure test [31]. The same solvent extract also reported to have good toxicity against medfly Ceratitis *capitata* adults (LC50 value of 70.6 and 147.1 lg/cm² for male and female respectively at 96 h in Petri dish residual bioassay) [32].

Moravvej et al. [33] tested fumigant toxicity of EOs from four citrus species namely *C. paradisi*, *C. limonium*, *C. sinensis* and *C. aurantium* among which *C. paradisi* was the most effective with LC50 value of 125 μ l/L and *C.sinensis* is the least effective with LC50 value of 269 μ l/L against *Callosobruchus maculatus*. However, fumigant toxicity of *C.sinensis* was reported against *Solenopsis invicta* (Hymenoptera: Formicidae) with 100% mortality at 3 mg/tube after 24 hrs [34]. Ethanolic extract of the same plant was found effective against *Anopheles stephensi*, *Aedes aegypti* and *Culex quinquefasciatus* with larval and adulticidal LC50 value of below 500 ppm along with more than 50% repellency at 150 ppm concentration till 180 min [35].

Ezeonu et al. [36] reported the insecticidal properties of the volatile peel extracts of *Citrus sinensis* and *Citrus aurantifolia* against mosquito, cockroach and housefly and recorded higher insecticidal potency of the peel extract of *Citrus sinensis* with maximum fumigant effect (85% at 60 min) against cockroach.

Zewde and Jembere [37] evaluated the solvent extract and essential oil of *Citrus sinensis* against *Zabrotes subfasciatus* (Coleoptera: Bruchidae) for their repellent, fumigant and protectant properties and recorded no progeny emergence on application of oil at low dose (30 mg of EO), prominent percent mortality at high dose (750 mg of essential oil killed 67% of Z. subfaciatus after 96 hours). Fumigant toxicity of *C. sinensis* EO was also reported effective against second instar larvae of *Musca domestica* (Diptera: Muscidae) with LC50 values of 71.2 μ l/L as well as percent inhibition of pupae with 46.4% at 40 μ l/L exposure concentration [38].

Orange oil extract was also recorded effective against the subterranean termite *Coptotermes formosanus* (Isoptera: Rhinotermitidae). Application of the oil extract at 5 ppm concentration resulted in 96% and 68% mortality respectively in closed container. At the same time termites did not show tunneling behavior on the 0.2 & 0.4% oil treated soil [39].

Majeed et al. [40] reported the insecticidal activity of the acetone, ethanol and aqueous extracts of seeds, leaves and fruit peels and leaves of *Citrus aurantium* and *Citrus sinensis* and two other plants against mealy bug *Drosicha mangiferae* (Hemiptera; Pseudococcidae) of which ethanol extracts of *C. sinensis* seeds and *C. aurantium* leaves were suggested as considerably toxic against the said insect.

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C. medica peel essential oil was reported as more effective in filter paper fumigation method against stored grain pest *Tribolium castaneum* (Coleoptera) with LC50 value of 29.5 mg/L air than the leaf EO [41].

Abdel-Kawy et al. [42] showed *Citrus trifoliata* essential oil loaded nanocubosome significantly enhanced insecticidal property of the essential oil against the second instar larvae of *Spodoptera littoralis*.

While working as biocidal and repellents, plant products including EOs and constituent terpene compounds are reported to act on cholinergic system [43], voltage – gated sodium channel of the nerve membrane, glutamate-gated chloride channel [44], GABA-system [45], Octopaminergic system [46], mitochondrial system [47], endocrine system disrupting the endocrinological balance and respiratory system of insect body. However, not much studies yet conducted on detailed study on the mode of action of EOs and their constituent compounds.

3. GC-MS profile of citrus EO

With the development of GC–MS technique, profiling of essential oil became easier. The composition of different citrus species from different parts of the world have been reported utilizing this technique. Most of the profiling results although detected average 20–50 numbers of compounds, a few compounds mostly occupy the major share of the bulk oil. The dominating compounds in most citrus species is limonene. In some species like *C. hystrix* the major compound is β -citronellal. The other common constituent compounds observe to be present in many GC–MS results of citrus species are linalool, pinene, β - caryophyllene, β -myrcene, terpinene, citral etc. Some of the reported constituent profiles of citrus species are mentioned below.

3.1 Citrus hystrix

From the leaf essential oil of *C. hystrix* in Malaysia, 29 compounds were reported, out of which beta-citronellal was the major compound (66.85%) [3]. The other compounds present in more than 1% total compositions were β -citronellol (6.59%), linalool (3.90%), 5,9-dimethyl-1-decanol (4.96%), methyl citronellate (1.90%), geranyl acetate (1.80%), citronellol (1.76%), 3-undecanol (1.04%). The same compound (beta-citronellal) with 86.43% amount was reported in another study from Indonesia along with 11.48% citronellol and 1.65% β -linalool [4].

3.2 Citrus maxima (synonym Citrus grandis)

In the leaf EO of *C. maxima*, 42 constituent compounds were reported to be present with citronellol as the major compound (28.26%) of the essential oil. The compounds comprised more than 1% of the oil compositions were β - caryophyllene (16.89%), –spathulenol (9.32%), α -caryophyllene (2.48%), γ -cardinol (3.16%), α -cardinol (2.51%), 2n-hexylcyclopentanone (2.22%), caryophyllene oxide (1.03%). Again, 34 compounds were identified from the peel essential oil of the same plant with limonene comprising bulk of the oil constituents (89.04%) [48]. The major compounds identified in more than 1% of the total composition were β -pinene (2.25%), β -myrcene (2.06%), β - copaene (1.76%). Compounds comprising more than 0.3% amount but below 1% were linalool, β -phellandrene, α -pinene, terpinene-4-ol. Earlier 35 compounds with limonene as the major constituent compound (93.2%) was reported from the fruit peel oil [49]. Myrcene comprised 2.9%

and other six compounds namely α -pinene, octyl acetate, germacrene -D, linalool, decanal, geranial comprised above 0.2% but below 1% of the total composition.

3.3 Citrus aurantium

Phytochemical profiling of essential oil of *C. aurantium* showed the presence of 25 compounds with limonene occupying 87.52% followed by linalool (3.365%) and β-myrcene (1.628%) as dominant compounds [25]. From Tunisian *Citrus aurantium*, limonene percentage in leaves, flowers and peel EOs were 6.52, 5.03, 73.6% respectively, linalool occupied 37.24, 41.82, 4.8% respectively, linaly acetate occupied 7.87, 13.75, 1.6% respectively and neral share was almost of similar (3.40, 4.80, 3.26% respectively) percentage. β-pinene composition for leaves and flowers were 9.68 and 9.21% respectively and α-thujene present below 0.5% in peel EO of the plant [28].

3.4 Citrus aurantifolia

In our recent studies 31 compounds from the leaf oil and 26 compounds from the peel essential oil was recorded from GC–MS analysis of EO of *Citrus aurantifolia* in India. Citral and limonene were noted as the major constituent compound of leaf oil and limonene and palatinol-1C as the major constituent of peel essential oil of the plant [19, 23]. The EO of the plant from Italy reported to comprise limonene (53.8%), γ -terpinene (16.5%), β -pinene (12.6%), β -Bisabolene (1.33%), Geranyl acetate (1.06%), Neryl acetate (1.12%), Geranial (1.84%), sabinene (1.74%), and α -Pinene (1.97%) [50]. Phytochemical analysis of leaf and peel EOs of the plant from Brazil [51] showed limonene as the dominant compound in both leaf (32.7%) and peel (77.5%) part. Other prominent compounds in the leaf EO were linalool (20.1%), citronellal (14.5%), citronellol (14.2%), trans- β -Ocimene (2.7%), geranial (2.6%), neral (2.1%), trans- β -Caryophyllene (2%), myrcene (1.4%) etc. The other major compounds of leaf EO were myrcene (4.4%), linalool (3.5%), citronellal (3.2%), citronellol (2%), β -Bisabolene (1.5%) etc.

3.5 Citrus sinensis

Phytochemical analysis of peel essential oil from three varieties of *C. sinensis* from Kenya showed presence of 56 components in Salaustiana variety, 73 in Valencia and 72 in Wshington varirty. Limonene occupied more than 90% in all the essential oil (Salustiana 94.6%, Valencia 92.5% and Washington 90.5%); alpha terpinene occupied 1.5% in Valencia and Washington and 1.7% in Salustiana [52]. Limonene with 90% share and β -myrcene, γ -terpine, linalool with 1.88%, 1.21% and 0.88% share out of 32 identified compounds in peel EO of *C. sinensis* has been reported from China [53]. GC–MS analysis of the plant EO from Argentina showed limonene (92.47%), linalool (1.43%), and β -myrcene (0.88%) as the major constituent compounds along with terpineol (0.28%) in lesser amount [26]. Almost similar constituents were identified from EO of *C. sinensis* with D-limonene (65.28–80.18%), Linalool (0.32–2.20%) and β - pinene (1.71 5.58%) as major part in another study [34].

3.6 Citrus nobilis

Phytochemical study on constituents from the peel of *C. nobilis* from China showed D- limonene (12,601 μ g/g) as the major constituent compound followed

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by β-myrcene (1600 µg/g), β-pinene (82.64 µg/g), p-mentha-1,8-dien-3-one (41.33 µg/g), α-pinene (25.41 µg/g), geranial (21.32 µg/g), sabinene (21.18 µg/g), E-β-ocimene (14.97 µg/g), linalool (10.38 µg/g), α-terpineol (6.36 µg/g). Other compounds are present in lower amounts (below 5 µg/g) [54]. In another study from Sri Lanka, 37 compounds were reported from peel part of which D-limonene (45%) was the major one followed by cyclopentane-2-methyl-1-methylene-3-(1-methylethenyl) (3.94%), p-mentha-4,8-diene (3.73%), α-terpinolene (3.03%), methyl-2-(methylamino) benzoate (2.4%), α-fernesene (1.1%). Other compounds were present in below 1% [30].

3.7 Citrus limon

Phytochemical analysis of citrus leaf EO from Iran showed presence of 27 compounds of which the major compound was linalool (30.62%). The other compounds present in significant amount were geraniol (15.91%), α -terpineol (14.52%), linalyl acetate (13.76%), geranyl acetate (6.75%), B-pinene (4.51%), neryl acetate (4.24%), p-Cymene (1.86%), and limonene (1.13%) [55]. Chemical composition of EO of *C. limon* grown in Iraq [56] showed presence of 24 compounds with limonene as principal compound with 29.52% share. Other major compounds were β -Pinene (23.89%), α -Pinene (2.25%), Myrcene (1.31%), (Z)- β -Ocimene (2.09%), Linalool (1.41%), (R)-Citronellal (15.10%), α -Citronellal (3.57%), (+)- α -Terpineol (1.57%), Neral (Z-Citral) (1.19%), Geranial (E-Citral) (1.73%), Thymol (9.79%), Citronellyl acetate (1.87%), Caryophyllene (1.36%), Phytol (1.36%). The analysis of the essential oil of *Citrus limon* from North-East India reported presence of 43 constituent compounds of which limonene (55.40%), neral (10.39%) trans-verbenol (6.43%) and decanal (3.25%) were the major constituent compounds [57].

3.8 Citrus paradisi

Phytochemical analysis result of *C. paradisi* peel EO from Turkey demonstrated presence of 25 constituent compounds of which limonene occupied the highest percentage (88.6%) of the oil. The other major compounds were α -terpinene (1%), and β -pinene (1.2%) [58].

From Nigeria, fifteen phytochemical constituents of the plant oil were reported. Among the compound limonene (94.2%) occupied the major share [59].

3.9 Citrus medica

A total of 19 constituent compounds were identified from leaf essential oil of *Citrus medica* from Bangladesh of which erucylamide (28.43%), limonene (18.36%), citral (12.95%), Mehp (8.96%), 2,6-octadien-10l,3,7-dimethyl-acetate, (Z) (5.23%) were the major compounds. From peel essential oil 43 compounds were reported out of which isolimonene (39.37%), citral (23.12%), limonene (21.78%) were the major constituents. Three other compounds namely β -myrcene, neryl acetate and neryl alcohol were reported to present at around 2% each in total composition and remaining compounds were present in traces amount [60]. In a study carried out by Li et al. [61], all total 28 compounds were reported to present in the fruit essential oil of *Citrus medica* of which limonene (45.36%), γ -terpinene (21.23%), dodecanoic acid (7.52%) were documented as major constituent compounds. Compounds like β -bisabolene, tetradecanoic acid, α -terpineol, terpinene-4-ol, hexadecenoic acid, α -bergamotene, α -pinene, β -pinene comprised between 5–1% range in total composition. In another study fruit peel EO of the plant was reported to comprise limonene (38.7%), γ - terpinene (28%) and o-cymene (15.2%) as major compounds [41].

4. Citrus EO compounds against insect sp

Essential oil composition of different citrus sp. across the globe although may vary but some of the compounds are observed as common in most of the oil profile. The most dominating and commonly present compound is limonene. Other common compounds are citronellal, citronellol, linalool, pinene, myrcene, ocimene, terpinene, caryophyllene etc. The bioactivity of EOs is often related to the activity of major compounds present in the crude oil and some of the studies have already established this fact. Individual assessment for insecticidal property of these common constituent compounds have been performed by different researchers and some of them were found active against insect pest. Limonene and other *Citrus limon*oids are reported as insect repellents, feeding deterrents, growth disruptors, and reproduction inhibitors against a wide range of pest complexes. Insecticidal activity of limonene was reported effective against Tuta asoluta (Lepidoptera: Gelechiidae) [25]. Yoon et al. [62] revealed repellent property of different citrus oil and its major compound limonene against different species of cockroaches like Blatella germanica, Periplaneta americana and Periplaneta fuliginosa. However, Karr and Coats [63] did not get significant insecticidal activity of d-limonene against Blattella germanica, Musca domestica, Sitophilus oryzae and Diabrotica virgifera virgifera. In contrast they reported enhanced growth of nymph of Blatella germanica after oral administration of d-limonene.

In another study against cat flea species *Ctenocphalides felis* (Siphonaptera: Pulicidae), d-limonene (LD 50 against larvae, adults 226, 160 μ g/cm² respectively) and d-limonene with piperonyl butoxide (PB) (LD 50 against larvae, adults 157, 49 μ g/cm² respectively) were reported effective against all the life stages except the pupal stage of the flea species [64]. Fumigant toxicity of d-limonene, α - terpineol etc. also reported against honey bee *Apis mellifera* and tracheal mite parasite species *Acarapis woodi* [65].

Fouad and da Camara [66] extracted the essential oil from *Citrus aurantiifolia* and *Citrus reticulate* and analyzed the phytochemical constituents using GC–MS and found limonene as the major constituent compound, 38.9% of the *C. auran-tiifolia* oil and 80.2% of the *C. reticulata* oil. They analyzed the enantiomers of limonene against the said insect. They found that *Citrus reticulate* was more toxic than *Citrus aurantifolia* towards the said insects. (R)-limonene was shown to have greater toxicity against *S. zeamais* than the (S)-limonene as found in the ingestion bio assay. Repellent bioassay showed (S)-limonene more susceptible to *S. zeamais* than (R)-limonene.

After identifying limonene as major compound in the EOs of *Citrus aurantiifolia* (38.9%) and *Citrus reticulate* (80.2%) Fouad and da Camara [66] tested enantiomers of limonene against *Sitophilus zeamais* and recorded greater toxicity of (R)-limonene than the (S)-limonene in the ingestion bio assay. But in the repellency test they found more susceptibility of *S. zeamais* towards (S)-limonene than (R)-limonene.

In a recent study, Sowler et al. [67] comparatively evaluated the effect of laboratory grade limonene and a commercial limonene-based insecticide against *Haematobia irritans irritans* in terms of deterrence, mortality, and reproduction. They showed that the egg viability was decreased in both the treatment, however, commercial limonene that caused loss of viability at 5.8% concentration was ovicidal in case of laboratory grade limonene. However, in terms of knockdown effect commercial limonene was better. Interestingly, at a concentration of less than 0.1%, both the commercial and laboratory grade limonene were acted as attractant.

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Giatropoulos et al. [68] tested essential oil of Citrus sinensis, Citrus limon, and *Citrus paradise* and their constituents and recorded γ - terpinene as the most toxic compound against Aedes albopictus larvae. They also reported that the constituent compound tested for repellency were better mosquito repellent than the parent essential oil. Similarly, Luo et al. [41] analyzed composition of leaf and peel essential oil of *C. medica* and tested both crude oil and major compounds viz. limonene, terpinene, o-cymene, β -caryophyllene against *Tribolium castaneum* and recorded γ -terpinene as the most effective insecticidal compound having LC50 value of 4.1 mg/l air and β - caryophyllene as the effective repellent compound. Limonene was reported to have almost similar fumigant toxicity like that of crude EO of C. aurantium against Tribolium castaneum, Sitophilus granarium and Cryptolestes ferrugineus [28]. In our earlier studies we found higher toxicity of citral, that is the major compound of the essential oil of *Citrus aurantifolia* than the crude oil as mosquito larvicidal, ovicidal and adulticidal against Aedes aegypti [23]. Plata-Reuda et al. [69] reported the insecticidal activity of citral and geranyl acetate against peanut beetle Ulomoides dermestoides. These compounds affected the survivorship, locomotor activity and reduced the respiration rate of the said species.

Nootketone and carvacrol, a phytochemical constituent present in essential oil of Citrus [70] acts as insecticidal compound against *Aedes aegypti* [71]. Pajaro-Castro et al. [72] recorded the neurotoxic effects of linalool and β -pinene on *Tribolium castaneum*. They observed that at low concentration both the compounds were attractant towards the insect and at higher concentration the compounds were repellent. Individual treatment of limonene and linalool was found to have fumigant toxicity against two ant species namely *Acromyrmex balzani* and *Atta sexdens* with LC50 values of 5.72 µl/L, 5.38 µl/L and 2.40 µl/L, 5.34 µl/L respectively [73]. Similarly individual treatment of β -caryophyllene is reported to have good contact toxicity against these two ant sp. but with low fumigant toxicity [74]. Fumigant toxicity of limonene, linalool and β - pinene were also reported effective against fire ant *Solenopsis invicta* (Hymenoptera: Formicidae) [34].

Linalool, α-terpinene was reported to show 100% fumigant toxicity against adult rice weevil *S. oryzae* at 3.9 mg/L [75].

Muller et al. [76] recorded 85.4%, 71.1%, and 29% repellency of the candles prepared with 5% geraniol, 5% linalool and 5% citronella against mosquitoes on human landing bioassay. They observed similar repellency against sand flies too. 78% repellency of *Culex pipiens pallens* was reported after using 30% citronellal [77]. Progeny deterrent, antifeedant, egg hatching inhibition activity was documented after application of 5–10 µl citronellol against *Callosobruchus analis* [78]. Compounds like limonoids act as synergists to enhance the activities of other biological and or synthetic insecticides [79]. We recorded medium larvicidal and adulticidal potential of limonene against *Aedes aegypti*, but found higher toxicity when combined with diallydisulphide and carvone respectively [80]. So, it is not always the individual compound that act as the most active but appropriate combinations of compounds having synergistic effects would be more fruitful as insecticidal against insect pests.

5. Citrus of North East India

North East India is enriched with Citrus species having documented 23 species and 68 varieties out of the 27 species of Citrus found in India [81, 82]. It is established that some of the citrus species are endemic and some are in endangered status [83]. According to Hore and Barua [84], there are eight citrus species indigenous to this region scattered in the form of semi-wild, wild state and some in cultivated state. Some of the species are naturally tolerant to viral and bacterial diseases and also for drought, cold and rainfall. For instance, *Citrus limon* is reported to resist scab, canker and gummosis, *C. indica* resistant to greening disease [84]. Due to diverse ecogeographical conditions the Citrus species and varieties of this region may bear specific traits including its aroma and essential oil constituents which needs to be investigated. However citrus crop and citrus essential oil-based industry is not yet flourishing in this part of the country. The information regarding Citrus essential oils extracted from the Citrus species grown in this particular area is relatively scanty. As the Citrus plants are rich in secondary metabolites to naturally defend an array of pathogens and pest complexes, it is expected that some of the key compounds for controlling insect pest may lie within the secondary metabolite compounds especially in the diverse aromatic essential oil part of the plants at least in the resistant citrus species.

Here we have attempted to evaluate insecticidal properties of essential oil extracted from the fruit peel of four citrus species namely *Citrus limon*, *Citrus max*ima, Citrus paradisi and Citrus medica grown in North Eastern part of India against one of the household ants Dolichoderus affinis (Hymenoptera: Formicidae). Fruits of Citrus limon, Citrus paradisi and Citrus medica were collected from Udalguri district, Assam (26.7210° N, 91.9906° E) and Citrus maxima from a daily market at Guwahati, Assam (26.1445°N, 91.736°E), in September, 2020. Essential oils from fresh peels were extracted by hydro-distillation using Clevenger's apparatus. After 6 hours, essential oils were collected, anhydrous sodium sulfate was added to absorb traces of moisture and were stored at 4°C till its use. The worker ants of Dolichoderus affinis from naturally existed colony located in the wooden frame of house wall were considered for the assessment. Fumigant toxicity of these oils were assessed following the method described by Hu et al. [34]. Six different concentrations viz. $0.25 \,\mu$ L, $0.5 \,\mu$ L, $1 \,\mu$ L, $2.5 \,\mu$ L, $5 \,\mu$ L and $7.5 \,\mu$ L. of each EO was individually loaded in 1.5 ml centrifuged tubes, evaporation of EOs was allowed by making five small holes of 1-1.2 mm diameter and placed into 500 ml properly cleaned borosilicate conical flask. Twenty worker ants were taken per flask in a replication and the flask was covered by aluminum foil and bound tightly by rubber band to prevent the loss of volatile compounds. For each concentration three replications were made. Equal number of controls set without oil were placed against each treatment. The environmental temperature range was 21–30°C and relative humidity range 56–99 during the experimental period. The Percent mortality [percent mortality = (Total no. of dead ants/Total no. of treated ants) × 100] data was recorded after 12 h and 24 h of treatment. The ants were considered to be dead if touched with a needle but did not show any movement. Based on the results sublethal concentration was determined using probit analysis with the help of SPSS and Minitab software. As shown in the figure, after treatment with *C. limon*, maximum 71.66% mortality was recorded at 12 h and 91.66% was recorded after 24 h at 7.5 µL treatment. The calculated LC50 for the oil was 2.66 μ l / 500 air volume. For the EO of *C.paradisi*, maximum 15% mortality was recorded at 12 h and 66.66% mortality was recorded after 24 h of treatment. LC50 for the oil at 24 h was 7.32 μ l / 500 air volume. For the EO of *C. maxima*, not more than 10% mortality was recorded even after 24 h at the highest dose applied and LC50 could not be computed for the oil. While for the EO of *C. medica*, maximum 56.6% mortality was recorded at 5 µl concentration at 12 h and maximum 88.33% mortality was recorded after 24 h at 7.5 µl. LC50 for C. *medica* at 24 h was recorded as 2.09 µl/500 air volume (**Figure 1**, **Table 1**). Highest toxic effect was recorded for C. medica followed by C. limon. Earlier Adusei-Mensah et al. [15] evaluated insecticidal properties of three citrus species viz. Citrus aurantifolia, Citrus sinensis and Citrus limon against Camponotus nearcticus (Formicidae) and recorded highest performance from C. limon with 95% mortality.

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Figure 1.

Relation between concentration of EOs and respective percent mortality.

Essential oils	Time	LC50 value µl/500 ml air [—]	95% confidence level		Regression equation	Chi-square
			Lower limit	Upper limit		value
Citrus paradisi	24 h	7.32	1.143	1.902	Y = 3.67118 + 1.53688X	31.750
Citrus limon	24 h	2.66	1.739	2.486	Y = 4.03677 + 2.26321X	49.452
Citrus medica	24 h	2.09	1.609	2.266	Y = 4.37759 + 1.94055X	52.747

Table 1.

LC50 values of the individual citrus oils against Dolichoderus affinis at 24 h.

But Guerra et al. [16] recorded only 15% mortality of *Camponotus pennsylvanicus* (Hymenoptera: Formicidae) on topical application of *C. limon*, the efficacy of which was comparatively lower than other eight EOs tested against the ant species. Not much studies on insecticidal activities of citrus EO against ants have been found to be reported. The findings showed the prospect of using *C. medica* and *C. limon* oil for controlling household ants.

6. Conclusion

With the increasing awareness of consumers for ecofriendly products and at the same time increasing resistance of insect pests against insecticides, the demand for novel, safe and effective products is increasing. As discussed above, the existing literature revealed presence of a good number of terpene compounds in different *Citrus* species which are present in different ratios although in most cases limonene is the predominant constituent. Both the crude oil as well as individual compounds possess good insecticidal and repellent properties against diverse insect pests, both indoor and outdoor. Our study also showed promising potential against *Dolichoderus affinis* while using four Citrus essential oils with higher efficacy of *Citrus medica* and *Citrus limon* essential oils. It is expected that in near future Citrus plant essential oils with their pleasant aroma and array

of chemical compounds shall take leading space in development of insecticidal and repellent products to be used in both indoor and outdoor pest management practices against insect pests.

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Chapter 11

The Orange Peel: An Outstanding Source of Chemical Resources

Gianfranco Fontana

Abstract

Citrus sinensis (L.) Osbeck is a very common cultivar belonging to the *Rutaceae* family. It is largely diffused in several areas of the world characterized by mild to warm climate conditions. Its abundant worldwide production (up to 10^7 Tons. per year) and consumption both as the edible part of the fruit and as several types of derivative products imply the production of a huge amount of waste, such as the fruit pomace. Several ways of recycling this material have been developed in recent years: employment as fertilizer, fodder ingredient, and even cloth material. However, the chemical added value of *Citrus sinensis* peel has been underestimated despite the diversified and significant content of useful chemicals, such as polyphenols, polymethoxylated phenols, glycosylated flavonoids, volatile and non-volatile terpenoids, pectins, enzymes, etc. This work aims to highlight the outstanding chemical potential of *Citrus sinensis* peel.

Keywords: biological activity, *Citrus sinensis*, essential oil, flavonoids, orange peels, polymethoxyphenols

1. Introduction

Citrus sinensis (CS) (L.) Osbeck is a perennial species growing in warm climate areas of the world and largely employed as food in form of fresh fruit, with a global production of ca. 6.7X10⁷ tons. per year (TPY) in 2016 [1], or as a processed derivative (ca. 1.85x10⁷ TPY) such as juice, marmalade, flavor, fragrance and coloring additive, pectin.

CS is an evergreen tree, 3 to 9 mts. high with sparingly barbed branches, alternate leaves with toothed blades differently shaped, oval or elliptical, connected to the stem by winged-petioles. Axillary flowers are present singly or in whorls of 6 and possess 5 white petals and up to about 25 yellow colored stamens. The pericarp of CS has a spherical or oval shape of 6–10 cm diameter with the color changing from green to yellow-orange during the ripening; the endocarp containing juice sac glands is enclosed within a wrinkled epicarp or exocarp or flavedo containing a great number of essential oil glands protected by a waxy epidermis. Below the flavedo is the albedo, also called mesocarp, a white filamentary tissue composed of tubular-like cells.

The principal industrial application of CS is the production of frozen concentrated juice. The procedure of juice extraction eventually accompanied by the extraction of the essential oil, implies the generation of a major "by-product" constituted by a pomace, mainly containing peels, accounting for up to around 60% w/w of the original fruit mass processed [2]. This huge amount of biomass does pose serious environmental concerns because of its high level of total organic carbon (TOC) and biological oxygen demand (BOD) that make disposal procedures rather complex and demanding from both the legal and industrial points of view. This is because there is an increasing trend to modify the way of approaching this problem by reconsidering the post-production orange pomace more like a by-product rather than a waste. In the last years, many producers have subjected this material to processings involving partial acidic fermentation, drying, and packaging to biologically and chemically stabilize the biomass before its application as animal feed in zootechnics, soil conditioners in agriculture, or the manufacturing of compost and biogas [2].

Beyond the standard workup of the *Citrus sinensis* peel (CSP) waste, new perspectives have been being opened in the context of the high chemical added value of the CSP [3–5] also by the complete knowledge of the rich metabolomics profile of this species. The use of CS peel has been proposed for a variety of purposes that include the production of antioxidant-enriched dietary supplements in veterinary [6], the preparation of human dietary supplements, and nutraceuticals such as citric acid [7] and flavonoids [8, 9]. The extract of CS peel is the source of a huge variety of phytochemicals and has been investigated on several applications including its chemotherapeutic and chemopreventive potential for several relevant human pathologies, such as cancer [10, 11] and obesity [12]. The extraction procedures vary in function of the main components that have to be obtained: from the simple cold pressing of pomace and the extraction with water to obtain highly hydroxylated compounds to the employment of mixtures organic protic solvent/water and finally low polar organic solvents such as Chloroform and Ethyl acetate to obtain polymethoxylated phenols (PMF, see below). New extraction technologies such as ultrasounds and microwaves may help to obtain better extraction yields.

In the following sections, the chemical structures and the biological effects of these compounds will be discussed.

2. The chemistry of Citrus sinensis peel

2.1 Essential oils

The essential oil (EO) is mainly obtained from the CS peel as a major by-product of the juice production process by a cold-pressing method that can provide the intact blend of compounds without losing the lighter, more volatile, components of the complex mixture that can be lost in the standard EO extraction procedure that is the hydrodistillation. The last one is mainly used in small scale applications, for example in research laboratories.

The chemical composition of CSP EO [13–15] is reported in **Table 1**. As it can be seen, the major component is D-Limonene, accompanied by several minor components belonging to the classes of monoterpene alkenes, oxygenated monoterpenes including alcohol aldehydes and esters, sesquiterpenes as well as linear alkanes and aldehydes. This rather complex blend accounts for the numerous deal of biological activities reported for the CSP EO [14–16], which include anthelmintic, anti-aflatoxigenic [17], antibacterial [18–20], anticarcinogenic, antifungal [21], antioxidant [17], anti-tumor [22], anxiolytic [23], food preservative [24], hepatocarcinogenesis suppressant, insecticidal and larvicidal [25], pain relief and relaxant [26]. It can be argued that the main effects are due to the presence of the major component Limonene that showed several bioactivities when tested as pure compound [27]. However, it is possible that synergistic effects due to the combination of Limonene with other minor components may be speculated and should have to be demonstrated.

Comp.	Comp. name	%	Compound.	Comp. name	%
1	Aromadendrene	0.01	21	β-Linalool	0.4–5.6
2	δ-Amorphene	0.05	22	β-Myrcene	1.3–3.3
3	D-Cadinene	0.01-0.03	23	Neral	0.1–1.3
4	δ-3-Carene	0.18	24	Neryl acetate	0.02
5	β-Citral	0.12-0.15	25	Nonanal	0-0.1
6	L-(+)-Citronellal	0.01–0.1	26	Nootkatone	0.01
7	Citronellyl acetate	0.01	27	<i>cis-</i> β-Ocimene	0.03-0.26
8	α-Copaene	0.04	28	Octanal	0.02–0.8
9	α-Cubebene	0.02-0.26	29	Perillaldehyde	0.03
10	β-Cubebene	0.03	30	α -Phellandrene	0.02–0.07
11	Decanal	0.04-0.4	31	α-Pinene	0.49-0.59
12	n-Dodecanal	0.06	32	(+)-Sabinene	0.2–1.0
13	β-Elemene	0.01-0.02	33	γ-Terpinene	0–1.21
14	Geranial	0–1.8	34	γ-Terpineol	0.04–008
15	Germacrene-D	0.02–0.08	35	α-Terpineol	0.07–0.42
16	β-Gurjurene	0.01	36	Terpinolene	0-0.08
17	Hexadecanol	0.04	37	α-Thujene	0.04
18	D-Limonene	Ca. 95			
19	L-Limonene	0.02			
20	trans-Limonene oxide	0.01			

Table 1.

Composition of C. sinensis essential oil obtained from peels.

2.2 Polyphenols

2.2.1 Flavanoids

Polyphenols extracted from the CS peel belongs to the general structural categories of flavanones (**Figure 1a**), flavones (**Figure 1b**), flavonols (**Figure 1b**), with and without sugar moieties attached to one or more of the hydroxyl groups [28]. It is worthy of particular mention the rarely occurring class of C-glycolflavones (**Figure 1b**, compounds **63–65**, **85**, **86**).

These compounds are produced *in vivo* from the biogenetic mixed pathway of the Acetate and Shikimate that implies the enantiospecific formation of the basic aromatic bicyclic framework of the flavanone, from which a huge number of flavonoids originate employing selective enzymatic hydroxylations, methylations, and glycosylation steps. As can be seen from the structures shown in **Figure 1**, most of the chemical entities found in the peel extract contain several methoxy fragments that decorate the carbon skeleton. This characteristic makes those molecules to get a rather apolar character that explains their presence in the hydrophobic environment of the waxy peel. On the contrary, compounds containing a major number of hydroxyl groups are less present in the peel and are instead more significantly concentrated in the juice of the pericarp. However, some glycosylated compounds are present in the peel. In these molecules, the aglicone bears a monosaccharide unit (mainly glucose) or a disaccharide, in most of the cases being Rutinose (91) – Rhamnosyl ($\alpha 1 \rightarrow 6$) glucose – or Neohesperidose (92)- Rhamnosyl ($\alpha 1 \rightarrow 2$) glucose (Figure 2).

The composition of the peel extracts described in the literature may slightly vary depending on the cultivar and the region of harvesting but some general points are



Cm	R_1	R_2	R ₃	R_4	R_5	R_6	Name
р.							
38	Н	Н	Н	Н	Me	ОН	Hesperetin
39	Н	Н	Rut	Н	Glu	Н	Narirutin-4'- glucoside
40	Н	Н	Rut	Н	Me	ОН	Hesperidin
41	Н	Н	Neohesp	Н	Me	ОН	Neohesperidin
42	Н	Н	Rut	Н	Н	Н	Narirutin
43	Η	Н	Rut	Н	Me	Н	Didymin
4 4	Η	Н	Glu-Glu	Н	Н	Н	Naringenin-7-0- bglucoside
45	M c	OMc	Ме	Н	Ме	Н	5,6,7,4'- tetramethoxyflavano ne
46	Н	OMe	Me	OMe	Me	OMe	5-hydroxy-6,7,8,3',4'- pentamethoxyflavano nc
47	Н	Н	Ncohesp	Н	Н	Н	Naringin



	R_1	R_2	R3	R4	R₅	R ₆	R7	Name
Cmp.								
48	Н	Н	Н	Н	Н	Н	Н	Apigenin
49	II	II	II	II	Me	11	II	Acacetin
50	Ме	OMe	Me	II	Ме	II	II	Tetra-0-
								methylscutellarein
51	Me	OMe	Me	Н	Me	OMe	Н	Sinensetin
52	Ме	OMe	Me	OMe	Me	Н	Н	Tangeretin
53	Me	OMe	Me	OMe	Me	0Me	II	Nobiletin
54	Me	OMe	Me	Н	Me	OMe	OMe	Hexa-O-
								methylquercetagetin
55	Ме	OMe	Me	OMe	Me	OMe	OMe	3',4',3,5,6,7,8-
								Heptamethoxyflavone
56	Н	Η	Н	Н	Me	Н	OH	Kaempferide
57	П	II	П	11	н	П	011	Kaempferol
58	П	II	П	I1	П	OH	II	Luteolin
59	Н	Η	Н	Н	Н	OH	OH	Quercetin
60	Me	OMe	Me	OMe	Me	OMe	OH	Natsudaidain
61	Me	OMe	Me	OMe	Me	Н	OH	3-hydroxy-5,6,7,8,4'-
								pentamethoxyflavone
62	Ме	OMe	Ме	I1	Ме	11	011	3-hydroxy-5,6,7,4'-
								tetramethoxyflavone
63	Н	Н	Н	C-	н	Н	Н	Vitexin
				Glu				
64	П	C-	11	C-	11	II	II	6,8-di-C-
		Glu		Glu				Glucosylapigenin
65	Н	C-	Н	C-	Me	OH	Н	6,8-di-C-
		Glu		Glu				Glucosyldiosmetin

66	11	11	Ме	11	П	OMe	0-	
67	II	II	Me	II	II	II	II	5,4'-dihydroxy-7-
68	II	II	II	11	II	OH	0- Chu	isoquercetin
69	Me	Н	Н	Н	Н	Н	0- But	5-Methyl-3- ruthinovylKaempferol
70	Me	Н	Me	Н	Me	OMe	H	5,7,3',4'-
71	Н	Н	Н	Н	Н	ОН	0- But	Rutin
72	П	H	[]	OII	П	11	II	Iscoscutellarein
73	Н	Н	Me	Н	Me	ОМе	OMe	5-hydroxy-3,7,3',4'- tetramethoxyflayone
74	Н	OMe	Me	Н	Me	OMe	OMe	5-hydroxy-3,6,7,3',4'-
75	П	II	Ме	OMe	Me	OMe	OMe	5-hydroxy-3,7,8,3',4'-
74		OMA	Mo		Ма	11	TT	pentamethoxyflavone
70	11	OMe	Me	11	Me	11	11	trimethoxyflavone
77	II	OMe	Me	11	11	II	II	5,4'-dihydroxy-6,7-
70	TT	OM-	Ma	OM ₂	Мо	TI	11	dimethoxyflavone
78	п	оме	ме	оме	Me	н	н	5-nyaroxy-6,7,8,4 - tetramethoxyflayone
79	Н	OMe	Me	Н	Me	OMe	Н	5-hydroxy-6,7,3',4'-
								tetramethoxyflavone
80	II	OMe	Me	OMe	Me	OMe	OMe	5-hydroxy-3,6,7,8,3',4'-
81	Н	OMe	Ме	OMe	Ме	OMe	Н	5-hydroxy-6,7,8,3',4'-
								pentamethoxyflavone
82	Н	Н	Rut	Н	Н	ОН	Н	Luteoline-7-0-
83	н	н	Rut	н	н	OMe	н	rutinoside Chrysoeriol-7-0-
00		••	mat	.,		ome		rutinoside
84	Н	Н	Rut	Н	Me	ОН	Н	Diosmin
85	11	C-	Rut	11	Me	OH	II	6-C-b-glucosyl diosmin
		Glu						
86	II	C-Glc	Rut	C-Glc	Me	OH	II	6,8-di-C-b-glucosyl
87	н	н	Rut	н	Н	н	Н	Isorhoifolin

Glu: Glucose, Neohesp: Neohesperidose, Rut: Rutinose.



Figure 1. Chemical structures of flavonoids from C. sinensis peels.

common, that is the presence of the high amount of bioactive polymethoxy-flavonoids [29, 30] (PMF) some of which are rather ubiquitous, e.g. Nobiletin **53**, Sinensetin **51**, 3',4',3,5,6,7,8-Heptamethoxyflavone **55**; some other compounds



Figure 2.

Chemical structures of the disaccharides most commonly bound to flavonoids of C. sinensis peel.

containing one to six methoxy groups in place of the hydroxyl groups are present at variable amounts. The presence of one or more residual hydroxy groups in the molecule may result in a higher bioavailability and in other general differences in their mechanism of biological and therapeutic actions [30, 31].

The biological role of these secondary metabolites in the plant is still matter of debate. It has been proposed their involvement in the mechanism of defense of the fruits exposed to the attack of phytopathogens, such as *Phytophthora citrophthora* [32].

Further, the composition of the PMF blend can be employed for the chemiotaxonomic characterization of the *Citrus* genus [33].

However, it needs to be stressed that in many cases the reported compounds were recognized by mass spectrometry and electronic spectroscopy. It is not always a matter of simplicity to discern the exact structure of a given PMF and to discriminate between different regioisomers, normally quite similar in terms of mass and electronic spectra, if an isolation procedure is not conducted and followed by a complete bi-dimensional NMR characterization. Significant differences in the extract composition do arise also in consequence of the extraction method; nonpolar solvents such as Methanol, Chloroform Ethyl acetate let to obtain PMFs-rich extracts while, on the other hand, hydroalcoholic and aqueous extracts do contain a low concentration of PMFs and a higher concentration of un-methylated polyphenols as well as glycosylated compounds.

The biological activities disclosed for the flavonoids extracted from CSP are variegated. They include antioxidant [9, 34–39], anti-inflammatory [40, 41], antimicrobial [39, 42–44], antimalarial [45], antitrypanosomal [46], cardio-protective [47], anti-osteoporosis [48], anti-ulcer [49], vascular protective [50], anti-diabetes [51, 52], hepatoprotective [53, 54], neurotrophic [55], anti-adipogenesis and anti-obesity [56–58], anti-hypertensive [59], cataract prevention [60], sun protection [61], metabolic syndrome control [62]. Further, it has been demonstrated [63] that while both flavonoid set **40**, **42**, **43** and the PMFs **51–53** were able to inhibit the anion transportin polypeptide OATP2B1 in HEK293 cells, only the PMF group displayed this inhibitory activity also for the OATP1B1 and OATP1B3 carriers.

The most abundant PMF occurring in CSP, Nobiletin **53**, was proven to possess sevral bioactivities, such as antioxidant, anti-inflammatory, cancer preventive [64] and also a significant protective effect *in vivo* against the endotoxic shock [65] and ethanol-induced acute gastric lesions [66] in mice. Further, compound **53** demonstrated the capacity to induce autophagy in human keratinocyte HaCaT cells [67], vasodilatator effect in the rat aorta [68] and to protect the intestinal barrier from the demages induced by dextran sulfate sodium [69].

PMFs can be considered as especially promising lead compounds for cancer therapy as asignificant cytotoxic activity has been demonstrated toward a number of cancer cells [70, 71] with several mechanisms of action [72, 73]; the cytotypes investigated include MCF-7 [73–76], Hs578T triple-negative breast cancer [73, 77]; colon cancer cells CaCo-2 [19], LoVo [78], HTC-116 [79, 80] and HT-29 [79, 81]; lung cancer cells A549 [80, 82], H460 [82, 83], H1299 [82, 83]; gastric cancer cell lines AGS, BGC-823, and SGC-7901 [84]; leukemia cells HL-60 [85]. However, data regarding a possible antitumor activity *in vivo* are still rather uncommon. An interesting example is the case of the significant reduction of the intestinal tumor mass in ApcMin/+ mice treated with a CSP extract containing various PMF [86]. Further, CSP extract and pure Naringin **47** were tested for their efficacy against a YM1 esophageal cancer in an animal model [87].

Given the development of pharmacological applications of CSP extract components, further investigations are needed to better understand the bioavailability, safety, and efficacy of these compounds in humans. Most of the data reported concern *in vitro* experimentations or animal model tests. For example, the toxicity of Hesperidin **40** was evaluated [88] in Sprague Dawley rats showing a 50% lethal dose (LD50) of about 5 g/Kg body weight (BW) and a lowest-observed-adverseeffect level (LOAEL) of ca. 1 g/Kg BW.

In general, it should be emphasized as the body of evidence concerning the actual efficacy of sweet orange-derived compounds in human health is still far to be exhaustive. For example, while this work is under typewriting, a severe acute respiratory syndrome pandemic due to a COVID-19 virus is in act and a big deal of research has been being directed toward antiviral remedies and therapies. Research on nutraceuticals is not an exception and in particular some authors have shown by computational and molecular docking methods how Hesperidin **40**, the most abundant polyphenol obtained from *C. sinensis*, would be able to bind the spyke protein of this virus thus inhibiting its activity [89]. Despite their undoubted interest, these results need to be further investigated with different experimental approaches.

The pharmacological potential of pure Hesperidin **40** was also investigated for several relevant human morbidity, such as cancer, hypertension, and ulcer [90].

2.2.2 Hydroxy-acids

Several hydroxylated carboxylic acids belonging to several structural sub-classes are present foremostly in the extract obtained with mixed hydro-organic solvents, such as MeOH/water and EtOH / water [37, 38, 51, 78]; these include the aliphatic Ascorbic, Citric, Kojic, Lactic, and L-Malic acids; the aromatic 4-Hydroxybenzoic, Protocatechulic, and Gallic acids. Further, the cinnamyl compounds (**Figure 3**) Cinnamic (93), p-Cumaric (94), Caffeic (95), Ferulic (96), Sinapinic (97) acids, and Artepillin (98) were identified in some CSP extracts that showed relevant biological activities, such as antioxidant [34, 37, 38] and antidiabetes [51].

These organic acids are mainly found in free form but in some cases, they are esterified with a variety of alcoholic compounds, such as Ethanol in Ethyl gallate **99** [51], 2-Phenylethanol in Phenylethyl ester of Caffeic acid **100** [51] and (–)-Quinic acid in Chlorogenic acid **101** [51]. An interesting ester derivative (**102**) in which the anomeric hydroxyl of Glucose is esterified with a O-Caffeylsinapoyl acid unit was found in the methanolic extract of a Greek cultivar of *C. sinensis* [34].

It was shown [38] that the antioxidant properties of a CSP extract better correlated with the total phenols content (TPC) of the sample rather than with its total flavonoid content (TFC), as it can be expected from the known relevant antioxidant character of hydroxycynamic derivatives.



Figure 3. Chemical structures of cinnamic acids extracted from C. sinensis peels.

2.2.3 Coumarins

Coumarins are aromatic compounds biogenetically related to the o-hydroxysubstituted cynamic acids from which originate by the intramolecular condensation between the carboxylic and the o-hydroxy groups. These compounds are most commonly encountered in other species of *Citrus* taxa [91], such as *C. aurantium* (bitter orange), *C. limon*, (lemon), *C. limetta* (lime), *C. paradisi* (grapefruit) and only a few molecules of this class were Isolated from extracts of CSP endowed with activity against osteoporosis [48] and antioxidant [92]; these compounds are shown in **Figure 4**. As coumarins are relatively less common in *C. sinensis* cultivars compared to other species of the *Citrus* taxa, their rarity can be considered as a chemotaxonomic marker for *C. sinensis*.

2.2.4 Catechins

The NADPH dependent bioreduction of flavanols is the biogenetic origin of this class of compounds, present as minor constituents in CSP extract possessing significant antioxidant activity [38]; they are the two enantiomeric forms Catechin **113** and Epicatechin **114**, together with Epigallocatechin **115** (**Figure 5**).



Figure 4.

Chemical structure of coumarins extracted from C. sinensis peels.

2.3 Pectins

Pectins [93] are chemically definable as complex mixtures of polyglyconic acids in which a linear polymeric backbone is structured by a series of α (1 \rightarrow 4) linkages (**Figure 6**). The main sugar monomer is always Galacturonic acid with the presence





Figure 5. Chemical structure of catechins from C. sinensis peels.



Figure 6. Minimal representation of a Homopolygalacturonic acid domain of the linear primary pectin structure with a 1/3 Mol. /Mol. Esterification degree.

of possible heterogeneous domains of other sugars such as Xylogalacturonan and Rhamnogalacturonan-I. A variable amount of the free carboxy functions may be esterified with methyl groups, while the hydroxy groups at C-2 and C-3 positions of the sugar monomers may be acetylated. Even though the primary structure of the main chain is linear, a possible degree of ramification, depending on the pectin source, may also be found. The differences in the pectins composition and structures, depending on their natural source, do confer them different physio-chemical properties, such as water solubility, sol–gel concentrations, etc. On the ground of the degree of methylation of the acid moieties, pectins are classified as "low methoxyl" (LMP, -COOMe/-COOH <50% mol.) or as "high methoxyl" (> 50% mol). A simplified representation of pectin structure is given in **Figure 6**.

Pectins find many applications in the food and drug industry as a thickening and gelling agents, excipients, and colloidal stabilizers [93].

As it has been already mentioned, the extraction method does affect the structure and the properties of the final product; the traditional acidic water extraction implies a certain degree of hydrolytic deterioration, so that new extraction technologies have been being investigated to improve the quality of the final pectins, that is microwave-assisted extraction (MAE) [94] and ultrasounds assisted extraction (USAE) [35, 95].

2.4 Enzymes

As it can be easily argued, the CSP cellular system, whose genomic profile has been fully characterized [96], is the site of a complex network of enzymatic activity. Some of the enzymes of CSP have been characterized and employed in many applications.

The acetylesterase (international enzymatic classification: EC 3.1.1.6) from CSP is known since 1947 [97] and was isolated and characterized [98]. The acetylesterase activity of the partially purified enzyme was used for the removal of the acetyl group at the 3 positions of β -lactamic antibiotics **116** [98] (**Figure 7a**). Further, the whole CSP, as well as pomace from the industrial waste of the orange juice production, was successfully employed to catalyze several relevant biotransformations [99] such as the conversion of Geranyl acetate **118** to Geraniol **119** (**Figure 7b**) and the di-acetoxynaphtalene derivative **120** to the vitamin k1 precursor **121** (**Figure 7c**).



Figure 7. Chemical reactions biocatalysed by enzymes from C. sinensis peels.

Recently, partial purification and functional characterization of a Uronic acid oxidase from CSP was accomplished [100]; this enzyme promotes the oxidation by O_2 of Galacturonic acid **122** to Galactaric acid **123** (Figure 7d).

2.5 Miscellaneous

2.5.1 Highly lipophilic compounds

The waxy environment of flavedo in CSP does contain several long-chain saturated and unsaturated compounds: alkanes, fatty acids, waxes, higher terpenoids.

Tetracosane, Tetratriacontanoic acid, and Ethyl pentacosanoate were identified in CSP of a Pineapple variety [101]. Further, some carotenoids were identified in the CSP extract obtained with a solvent mixture composed of Ethanol, Ethyl acetate, Petroleum ether 1: 1:1 [102]. This complex blend of carotenoids includes α - and β -Carotene, Phytoene, Phytofluene, (all-E)- and (9Z)-Violoxanthin, (all-E)-Neoxanthin, (13Z)-, (13Z')- and (all-E)-Lutein, (9Z)-Zeaxanthin, (all-E)-Zeaxanthin; the mono and di-esters of violaxanthin, antheroxanthins, Xanthophyll, β -Citraurin with various fatty acids, including Lauraic, Myristic, Oleic, Palmitic, Stearic. The composition of the blend has been correlated with the maturity stage of the fruit.



Figure 8.

Primary structure of cyclic peptide isolated from the C. sinensis peels.

2.5.2 Peptides

Three cyclic peptides have been isolated from the hot water extract of CSP and were structurally characterized by FAB-MS and 2D-NMR techniques [103]. Their amino-acidic sequences, including a mostly lipophlic heptapeptide **124**, a di-hydroxylated heptapeptide **125**, and a Glutamate-rich octapeptide **126**, are reported in **Figure 8**.

3. Conclusions

The chemical richness of the primary and secondary metabolome of *C. sinesnis* species is undoubtedly impressive. Thousands of different compounds belonging to dozens of structural classes have been isolated and described. The most deeply investigated are sure, on one hand, the mixtures of volatile compounds composing the blend of the essential oil and, on the other hand, polyphenols, especially flavonoids.

The chemical composition of the extract from the exocarp of *C. sinensis* does differ from the composition of juice, or leaf extracts for some aspects [104]: the presence of a higher amount of more lipophilic compounds such as polymethoxy-flavonoids, r carotenoids, higher alkanes; a lesser extent of lighter terpenoids, a lower content of glycosylated compounds, the absence of cyanidins and sterols.

It is also a matter of fact that several interesting bioactivities were disclosed in the last years for the *C. sinensis* extracts that have been variously associated with the well-recognized beneficial effects that regular sweet oranges consumption may have on human health. However, a great deal of research work is still needed to clarify the molecular basis and the mechanism of these chemopreventive effects and to relate them with precise chemical entities that can be recognized as valuable nutraceuticals, as it is already the case for the well-established antioxidants Ascorbic acid, Hesperidin, Hesperetin, Quercetin, etc.

Conflict of interest

The author declares no conflict of interest.

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Chapter 12

Physiological Functions Mediated by Yuzu (*Citrus junos*) Seed-Derived Nutrients

Mayumi Minamisawa

Abstract

This section is focused on the physiological functions of yuzu (*Citrus junos*) to improve health. The modern lifestyle involves number of modern lifestyles involve various factors that may increase the production of active oxygen species. Nutritional supplements and medicines are commonly utilized to maintain health. Yuzu seeds contain >100-fold the limonoid content of grapefruit seeds and are rich in polyamines (PAs), including putrescine, spermidine, and spermine. Limonoid components mediate the antioxidant properties of citrus. Limonoids and PAs convey various bioactivities. PAs are closely associated with maintaining the function of the intestinal mucosal barrier, which might be involved in the metabolic processes of indigenous intestinal bacteria and in the health of the host. After ingestion, food is digested and absorbed in the intestinal tract, which is also responsible for immune responses against food antigens and intestinal bacteria. Detailed investigations of the physiological functions of extracted yuzu seed extracts may help to develop new treatment strategies against diseases associated with inflammatory responses.

Keywords: Yuzu (*Citrus junos*), limonoids, polyamine, gut microbiota, anti-inflammatory, short-chain fatty acid (SCFA), central neurodegenerative disease

1. Introduction

In 1997, the World Cancer Research Fund published 14 articles concerning dietary recommendations in addition to smoking cessation for the prevention of cancer in *Food*, *Nutrition and the Prevention of Cancer: a Global Perspective* (2007 revised edition) to promote international awareness of the relationship between nutrition, diet, and cancer. Articles 1, 4, and 5 strongly recommend the consumption of foods of plant origin, and especially emphasized the importance of fruits and vegetables for the prevention of many types of cancer [1].

There are more than 1000 species of citrus and various varieties account a major part of all fruit production worldwide. In particular, citrus species native to Asia are believed to have originated in the Assam area of India around 30–40 million years ago and were propagated in China, Thailand, Malaysia, Indonesia, and Taiwan before being brought to Japan [2]. The tachibana orange, which is the oldest variety of mandarin orange in Japan, was introduced in Japan from Taiwan via the Korean Peninsula from mainland China, and is listed in the *Manyoshu*, the oldest extant collection of classical Japanese poetry compiled sometime after 759 AD during the Nara period, as the only citrus fruit that existed in the wild. After that, it is estimated that the daidai, an Asian variety of bitter orange, and other small oranges arrived in Japan around 2 to 300AD. Yuzu (*Citrus junos Sieb. ex Tanaka*) originated in China and was introduced to Japan and other countries around the 4th to 8th centuries, as this fruit is mentioned in the *Shyoku-Nihongi*, an imperially-commissioned Japanese history text completed in 797 AD.

The traditional Japanese meal *washoku* was recognized as a UNESCO Intangible Cultural Heritage of Humanity in 2013. The Japanese have the highest life expectancy of any other ethnicity. Therefore, washoku has attracted attention as a healthy diet. Especially, yuzu is an essential ingredient of the Japanese diet in the winter months. A traditional 0sechi dish, including yuzu, to be eaten on New Year's Day is shown in the photo in **Figure 1**.

Yuzu is a commercially important fruit, as compared to other sour citrus fruits, and has become very popular in Japan. Although rarely eaten as a fruit, yuzu is a common ingredient in Japanese cuisine, where the aromatic zest (outer rind) as well as juice are used much in the same way as lemons in other cuisines. The yuzu fruit and juice are traditionally used in making vinegar and seasoning yuzu peel and juice, and along with sudachi, daidai, and other similar citrus fruits, are integral ingredients in the citrus-based sauce ponzu. In addition, yuzu is often used as an ingredient in alcoholic drinks, such as the yuzu sour. Recently, yuzu kosho "yuzu and pepper" has become a very popular spicy Japanese sauce made from the peel (zest) of green or yellow yuzu, combined with green or red chili peppers and salt. Yuzu is also well-known because of its pleasant aroma and essential oil of the outer rind. In fact, in Japan, it has been customary since ancient times to take a bath with yuzu in hot water during the winter solstice. The yuzu peel is particularly high in aromatic compounds and pectin; therefore, the waste peel from juice extraction is sometimes used to produce essential oils and flavorings as well as for medicinal purposes. Similarly, yuzu is industrially used in the production of sweetened beverages, cosmetics, and perfumes, as well as oils for aromatherapy [3]. Only a small portion of produce is used for natural medicine, while satsuma mandarins, oranges, and grapefruits are commonly used for the production of fruit juices.

There is a reason why yuzu is not often eaten as a fruit in Japan because it contain large seeds that convey a bitter taste to the juice. The well-known constituents of citrus fruits include essential oil components, flavonoid glycosides, and other basic substances with biological activities, including limonene, a major component of essential oils found in the juice and rind, polymethoxyflavones, coumarins, carotenoids, which are pigments, vitamins, and terpenoids [4–7]. In the past, the bitterness of citrus juices, skins, and seeds hindered the demand for citrus fruits.



Figure 1. *A typical Osechi package for New Year's day in Japan.*

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Much research has been conducted to produce bitter-free citrus fruits. At the time when there was very little demand for bitter fruit and juice, Hasegawa et al. [8–10] reported that the high physiological activity of limonoids was responsible for the bitter taste in citrus juice. Limonoids are a group of triterpene derivatives found in plants of the Rutaceae and Meliaceae families. So far, more than 300 types of limonoids have been reported, and about 100 types have been isolated from the neem and sendan plants of the family Meliaceae.

Limonoids are characterized by a furan ring at C-17, a lactone ring at C-3 or C-6, and an epoxide between C-14 and C-15 (**Figure 2**). Of the four basic structures of limonoids, reversible opening and closing occurs in A, D lactone rings. For example, in the case of limonin, one of the major limonoids, a closed D-ring creates a bitter taste, while the open D-ring form (limonate A-ring lactone) has no bitter taste. Furthermore, the limonate A-ring lactone at C-17 is converted into the limonoid glycoside 17- β -D-glucopyranoside, which is D-glucose bound with β -glucoside.

Until recently, not much had been known about the metabolism of limonoids found in fruit. In 1991, Hasegawa et al. [11] discovered that limonate A-ring lactone, an open D-ring form of a limonoid aglycone, was metabolized to a glucoside derivative in the late stages of fruit growth and maturation, and suggested that this occurred independently in both the seeds and fruit. It is known that the limonoid aglycone and glycosides accumulate in the seeds [12].

To date, 36 types of aglycones and 17 types of glycosides have been identified mainly in citrus fruits of the family Rutaceae, which is composed of 160 genera and about 2,070 species [13]. The first study of the physiological effects of *Citrus limon*oids reported inhibitory effects on the eating behaviors of armyworms and predatory insects [14]. Strong inhibitory effects on eating were subsequently observed in termites. It has been reported that the ligand activity of the bile acid receptor TGR5 increases the inhibition of tumor formation and the activity of glutathione S-transferase, which is a detoxification enzyme that assists with the excretion of toxic substances by the liver and digestive organs, as well as increasing anti-obesity effects via insulin and increased heat production [15]. The seeds of citrus fruits contain particularly potent limonoids. The metabolic pathway of limonoid biosynthesis in citrus fruits has been nearly elucidated by Hasegawa et al. [16] with the use of 14C-labeled radioisotopes as tracers (**Figure 3**). Within the phloem of the stem, nomilin is synthesized through the metabolism of acetic acid, mevalonic acid,



Figure 2. D lactone ring structures of limonin.



Figure 3. The limonoid biosynthesis pathway of yuzu seeds.

and furnesole phosphate. Especially in the stems of seedlings, nomilin synthesis becomes very active [17].

Nomilin is synthesized only in the stem and then transferred to the leaves, fruits, and seeds where it is metabolized into other limonoids. Since metabolism proceeds with the D-ring open form, limonoids exist in the D-ring open form in the stems, fruits, and leaves, and mostly form glycosides. In seeds, metabolism to limonoid aglycone with a closed D-ring also proceeds at the same time, so that aglycone and glycoside are accumulated simultaneously. Therefore, the aglycon in the fruit tissue decreases during maturation, but continues to accumulate in the seeds.

2. Bioactive substances of yuzu seeds

Minamisawa et al. [18, 19] has been searching for new antioxidants to maximize the original functions of living organisms with the use of waste resources derived from natural products, including yuzu seeds, that can be regenerated as many times as possible in the human life span. In 2011, the oldest original species of yuzu in Japan was discovered in the village of Mizuo, which is located in the north-west of Kyoto city [2]. This yuzu is characterized by "seedlings" grown from seeds in the land associated with Emperor Seiwa (9th century) and Emperor Hanazono (15th century), and it takes about 20 years to harvest. Most of the citrons cultivated in Japan today originate from the yuzu of Mizuo, which is cultivated mainly by grafting, and the growth is faster than that of seedlings. Since the yuzu of Mizuo is considered to be the finest quality, the fruit is highly demanded by high-end restaurants serving Japanese cuisine in Kyoto. However, yuzu seeds, which are closer to the ancestral citrus, account for 20–30% of the fruit weight, but are discarded as waste after the juice extraction process.

Hence, our team chose to evaluate the this development of active natural resources would encompass the application in nutrition and environmental attributes of yuzu seeds as natural resources with bioactivities [20]. In 2014, we reported the development of a relatively simple technique to simultaneously extract secondary metabolites of yuzu seeds, including expensive limonoids and yuzu seed oil with high total antioxidant capability, from the waste of fully ripe fruits [2]. Yuzu seeds contain higher amounts of fat-soluble limonoid aglycones, water-soluble limonoid glycosides, and oil than other citrus fruits (**Figure 4**).

Analysis of the components of limonoids from yuzu seeds by high-performance liquid chromatography-mass spectrometry identified five limonoid aglycones (deacetylnomilin, limonin, nomilin, obacunone, and ichangensin) and eight

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Figure 4.

The extracted components from Yuzu seed.

limonoid glycosides (limonin glucoside, ichangin glucoside, deacetyl nomilinic acid glucoside, deacetylnomilin glucoside, nomilin glucoside, nomilinic acid glucoside, ichangensin glucoside, and obacunone glucoside) (**Figure 4, Table 1**). Yuzu seed oil extracts (**Table 2**) contain large amounts of oleic and linoleic acids ([2], in preparation). The contents of limonoids extracted from yuzu seeds compared with the results of previous studies are shown in **Table 1** [21–23].

As compared to other citrus seeds, the concentrations of limonoid aglycones extracted from the seeds of yuzu fruit from Kyoto were two- or three-fold greater than in fruits from Tokushima and California (334 vs. 167 and 0.94 mg/g, respectively). According to Nogata [22], the iyokan fruit (*C. iyo*), Valencia orange (*C. sinensis Osbeck*), and hyuganatsu (*C. tamurana Hort. ex Tanaka*) belong to the same family as the *daidai* (*C. aurantium group V*). Hence, the limonoid compositions of these varieties are similar (**Table 1**). Although the amount of nomilin in the Valencia orange is similar to that in the *iyokan* and *hyuganatsu* varieties, the amount of limonin is approximately two-fold greater, while the amount of deacetylnomilin is higher and that of obacunone is significantly lower.

The yuzu and hanayu (*C. hanaju*) varieties are classified to yuzu group VI. However, both the compositions and amounts of the limonoid aglycones differed markedly between these two species in the present study, which may be attributed to differences in the metabolism of the seeds and fruits [24, 25]. For this reason, the ratio of aglycone to glycosides in mature fruit tissues is mostly due to glycosides, whereas the glycoside content in seeds may be the same or lower than that of aglycones (**Table 1**). These findings indicate that limonoids are biosynthesized completely independently of fruit tissues and seeds.

Nogata et al. [22] pointed out that the high amounts of glycosides in seeds of the iyo and shiikuwasha fruits could be due to the high activity of uridine diphosphate-D-glucose transferase, and perhaps in the yuzu seeds as well. The high limonoid content in the seeds of yuzu fruit grown in Kyoto is thought to be related to the seedling cultivation method. Similar to yuzu seeds, the glycosides deacetyl nomilinic acid glucoside and deacetylnomilin glucoside, but not ichangensin glucoside, accumulate in hanayu seeds. Ichangensin is reportedly metabolized from nomilin through the intermediaries deacetylnomilin and deacetyl nomilinic acid [26, 27].

				Content (1	ng/g of dry seeds)			
Seeds		Lim	onoids aglycone	SS			Total	
	Nomilin	Deacetyl-nomilin	Limonin	Obacu-none	Ichan-gensin	Aglycones	Glucosides	Aglycones/glucosides
Yuzu (Kyoto, Japan)	114	106	95.0	16.7	2.10	334	452	0.74
Yuzu (Tokushima, Japan)	58.4	48.3	54.0	5.20	1.60	167	192	0.87
Yuzu [*] [21] (California)	0.25	0.22	0.47		0.23	0.94		
Hanayu [22]	0.12	0.86	0.54	0.07		1.59	15.9	0.10
Shiikuwasha [22]	0.96	I	1.87	0.45		3.28	12.7	0.30
Iyo [22]	2.53	0.72	4.57	0.91	1	8.73	4.46	2.00
Hyuganatu [23]	3.73	0.35	4.68	0.28	1	9.04	3.12	2.90
Grapefruit [23]	1.84	1.10	19.1	1.86		23.9	6.98	3.40
Lemon [23]	3.03	I	8.95	0.58		12.6	6.37	2.00
Valencia [23]	2.30	1.24	10.0	0.08	1	13.6	8.71	1.60
Units are mg/g of fresh weight.								

Table 1. Limonoids in various citrus seeds.

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Fatty acid	16:0	16:1	18:0	18:1	18:2n-6	18:3n-3	20:0	20:1	24:0
%	19.7	0.5	3.8	37.9	34.7	1.6	0.3	0.2	0.1

Table 2.

The FA content was determined by gas chromatography with the use of a GC-14 gas chromatograph (Shimadzu corporation, Kyoto, Japan) equipped with a DB-1 column (30 m, 0.25 mm, Agilent Technologies, Inc., Santa Clara, CA, USA), which was maintained at a constant temperature of 300°C. The yield of yuzu seed oil was 100 mg/g of dry seeds.

The hanayu are, therefore, different from other citrus varieties. Although both belong to the same yuzu group, there are differences in characteristics, such as aglycones contents.

Several *in vitro* studies have shown that limonoid components mediate the antioxidant properties of citrus. Reactive oxygen is believed to be a factor in diseases with underlying cellular disorders [28]. Modern lifestyles and diets involve a number of factors that can increase the production of active oxygen species, which can overwhelm the body's self-regulating defense mechanisms [29, 30]. One way to protect oneself from active oxygen species is to consume foods containing antioxidants. One of the main reasons for our interest in antioxidants is the link between active oxygen species and aging.

Limonoids components of *C. junos* are known to possess a vast range of biological activities, including antioxidant functions, protective effects on vascular endothelial cells [31], and anti-carcinogenic activities [15, 20, 32–34].

Study also evaluated the *in vitro* antioxidant activities of yuzu seed aglycones, glycosides, and oil extracts. Notably, the yuzu seed oil, the potential extracts had high antioxidant activities due to the presence of lipophilic aglycones (**Figure 5**, new unpublished data). Yuzu seed oil is a semi-drying oil that contains large amounts of unsaturated fatty acids (FAs), mainly oleic acid and linoleic acid, in addition to a lot of palmitic acid.

Virgin yuzu seed oil, which is obtained by a pressing process without heating, contains about 2% of limonoid aglycones. Pure oil with the composition shown in **Table 2** can be obtained by heating and drying roasted yuzu seeds, followed by extraction with an organic solvent, such as hexane, and purification.

Lipophilic limonoid aglycones, which were extracted from the residual extracts of yuzu seed oil [2], were composed of the following concentrations of limonoids per gram of dry seeds: deacetylnomilin, 105 mg; limonin, 95 mg; nomilin, 115 mg; obacunone, 17 mg; and ichangensin, 2.1 mg.



Figure 5.

Total potential antioxidant activities of various plant seed oils and yuzu aglycones by the total potency of antioxidants that are soluble in oil method [35]. Measurements were performed 4-6 times or more. The inhibition ratio is presented as the average value \pm standard deviation (S.D.).

The total potential antioxidant capacity of yuzu seed oil and lipophilic limonoid aglycones was measured by utilizing the reduction reaction of copper (Cu⁺⁺/Cu⁺) [35]. Many other plant oils, including olive oil [36], tea tree oil, grape seed oil [37], and neem seed oil [38], which have strong antioxidant activities, were measured at the same time for comparisons. Among all of the tested plant seed oils, limonoid aglycones extracted from yuzu seeds had the highest antioxidant capacity, followed by yuzu seed oil, neem seed, grape seed, tea tree, olive oil, and pure yuzu seed oil. The antioxidant capacity of pure yuzu seed oil was approximately 6–9-fold greater than that of palmitic acid, oleic acid, and linoleic acid.

While water-soluble antioxidants are rapidly excreted through the urine if an excessive amount is ingested, fat-soluble antioxidants are adsorbed onto lipoproteins and cell membrane lipids, and are therefore considered to exhibit a higher activity in the body. For this reason, fat-soluble antioxidants are expected to be beneficial in preventing diseases caused by oxidative stress. Vitamin E, oryzanol, and carotenoids are well-known examples of fat-soluble antioxidants. Neem (*Azadirachta indica*) seed oil, which has the same total antioxidant capacity as yuzu seed oil, contains the triterpene derivative azadirachtin, which is similar to the triterpene limonoids of yuzu seed oil, which is a potent insect repellent [39]. Press-extracted virgin olive oil contains oleocanthal that has a potent anti-inflammatory effect strikingly similar to that of ibuprofen. Both of these molecules inhibit the same cyclooxygenase enzymes in the prostaglandin-biosynthesis pathway [40].

The result in **Figure 5** suggest the presence of other types of fat-soluble antioxidants. Limonoid aglycones also contribute to the high antioxidant capacity of yuzu seed oil.

3. Yuzu seeds contain arginine and polyamines (PAs)

Atherosclerosis has become a serious health concern worldwide, as one-third of the global population is at risk for diseases associated with arteriosclerosis, which accounts for about half of deaths in developed countries. In particular, cardiovas-cular disease (CVD), which is a consequence of atherosclerosis, is the leading cause of death in industrialized nations. Besides lifestyle habits, body weight, socio-economic factors, and certain pre-existing conditions, a number of foods seem to play a role in the incidence of CVD [41, 42]. In addition, many studies have suggested the importance of inflammation in atherosclerosis and CVD [43, 44].

Some food components with anti-inflammatory properties can decrease the risk of CVD [44, 45]. Many foods contain wide-ranging concentrations of natural PAs, such as spermidine (Spd) and spermine (Spm), which suppress the synthesis of pro-inflammatory cytokines [46, 47]. In particular, an epidemiological survey of Westerners found that "people who eat cheese or yogurt every day are less likely to have myocardial infarction." [48]. The Japanese consume a lot of traditional fermented foods, mainly soybeans, which are thought to suppress arteriosclerosis [49]. PAs concentrations are relatively high in yogurt, cheese, and traditional Japanese foods. PAs are aliphatic amines that are essential for the growth of all living cells [50]. PAs exist primarily in association with RNA and are involved in promoting the synthesis of specific proteins and overall protein synthesis via the ribosome activation. As shown in **Figure 6**, the PAs comprising Put (NH₂ (CH₂)₄NH₂) \rightarrow Spd (NH₂ (CH₂)₃NH(CH₂)₄NH₂) \rightarrow Spm (NH₂(CH₂)₃NH(CH₂)₄NH₂) are produced from arginine via ornithine or agmatine [51].

PAs have been implicated in the regulation of several growth and development processes in plants, including cell division, morphogenesis, flower initiation, pollen

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Figure 6.

PA synthesis and degradation [50, 51]. AcPAO, acethylpolyamine oxidase; ODC, ornithine decarboxylase; SAMDC, S-adenosylmethionine decarboxylase; SMO, spermine oxidase; SPDS, spermidine synthase; SPMS, spermine synthase; SSAT, spermidine/spermine N1-acetyltransferase.

Citrus			Arginine		
-	Putrescine	Spermidine	Spermine	Total PAs	
Yuzu (<i>Kyoto, Japan</i>) [58]	294.8	117	34.6	446.4	36625
Only seed	79.4	117	34.6	231	30540
Yuzu (Tokushima, Japan) [58]	466	144.6	24.7	635.3	20207
Only seed	125	89.5	24.7	635	14007
Lemons (Kamogawa, Japan)	320.0	56.2	3.8	380.0	37933
Only seed	71.1	56.2	3.8	371	33950
Grapefruit [54]	436	19.5	1.5	457	
Lemons, Limes [54]	390.5	19	4.5	414	
Oranges, Mandarins [54]	432	15.5	0	448	
Bean	Putrescine	Spermidine	Spermine	Total PAs	
Soybean [54]	297	909	235.5	1442	
Beans [57]	147	644.8	299.7	1092	
Sesame seed [54]	29	126	22	177	

Table 3.

PAs and arginine contents in several citrus (nmol/g).

tube growth, and senescence [52]. Analyses of the PA contents of various fruits have mainly been conducted in Europe [53, 54].

Recent studies have indicated that citrus limonoids have antitumor, detoxification, and anti-obesity effects [55, 56], which may indirectly contribute to the suppression of acrolein production, which is a side reaction product of PA metabolism (**Figure 6**). Hence, we measured the PA and arginine contents of yuzu, which produces very high concentrations of limonoids, as well as lemons produced in Japan for comparison. The PA contents, as determined by high-performance liquid chromatography, as well as the arginine and free arginine contents, as determined by automated amino acid analysis, of various citrus fruits are shown in **Table 3**.

As compared with the juice and peel of yuzu and lemons, the seeds contain higher quantities of PAs and arginine. The Put contents are high in all citrus fruits, but the quantities of Spd and Spm in yuzu seeds were 5–23-fold greater than the reference values. The PA contents of yuzu and lemon fruit are not high as compared to legumes [57], but when the limonoid and arginine contents are also considered, these fruits have high levels of functional constituents.

As mentioned earlier, limonoids and PAs have various bioactivities and reportedly have strong anti-inflammatory capabilities. Hence, the potential antioxidant activities (i.e., H₂O₂-scavenging activity, 2,2-diphenyl-1-picrylhydrazyl radicalscavenging activity, and inhibition of superoxide dismutase [SOD] and antioxidants with SOD-like activities) of PAs (Put, Spd, and Spm) and arginine were investigated (in preparation). The results showed that these compounds have no antioxidant activities or only weak (less than 10%) inhibitory potential. As reported in many studies, the anti-inflammatory activities of PAs and arginine are due to factors other than antioxidant capacity.

4. Yuzu seed limonoids or Spm increased survival of mice with Sandhoff disease

In our previous study, we investigated the life-extending effect of limonoids (lipophilic limonoid aglycones) and Spm as an exogenous anti-inflammatory component in a mouse model of Sandhoff disease (SD), which is a lysosomal disease [58]. Lysosomal storage disorders are caused by functional defects of proteins that are essential for normal lysosome function, such as enzymes that play critical roles in the intracellular digestion of glycoproteins, glycolipids, glycosaminoglycans, and other macromolecules [59]. SD is an autosomal recessive hereditary disease [60]. The gangliosides GM2 and GA2 accumulate in the nervous system, resulting in severe developmental and neurological disorders, and death, which usually occurs during infancy because of the lack of effective treatment methods. Neurological dysfunction is the major clinical manifestation of GM2 gangliosidosis [61–64].

SD mice present with trembling, startled responses, and decreased motor activities from 11 to 15 weeks of age (105 days) due to damage caused by microglial activation, macrophage infiltration, and oxidation associated with the accumulation of glycolipids. It has been suggested that inflammation may be fatal [51]. The therapeutic effects of enzyme replacement therapy and anti-inflammatory drugs have been reported [65]. Inflammation due to the accumulation of lipids is inhibited by antioxidant and anti-inflammatory treatments, which can delay disease progression, but no cure exists at present.

We consider the degeneration of the nervous system might be rooted in oxidative stress and inflammation. Given that dietary interventions can moderate these phenomena, consuming foods with antioxidants and anti-inflammatory components, such as limonoid aglycons (limonoids) and Spm, could effectively combat or minimize neurological damage. Therefore, the inhibition of SD pathologies could be promoted by factors other than suppressing the storage of gangliosides.

Preventing inflammation appears to be one of the most effective approaches for increasing longevity [66, 67]. To test this hypothesis, the life spans of SD mice treated with limonoids or Spm were assessed. The prognostic outcomes of SD mice, a typical model of abnormal glycolipid metabolism in humans, were observed after administration of limonoids extracted from yuzu seeds and Spm. The treated mice
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Figure 7.

H&E-stained thalamus sections of control SD mice (A and C) and SD mice administered yuzu limonoids (B) and Spm (D). Enlarged cells with ganglioside storage are indicated. The numbers of neurons in the thalamus and midbrain of SD mice administered limonoids (B) and control SD mice (A). Data are presented as the mean ± S.D. *p ≤ 0.05 (Student's t-test).

lived significantly longer than untreated littermates (9–10%, p < 0.01) and had a slower rate of disease progression (p < 0.01) [58]. When limonoid treatment was combined with Spm therapy, synergy resulted in a maximum improvement of 12% in survival (p < 0.001) (in preparation). The hematoxylin and eosin (H&E) staining results of thalamus sections of SD mice following administration of limonoids or Spm are shown in **Figure 7**.

H&E staining results of the neural tissues of the SD control mice (A) and(C) correspond to SD mice treated with limonoids (B) and Spm (D), respectively.

Gangliosidosis and inflammatory/autoimmune diseases are characterized by degeneration and the accumulation of fat, granulovacuolar degeneration, rod-shaped microglia, and neuronal inflammation in metabolic diseases, as determined by analyses of pathological tissues (Tokyo Metropolitan Institute for Medical Science). The characteristic degeneration was clearly decreased in SD mice treated with limonoids or Spm. The numbers of neurons in the thalamus and midbrains of SD mice treated with limonoids were higher than those in the control SD mice. These results demonstrate that inflammation contributes to disease progression and the anti-inflammatory effects of Spm and limonoid therapies as a potential adjunctive approach to slow the clinical course of inflammatory diseases.

5. Bacterial flora analysis of SD mouse feces by the 16S ribosomal DNA (16S rDNA) terminal restriction fragment length polymorphism (T-RFLP) method

PAs possess anti-inflammatory activities by inhibiting the synthesis of inflammatory cytokines by macrophages and the regulation of nuclear factor- κ B activation, which are closely associated with maintaining the intestinal mucosal barrier function [68]. Bilateral signals between the intestine and brain are involved in the control of nerve, hormone, and immune activities, as well as prolonging longevity [69]. Recent studies have shown that bilateral signals between the brain and intestine are important for maintaining homeostasis and extending the life span [70]. In particular, the functions mediated by PAs may be involved in metabolism by indigenous intestinal bacteria and the health of the host [71].



Figure 8.

Estimated ratios (%) of the taxonomic categories of the bacterial flora at the order level were identified by T-RFLP analysis of 16S rRNA in the feces of SD mice at 12 weeks of age in the control, Spm, and limonoids groups (n = 9/group). Values are presented as the mean \pm S.D. data of the treatment groups are plotted against those of the control group. * $p \le 0.05$, ** $p \le 0.001$, and *** $p \le 0.001$ vs. the untreated control SD mice or each (dashed line----). All experiments were performed at least three times.

At 12 weeks of age for which the survival period was extended by limonoids or Spm, T-RFLP analysis of 16S rRNA was performed to classify the intestinal microbiota at the order level for each mouse group (**Figure 8**). The results showed that the taxonomic groups of the bacterial flora in feces after administration of limonoids or Spm were completely different from those of the SD control mice.

The bacterial flora in feces after administration of limonoids or Spm had increased proportions of *Bacteroidales* and *Clostridiales*. However, *Lactobacillus* was remarkably prevalent in feces of the SD control mice. The abundance of *Clostridiales* were significantly increased in the feces of SD mice treated with Spm, whereas *Bacteroidales* were increased in the feces of SD mice treated with limonoids. The administration of Spm or limonoids slightly increased the proportion of *Erysipelotrichales*.

It is generally known that the abundance of *Erysipelotrichaceae* is increased due to fat accumulation in mice [72]. In this case, it was possible that the bacterial flora of the SD control mice caused dysbiosis [73]. In the SD control mice, dysbiosis may have been due to suppressed absorption of dietary fats and other nutrients. Even more interesting was the significant appearance of *Verrucomicrobiaceae* in feces after the administration of limonoids or Spm, which were not found in feces of the control SD mice. *Verrucomicrobiaceae* include mucin-degrading bacteria that are also present in the human intestine, and especially *Akkermansia*, which promote the suppression of obesity, diabetes, and inflammation [74, 75].

It will be necessary to investigate the specific bacteria involved in more detail. Unfortunately, the T-RFLP method made it difficult to analyze the bacterial flora in more detail, and it was not possible to identify particular species. We are currently preparing a report of the findings of next-generation sequencing that allowed for more detailed classification.

6. Short-chain fatty acid (SCFA) production in SD mouse feces

There have been many reports of the relationships between chronic inflammatory diseases and the intestinal bacterial flora that have helped to clarify the balance

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between the intestinal ecosystem and diseases related to the intestinal tract. For example, genetic abnormalities and the breakdown of the intestinal ecosystem have been detected in inflammatory bowel disease [76]. In particular, members of the genus *Clostridium* promote the production of butyric acid, induce an immune response in the intestinal mucosa, and promote the differentiation of regulatory T cells (Tregs) that contribute to suppression. Thus, changes in intestinal *Clostridium* are considered to be closely related to the onset of inflammatory bowel disease [77, 78]. It has been reported that the SCFAs produced by intestinal bacteria may function as bio-modifying factors. Hence, the SCFA composition of feces from the same mice at 12 weeks were determined (**Figure 9**).

The production levels of SCFAs comprising acetic acid, propionic acid, and butyric acid were increased in mice administered limonoids or Spm as compared to SD control mice. In particular, the production levels of all SCFAs were higher in SD mice following administration of limonoids or Spm. The experimental results demonstrated differences between the fecal microflora composition and these metabolites after administration of limonoids or Spm. Butyric acid is a SCFA that is produced by clostridia [78].

As shown by the results presented in **Figures 8** and **9**, the addition of Spm to the diet clearly increased the proportion of *Clostridiales* and butyric acid in feces. Previous metabolomic analyses have shown that butyric acid contributes to the induction of Treg differentiation in the colonic mucosa. Thus, butyric acid functions as a histone deacetylase inhibitor and as an immunomodulator responsible for inducing Treg differentiation in the colonic mucosa, as well as the activation of dendritic cells [79]. Acetic acid produced by intestinal bacteria suppressed colitis in a mouse model by promoting apoptosis via the GPR43 receptor expressed by neutrophils and plays a central role in the inflammatory reaction [80, 81]. Furthermore, the addition of limonoids seems to contribute to the production of acetic acid and propionic acid as well as butyric acid.

Acetate, butyrate, and propionate are produced by members of the intestinal microbial community through fermentation of dietary fibers and starches, which are unable to be broken down by host metabolism [82]. In turn, these metabolites are sensed by host cells through various G-protein coupled receptors, known as free



Figure 9.

SCFA contents (μ mol/g) in feces of SD mice in the control, Spm, and limonoid groups. SD mice were treated at 12 weeks of age (n = 9/group). Values are presented as the mean \pm S.D. data of the treatment groups are plotted against those of the control group. * $p \le 0.05$, ** $p \le 0.001$, and *** $p \le 0.001$ vs. the untreated control SD mice or each (dashed line----). All experiments were performed at least three times.

fatty acid receptors, and intracellular peroxisome proliferator-activated receptor gamma. Furthermore, SCFAs can also regulate cellular responses through inhibition of histone deacetylases. Examples of the effects of SCFAs on the host include differentiation of Tregs and macrophages, and downregulation of pro-inflammatory mediators. These effects underline the fine balance that SCFAs help to maintain between intestinal immunity and inflammation [82–84].

These results suggest that yuzu conveys anti-inflammatory and lipid metabolism-promoting activities in mice following administration of limonoid aglycones and Spm. Thus, the metabolites of intestinal bacteria may be indirectly involved in suppressing the inflammatory mechanism to directly enhance the health of the host. Furthermore, administration of limonoids or Spm improved the proportions of beneficial bacterial in the intestinal flora and associated metabolites. In the healthy intestinal tract, the microbiota and gut-associated immune system are assumed to be at a dynamic homeostatic equilibrium [85], but the inflammation process may undermine this balance. We consider that the human lifespan can be extended by inhibiting inflammation via control of the intestinal microbiota.

However, it was not possible to elucidate the mechanisms underlying the effects of limonoid aglycones and Spm on the extended life span of SD mice. Thus, in order to clarify the anti-inflammatory effects of yuzu seed extract, limonoids, and Spm, as well as to widely apply yuzu to promote health and enhance longevity, it will be necessary to determine the composition of the bacterial flora based on detailed metagenomic analyses of 16S rRNA. Furthermore, it will be necessary to analyze the anti-inflammatory effects of limonoids and Spm in yuzu seed extracts at the gene level.

PAs quantities have reference values. The reference values for the PAs contents of legumes are also shown in **Table 3**. The values for Japanese produced yuzu and lemons are shown, as well as the reference values for other citrus fruits. No previous studies reported the quantities of arginine in citrus fruits, so only the compared PA quantities are indicated by reference values. The reference values for the PA contents of legumes are also shown.

7. Conclusions

Yuzu is a natural and renewable resource of limonoids, arginine, and PAs. The results of the present study suggest that yuzu limonoids and Spm improved the proportions of beneficial bacteria and their metabolites in the intestinal flora. Thus, the ingestion of fruits that contain high concentrations of specific ingredients may be a simple method to suppress inflammation, thereby enhancing immune function, improving intestinal health, and increasing lifespan. In other words, our this study demonstrated the possibility that bilateral signals between the brain and intestine are important for maintaining homeostasis and extending lifespan. However, it was not possible to examine the physiological effects of limonoids and Spm. Thus, future studies are needed to evaluate the effects of limonoids and Spm on metabolism and the immune response, and to explore the potential of these molecules as natural antioxidants/antibiotics for lysosomal diseases, such as SD.

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Conflict of interest

The authors declare no conflict of interest.

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Chapter 13

Nondestructive Assessment of Citrus Fruit Quality and Ripening by Visible–Near Infrared Reflectance Spectroscopy

Ana M. Cavaco, Dário Passos, Rosa M. Pires, Maria D. Antunes and Rui Guerra

Abstract

As non-climacteric, citrus fruit are only harvested at their optimal edible ripening stage. The usual approach followed by producers and packinghouses to establish the internal quality and ripening of citrus fruit is to collect fruit sets throughout ripening and use them to determine the quality attributes (QA) by standard and, in many cases, destructive and time-consuming methods. However, due to the large variability within and between orchards, the number of measured fruits is seldom statistically representative of the batch, resulting in a fallible assessment of their internal QA (IQA) and a weak traceability in the citrus supply chain. Visible/near-infrared reflectance spectroscopy (Vis-NIRS) is a nondestructive method that addresses this problem, and has proved to predict many IQA of a wide number of fruit including citrus. Yet, its application on a daily basis is not straightforward, and there are still several questions to address by researchers in order to implement it routinely in the crop supply chain. This chapter reviews the application of Vis–NIRS in the assessment of the quality and ripening of citrus fruit, and makes a critical evaluation on the technique's limiting issues that need further attention by researchers.

Keywords: nondestructive, Vis-NIRS, citrus fruit, quality, ripening

1. Introduction

Citrus fruit are grown commercially in more than 50 countries around the world and are major commodities in the international trade [1, 2]. In Europe, the exceptional characteristics met by some of these produces have granted them the Protected Geographical Indication (PGI), such as the lemons (*Citrus limon* (L.) Osbeck) of Menton in France, Sorrento, Amalfi and Syracuse, and the Sicilian blood orange (*Citrus* × *sinensis*) in Italy, the "Algarve Citrus" in Portugal, or the "Valencianos Citrus" in Spain.

As non-climacteric, citrus fruit are only harvested at their optimal edible ripening stage, and are required to meet the expectations of the current consumer who demands for fruit not only with the best appearance, flavor, and nutritional properties, but that also comply with safety, traceability, and the sustainability of the cultural practices used. Like any other commodity, citrus fruit are subjected to worldwide standard specifications within the value chain [3] on their quality attributes (QA). Additionally, there are also adjustments to these requirements on quality and commercial ripening indices, that arise from the respective PGI normative of each commodity, growing regions and destination markets [4]. The main external quality attributes (EQA) accounted for citrus fruit are general appearance, size, weight, and color. Among the internal quality attributes (IQA), soluble solids content (SSC), titratable acidity (TA), juiciness, maturity index (MI; MI = SSC/TA), and the absence of internal defects are the most relevant. Although firmness is not defined quantitatively, it represents an important IQA, since it is a limiting factor regarding postharvest handling, transport and shelf-life, fruit being expected to maintain a good consistency through the whole supply chain.

Once fruit attain the expected IQA, additional factors will condition the harvest of citrus fruit: orchard yield and size, ripening variability, harvest cost, storage conditions, market prices and consumers' demand. Although dependent on the country, producers are provided with three options to handle these constrains: (i) immediate harvest and marketing; (ii) immediate harvest and cold-storage; or (iii) delayed fruit harvest. Opting for immediate harvest may result in minimal organoleptic quality and low prices, whereas postponing it until favorable market conditions, risks fruit drop, decay and spoilage caused by extreme weather events, pests and diseases [5]. To prevent some of of these consequences, producers resort to the regular use of pesticides, which increase the production costs and impact negatively the environment [6, 7]. Cold-storage is used in some of the major citrus producing countries, such as Spain or South Africa, and require very strict conditions to avoid fruit loss caused by chilling and/or freezing injury [8]. Both, cold-storage and harvest delay may lead to adverse alterations in the citrus-like flavor, and thus fruit quality deterioration, even if MI or SSC remains acceptable for marketing [9]. In all cases, fruit become more susceptible to the occurrence of physiological disorders that cause internal and/or external defects. Among the most typical physiological disorders registered through the supply chain of citrus fruit, there is the section drying, the rind breaking disorder (RBD), the rind pitting disorder (RP), freezing damage, and granulation, as reported for tangerine (*Citrus tangerine* Tanaka [10], 'Nules Clementine' mandarin (*Citrus* \times *clementina*) [11], 'Marsh' grapefruit (*Citrus* \times paradisi Macfad.) [12], sweet lemons (*Citrus limettioides* Tan.) [13], and 'Honey' pomelo (Citrus maxima Merr.) [14], respectively. These disorders are difficult to sort out by visual inspection at harvest, but lead to posterior fruit deterioration, limiting their quality, shelf-life, price and acceptance by consumers. In fact, there are strict standards for fruit sorting and grading, which require the detection of some of these disorders, throughout the supply chain, as established by the California Department of Food and Agriculture (CDFA). For exemple, it is not permitted to sell oranges (Citrus sinensis (L.) Osbeck) if, generally, more than 15% of fruits per batch have considerable freezing damage [15].

Therefore, the ripening of citrus fruit at harvest is a major determinant of their final quality after the whole postharvest handling processes, the occurrence of storage disorders, and the produce shelf-life span [16]. It also affects the rate of fruit loss between the tree and the consumers' home. Thus, the management and the decision capacity of the optimal harvest date (OHD) is a critical step in the supply chain. The current approach followed by producers and packinghouses to establish it and therefore, to decide on the harvest, is to collect small fruit sets from the various orchards by the beginning of each variety harvest season, and to use them to determine QA through standard methods, that in most cases are destructive, subjective and very time-consuming.

However, all QA vary greatly inside the same orchard, either in terms of absolute values and/or in terms of spatial and temporal distribution, and even in the same tree. This has been shown in citrus orchards of 'Shiranuhi' mandarin (C. unshiu \times C. sinensis) \times C. reticulata [17], 'Ortanique' (Citrus reticulata Blanco x Citrus sinensis (L) Osbeck) [18], mandarin (Citrus reticulata Blanco) [19], and 'Newhall' and 'Valencia Late' orange [20]. Multiple factors, such as the level of sunlight exposure and the associated fruit temperature on the tree, fruit yield and size, tree vigor and age, rootstocks, site-specific nutritional requirements and micro topographies within the orchard, are reportedly associated to this variability [21–24]. Furthermore, the location of the orchards and their edaphoclimatic conditions, as well as the cultural practices also induce variability on the fruit maturation process, leading to different levels of QA and different ripening rates observed for the same cultivar at different sites [20, 21]. Consequently, the number of tested fruits with the standard methods is seldom statistically representative of the orchard, leading to the sub-representation of the effective ripening stage of the fruit within and between orchards, which results in a limited assessment of their ripening, heterogeneous fruit quality, a deficient OHD management and a weak traceability in the citrus supply chain [25–27].

Overall, there is the need to upgrade the management and the sustainability of citrus fruit supply chain with smart and nondestructive technologies that allow a fast, objective, accurate and extensive assessment of fruit QA and ripening on-tree and in the following postharvest, to replace conventional methods. Their aim would be to deliver the best produce to the markets, and contribute to reduce the current level of food loss around the globe, that involves a large portion of fruit and vegetables [28–30]. Considering how much of the world's population lacks food security, and the importance of these commodities in the provision of essential nutrients and vitamins, which could prevent malnutrition, that kind of technologies would comply with the sustainable development goals (SDGs) proposed by the Food and Agriculture Organization (FAO), International Fund for Agricultural Development (IFAD), and the World Food Programme (WFP), in the 2030 Sustainable Development Agenda, which supports a global commitment to end poverty, hunger and malnutrition by 2030, creating a #ZeroHunger world [31, 32].

The large number of reports published in the past two decades, show an active, and highly motivated research concerning the development of various nondestructive technologies for the assessment of quality and ripening parameters of a wide variety of fruit, including citrus [16, 33–37]. These techniques are used on inline sorting systems, on the bench or in the field and come in many forms, prices and commercial brands. Among them, the visible–near infrared reflectance spectroscopy (Vis–NIRS), is conceivably one of the most suitable and advanced nondestructive technologies currently used to monitoring several horticultural produces. It has been implemented in applications ranging from the inline automated grading systems, assessing up to 10–12 fruit per second, to handheld units suitable for field use, operating in full sunlight and varying ambient temperature [38, 39]. Additionally, it continues to grow stronger as a major investigation topic worldwide, with a major potential for improvement and contribution to the state of the art of precision agriculture and agronomic systems management [40].

This chapter comprises a brief explanation of Vis-NIRS fundamentals and a review of the various reports on its application published since 2012. Reports published before 2012 were already covered in the last review by [41] and will not be repeated here, with a few exceptions that represent relevant breakthroughs in the area. It will further attempt a critical evaluation on the limiting issues that need further research, to implement it as an effective nondestructive method to assess these commodities' quality and optimal ripening.

The authors invite the reader to complement this chapter with some of the most outstanding reviews published throughout the years, by the main researchers working on the subject (but not only in citrus). These reviews comprise the principles of the technique, its various methods and the listing of fruit and the respective QA for which it has provided calibration models [41–45], the overview on the publications and main research groups in the field [40], various recommendations for future research activity in the area regarding the adequate experimental design and the reporting requirements [38], as well as the current real-life applications available on the market that seem to comply with the warranted robustness for the technology to be integrated in the supply chain of many crops, including citrus [38, 39].

2. Fundamentals of visible-near infrared reflectance spectroscopy (Vis-NIRS)

In this review we will adopt the most common definition that Vis–NIRS covers the wavelength range 400–2500 nm of the electromagnetic spectrum. The lower limit is consensual, since it is the onset of the visible range, but the upper limit is mainly defined by the spectral response of the most common spectrometers. It comprises the visible (Vis) region (400–750 nm), the more penetrative short wave NIR (SWNIR), or Herschel region (750–1100 nm), and the near infrared region (750–2500 nm) of the spectrum [38, 39, 45]. The NIR radiation was discovered by Friedrich Wilhelm Herschel in 1800, and was first used in agricultural applications to measure the moisture in grain in the late 1960s [45]. The first Vis–NIRS application was commercialized in Japan in 1989 to sort peaches based on SSC in an automated grading line, but the research on its principles, applications and on the development of new customized systems, have only followed some decades later, being quite active nowadays [38–40].

2.1 Interaction of radiation with the fruit

When a light beam from the sun or a tungsten lamp, hits a fruit or any other sample, the incident radiation may be specularly reflected, absorbed or transmitted, and the relative contribution of each phenomenon depends on the chemical constitution and physical parameters of the sample (Figure 1) [46]. The spectral distribution of the radiation that penetrates the product change through wavelength dependent scattering and absorption processes. The photons that enter the fruit may emerge through multiple scattering in the tissue. Light emerging on the same side of incidence is described as *diffuse reflection*, while light emerging on the opposite side is described as *diffuse transmition*. Both diffuse modes may be understood in a general sense as 'transmitted', according to the initial description. The emerging diffuse light is collected by a spectrometer, originating the term *diffuse* reflection spectroscopy. The spectral features depend on the chemical composition of the product, as well as on its light scattering properties which are related to the sample microstructure. Fruit and vegetables are turbid media, in which scattering events dominate over absorption in the visible (400–750 nm), and particularly in the SWNIR and NIR ranges of the electromagnetic spectrum (750–2500 nm) [44] (see Figure 1).

In thin rind fruit most of light interaction takes place on the flesh and the skin has mainly a modulation effect upon the spectra. In most citrus, however, most of light interaction occurs in the thick rind and few photons probe the flesh. Thus, the



Figure 1.

When light hits a fruit, the photons may be reflected at the fruit surface (specular reflection) or enter the fruit tissue. In the latter case, a succession of scattering events takes place, where the photons change direction. Some of them reemerge, originating diffuse reflection (or transmission, not represented), and the other are eventually absorbed.

assessment of IQA depends on the interplay between pulp and skin biochemistry and their optical properties [47–49].

Absolute quantification of diffuse reflected light (for example, as spectral radiance $[W \cdot sr^{-1} m^{-2} Hz^{-1}]$) is of little use, because it depends obviously on the characteristics of the light source. The calculation of reflectance avoids that subjectivity, since it normalizes the absolute measurement of the sample's reflection by that of a reference material, usually a near perfect reflector ('white') in the wavelength range under study. It should be stressed, however, that even the reflectance depends on the collection geometry (solid angle of collection, viewed area, etc.). Common choices for the reference material include Spectralon or Teflon, with nearly 100% reflection in the Vis-NIR. Reflectance $R(\lambda)$ is calculated according to the Eq. (1) presented below.

$$R(\lambda) = \frac{S(\lambda) - D(\lambda)}{Ref(\lambda) - D(\lambda)} \times 100,$$
(1)

where *S* stands for the Sample counts, *D* for the Dark counts, *Ref* for the Reference counts and λ is the wavelength. Here, *counts* refer to the digitized output of the spectrometer, which are proportional to the spectral radiance measured in a specific geometry. The dark counts are obtained with the spectrometer closed and represent the electronic noise, which must be subtracted from the sample and reference measurements.

2.2 'Point' and imaging measurements

Vis-NIRS is most commonly applied on specific 'points' of the fruit, by observing a small area, which produce an average spectrum for that specific site. This is called *point measurements*. But Vis-NIRS can also be applied on extensive sections across the fruit, through *multispectral* and *hyperspectral* measurements, which create an image of the measured sections for each wavelength band [44]. The main difference between multi- and hyperspectral modes are the number of wavebands used. Multispectral imaging uses a set of filters and a common digital camera to deliver typically no more than ten bands, while hyperspectral cameras merge imaging and spectral separation in the optical hardware to produce hundreds of contiguous wavebands. Another way to look into hyperspectral images is to think that it yields the reflectance spectrum for each spatial position of a sample (*i.e.*, for each pixel of the image) [44]. Both techniques, although costly, have been shown to successfully assess several IQA, diseases and defects in several fruit, including citrus fruit [50, 51]. Yet, extensive investigation is needed to allow both the acquisition and image processing software to be implemented in real-time systems. Thus, this chapter will only address the systems that perform 'point' measurements, based on their much wider spread, cost-effective and friendly use through the supply chain, and particularly under field conditions. Further information on the principles and applications of multispectral and hyperspectral Vis-NIRS technologies could be found in the reviews by [44, 50].

2.3 Instrumentation and measurement setup

There are currently, a large variety of both commercial and lab-made customized Vis-NIRS systems, with various shapes, sizes, and prices, that operate in many spectral ranges, and have reportedly allowed the assessment of several QA, the content of critical compounds and the diagnostic and/or prediction of disorders in a wide sort of fruits, including citrus (**Tables 2–4**). Nevertheless, most of the current commercial fruit applications of Vis-NIRS are based on the use of silicon-based spectrophotometers comprising the Vis-SWNIR region (400–1100 nm), because of their accessible prices and the larger light penetration depth in this band, in comparison to the significantly more expensive InGaAs-based devices (900–2500 nm), that do not add too much value to the quality assessment procedure [39].

From handheld, benchtop, to inline automated grading system, all Vis-NIRS devices comprehend the following fundamental components: an optical spectrometer, a light source (usually a tungsten halogen light bulb) and collection optics (optical fibers, lenses, integration spheres, dedicated probes). The current spectrometers typically include a connection for an optical fiber, an entrance slit (that defines the spectral resolution), a diffraction grating to separate the light into its spectral components, mirrors for collimation and focusing, and a light sensing device that is usually a one-dimensional CCD (Charge-Coupled Device) or CMOS (Complementary Metal-Oxide Semiconductor).

The Vis-NIR spectra may be acquired according to three principal geometrical configurations, as depicted in **Figure 2**: the reflectance mode (a), the transmittance mode (b), and in the interactance mode (c). The reflectance mode is susceptible to receive specularly reflected light, which may be a disadvantage, since only a fraction of the collected photons probes the fruit interior. In the transmittance mode the photons probe necessarily the fruit interior; however, the optical signal may be weak and noisy. The interactance mode is a tradeoff between the two previous modes: by using a contact probe it avoids specularly reflected photons and receives only those traveling through the fruit flesh. Also, the distance between light injection and collection is small, insuring a good optical signal. However, this is also a disadvantage, since the probing depth into the fruit pulp is shallow.

The choice of the geometry is thus of the utmost importance for obtaining good results, and should account for the fruit and the assessed QA. The penetration of NIR radiation into fruit tissue decreases exponentially with the depth, which is quite critical in thick rind fruit such as citrus [50]. Furthermore, the choice of the detection mode might be influenced by the spectral range used, as report by [48], in which both interactance and reflectance modes produced similar models to assess the SSC of 'Sunkist' navel oranges in the Vis-SWNIR range, but the participation of Vis region degraded this assessment in the transmittance mode. in general, to detect



Figure 2.

Setup for the acquisition of Vis-NIR reflectance spectra in (A) reflectance, (B) transmittance, and (C) interactance modes. Based on [45].

the internal defects, the transmittance mode should be chosen, while the other two modes are quite reliable regarding other IQA (**Tables 2–4**).

2.4 Typical Vis-NIRS spectrum and its interpretation

All studies on the use of Vis-NIRS to assess the fruit QA start by acquiring the reflectance (R) spectra, which is then converted to the respective absorbance log (1/R) spectra (**Figure 3**). The main spectral differences observed in a wide variety of fruit are in the visible region, namely in the 400–750 nm range. This is due to changes in the pigments' content through ripening, namely chlorophylls, carotenoids and anthocyanins present on the fruit rind [38]. In fruit that change from green



Figure 3.

(a) Average reflectance spectra of a set of 255 'Valencia Late' oranges and 239 'Rocha' pears acquired in the Vis/NIR range in the interactance mode; (b) Average absorbance spectra of the same set of fruit. The nominal positions of the most important absorption bands are indicated in the curves. The number is the order of the transition, v stands for stretching vibration, δ for bending vibrations and the sum indicates combination bands (for example, $3v+\delta(O-H)$ represents the combination band of the second overtone of stretching with the fundamental bending in O-H); (c) Savitzky–Golay [52] filter of second derivative order applied to the absorbances. The bands are again indicated; (d) to (f) Same as in (a) to (c) but in the NIR range, with the spectra acquired in reflectance mode.

to yellow/orange/red colors through ripening, the spectral information on the pigments' absorption range, may provide accessory indirect correlations with IQA such as firmness, as found in 'Rocha' pear (*Pyrus communis* L.) [53]. This, however, is not so clear in citrus fruit because their color change do not correlate with their maturity and depends on the orchards' location climate [6]. Otherwise, the pattern of the absorption spectra in the NIR range is quite similar among the various fruit species, although position and magnitude of the peaks are fruit specific, even among citrus fruit varieties [41]. The magnitude of the peaks and minima are also dependent on the acquisition mode used, but in general the same features are present and the landscape of the spectra is similar among the same fruit as reported for 'Sunkist' oranges [48].

The spectra in the NIR range convey information mainly related with vibrational bands (stretching and bending) of the relevant functional organic groups, such as O-H, C-H, C-O and C=O. The compilation of the main wavebands present in citrus fruit, namely, O-H and C-H vibration absorptions are presented graphically in Figure 3. These groups exist in all fruit organic molecules, but variations associated with water and storage reserves may induce slight changes in the spectra that may be related with the IQA. Vibration states are quantized and the transitions between states are said to be fundamental or overtones. The fundamental transitions (corresponding to a fundamental band) refer to the transition from the ground state to the first excited state, and take place mainly in the infrared range, that is above 2500 nm. In this range, the absorption peaks are distinguishable and correlate directly to specific compounds, allowing a better assessment of organic compounds such as vitamin C, citric acid or sucrose, as reported in 'Valencia' orange [54]. In contrast, the overtone bands correspond to transitions to higher excited states, with a large number falling in the NIR range. For example, the first overtone band corresponds to the transition from the ground state to the second excited state. A very crude approximation is that the n-th overtone frequency is close to (n-1) times the fundamental frequency. Thus, a general rule is that overtones have higher frequencies and lower amplitudes than the fundamental. For example, the fundamental frequency for O–H stretching (1ν) is around 2700 nm, which means that is beyond the range of the most common NIR dispersive-type spectrometers. However, the overtones are within their instrumental range: 2ν at 1420 nm (first overtone, strong intensity band), 3ν at 970 nm (2nd overtone, medium intensity) and 4ν at 750 nm (3rd overtone, low/very low intensity band). The quoted values are only indicative of a typical band central value. Indeed, the vibrations are dependent on the chemical environment, which results in a frequency spread of the bands. Finally, it is important to refer the combination bands. These correspond to the superposition of vibration motions. For example, the fundamental bending mode (1δ) of water at 6300 nm (infrared range) may combine with the fundamental stretching mode at 2700 nm (1 ν), to generate a combination band around 1900 nm $[\nu+\delta (O-H)]$. Having in mind that the fruit tissue is composed by many different organic molecules, it is easy to understand that the spectral landscape of fruit NIR reflectance is a continuum, due to band superposition, as previously shown by [54], when comparing NIR and medium infrared spectroscopies (MIR), to assess several compounds in 'Valencia' oranges. Summarizing, the NIR spectra of a fruit contains mainly overtones and combination bands of stretching and bending vibrations of the main functional organic groups of relevant organic compounds regarding the fruit IQA, such as O-H and C-H. The large number of possible vibrations and corresponding bands originates a spectral landscape with very broad and unspecific features, from which it is nevertheless possible to retrieve useful information. For instance, [54] obtained better prediction of fructose and reducing sugars when using NIRS than MIRS.

The typical Vis–NIR and NIR reflectance and absorption spectra of 'Valencia Late' orange and 'Rocha' pear are depicted in **Figure 3** [55]. Both fruit spectra were relatively flat from 700 to 910, followed by strong water absorption peaks around 970, 1450 and 1940 nm. However, the C-H bands may distort slightly the water peaks, and the analysis of this distortion conveys more information than the main peaks alone. It is from these patterns associated with the OH and CH vibrations that it is possible to retrieve the information about sugars. Even the water bands by themselves may convey information about the sugars, because the concentration of sugars and water are interdependent [41, 56–58].

In the Vis–NIR range the most prominent feature is the 3ν (O–H) peak at \sim 975 nm. This peak is reported in the literature in the range 960–980 nm [57], but the actual location depends on multiple factors: (i) the degree of OH bonding; (ii) the temperature; (iii) the presence of other close bands. In other words, within different chemical environments, the OH group will peak at different wavelengths. Effect of (iii) is more clearly observed in **Figure 3c**. Indeed, the smooth 975 nm peak observed in absorbance has a more fine structure disclosed upon derivation. Thus, the peak 4ν (CH₂) at 930 nm is actually coalescing with the main water peak, causing a depression in the 2nd derivative left positive peak. The form of this overlap is a source of information about all organic compounds content, namely sugars, acids, proteins, etc.

A minor feature, but consistently observed in most fruit, is the slight inflection around 840 nm, which is caused by the band $3\nu+\delta$ (O–H). In this case it is more clearly observable in the reflectance spectrum, **Figure 3a**. In this discussion is important to have in mind that second derivation of a symmetric peaks yields a negative peak at the same position, with two lateral smaller positive peaks. This is clearly observed for the 840 nm feature in the 2nd derivative plot, with the negative peak coinciding with the $3\nu+\delta$ (O–H) absorption wavelength. On the contrary, the peaks of the 3ν (O–H) features do not coincide in the absorbance and 2nd derivative plots, which clearly indicates spectral overlap.

Similar curves are observed in other cultivars. For example, in 'Newhall' orange [56] the same structure for the second derivative plot was observed, although a different technique was used, namely the Norris derivative [59].

Concerning the NIR spectra, those from the oranges show three main peaks at 1190, 1450 and 1940 nm, whose origin may be traced to the 3ν (C–H), 2ν (O–H) and ν + δ (O–H) bands, respectively. However, satellite bands overlap, as in the Vis/NIR case. The most 'pure' peak is the first, around 1200 nm, corresponding to the 3ν (C–H) band. As is the 840 nm band, absorbance and 2nd derivative peaks coincide. The other two main peaks are more complex blends of two or more bands. For example, the second peak around 1450 nm, although dominated by the stronger 2ν (O–H) band, has contributions from 2ν + δ (C–H) and 2ν + 2δ (C–H). Consequently, the 2nd derivative feature associated with this mix is more complex. The same could be said about the third peak.

Furthermore, due to several causes, the various peaks, even when they coincide among different fruit, may present different levels of importance, signified by their infrared values, with the various IQA. For instance, the combination band of OH reported at 839 nm correlated highly with SSC in 'Rocha' pear samples, but not in 'Valencia Late' oranges [55].

3. Chemometrics

As it has been mentioned earlier, Vis–NIR spectra of fresh fruits tend to be composed by a large superposition of absorption bands. The presence of a substantial amount of water in fresh fruit has a big impact in the spectra, dominating most of the spectral landscape. Therefore, the signals corresponding to the absorption bands of key chemical compounds such as sugars and acids, become masked by water and are only discernible as weak fluctuations in the spectrum. Given the complex interplay between the multiple absorption bands and their weak amplitudes, most of the times the linear relation between chemical compound concentration and absorption (Lambert-Beer law) is almost lost. In order to be able to extract information about chemical concentrations from this type of spectrum, we have to look for relationships' patterns between different wavelengths. This is done resorting to multivariate statistical techniques, that when applied in the field of analytical chemistry is called Chemometrics. This research field can be considered as a subset of the broader area of Machine Learning, and pursues the same goals, i.e., infer critical information from high dimensional data. There is a vast literature on this subject for those who wish to learn more about Chemometrics. Here are some suggestions for introductory and advanced levels [60–64]. In this section, it is presented a brief introduction to the scientific language used in this area, with the sole objective of familiarize the reader with the main concepts that are often presented in the literature. In Vis-NIRS of fresh fruit, the input data are spectra, i.e. one-dimensional arrays of values, each one corresponding to the intensity of light (diffusively reflected or absorbed) at a specific wavelength. Each spectrum X_n (n = 1, ... number of samples) represents a measurement or sample and each point x_i (i = 1, ..., number of variables) of the spectrum is usually referred to as input variables, spectral features or simply as 'wavelengths'. The macroscopic properties or QA features that are obtained through laboratory testing (e.g. SSC, firmness, etc.) are commonly defined as target variables Y_n or simply attributes. Chemometrics consists on the application of mathematical/statistical methods that allow mapping the spectral features x_i into the target variables Y. These methods can be subdivided into two broad subcategories: unsupervised and supervised. In the former type of method, only the input variables x_i are used and, the main purpose is to find, for example, trends within the data, clusters that can be used for classification or other general characteristics of the data set. On the other hand, supervised methods use both input and target variables and can be used for classification tasks (e.g. discriminate between different fruit sub-species or origins) or for quantitative (regression) prediction of attributes that have continuous distributions (e.g. SSC, firmness, TA, etc.).

3.1 Spectral pre-processing and outlier detection

In order to implement the best calibration model possible to predict the expected QA, often the spectral data has to be preprocessed before being used. Preprocessing techniques are used to remove irrelevant information (noise, systematic errors and faulty samples) that can degrade the performance of the numerical algorithm used to develop the calibration model. Several preprocessing methods have been created for this purpose and reviews, such as [61, 65], present a wider scope of these techniques. A brief summary of the most commonly used methods is presented in **Table 1**. For Vis-NIRS, the most common forms of spectral preprocessing can be stacked into two groups: scatter corrections (SC) and derivative techniques (DT). Scatter-corrective methods are used to remove the influence of scattered light that can contaminate the diffuse reflectance spectra. The rationale behind SC techniques is to remove effects that are unrelated to the chemical composition of the samples and that just depend on the measurement geometry or samples morphology. On the other hand, DT are designed to improve signal to noise ratios, eliminate systematic baseline biases and enhance spectral variations. Another type of preprocessing

Technique	Description
MSC (Multiplicative Scattering Correction) [66]	Each spectrum X_n is regressed against a reference spectrum (usually the mean spectrum, X_m) using a least square method. Then, the corrected spectrum is computed based on these regression coefficients. There are more sophisticated variants of this method (e.g. Extended MSC [67] that try to correct for additional additive effects).
SNV (Standard Normal Variate) [68]	Each individual spectrum X_n is normalized to have zero mean and unit variance. In some cases, this can also appear as sample standardization.
Smooth derivatives (Savitzky–Golay (SG) [52] and Norris-Williams (NW) [59]	The spectrum derivative in relation to the wavelength is computed. Usually the first derivatives remove baseline effects and the second derivative remove baseline and linear trends in the spectrum. Smooth derivatives are computed using an averaging multi point window in order to be more robust against spurious noise. SG and NW algorithms are the most common methods to compute these derivatives.
De-trending	This process consists in fitting a polynomial (usually 1st or 2nd order) to the spectrum and subtract it from the signal. This provides baseline correction.
Scaling	Scaling is a sort of umbrella under with we can find multiple types of data manipulations. The most common ones are: baseline subtraction, sub-sampling, normalization on columns (all spectral features in the data set are scaled between a max and min values) and normalization on rows (the individual spectral features are scaled between min and max).

Table 1.

Most common spectral preprocessing techniques.

commonly mentioned in the literature is outlier detection. This process consists in identifying and removing from the data set, samples that are very different from the rest of the samples. These outliers can be for example, reflectance spectra that were defectively acquired or fruit with odd properties. The idea is to remove these samples from the data set, using some pre-defined metric in order to feed the model only with the most representative samples in the data set that lead to a correct mapping of the attribute being predicted by the calibration model constructed.

3.2 Clustering

In Vis–NIR spectra of fresh fruit, the common spectrum is often described by smooth mounds and soft depressions. This means that adjacent wavelengths can be highly correlated. Therefore, in order to reduce redundancy of information provided by neighbour features $[x_i, x_{i+1}, x_{i+2}, ...]$ (also called co-linearity), sometimes it is beneficial to restrict the number of used input variables. This is often called dimensionality reduction and is very important for the right operation of certain calibration models. The simplest way to deal with this problem is by using subsampling, where a certain number of spectral features are discarded, e.g. every 3rd or 5th point in the spectra. To deal with this problem of dimensionality reduction, some 'clever' algorithms were introduced, the most common being the Principal Component Analysis (PCA) algorithm, the Hierarchical Clustering Analysis (HCA) and K-Means. the latest two methods can be used for classification tasks (mapping a

cluster to a class) and for outlier detection as well. If samples are too far apart from the defined clusters (according to some metric such as the Euclidean distance or the Mahalanobis distance), then this suggests that it might be an outlier.

3.3 Classification and regression models

As we mentioned earlier, depending on the problem at hand, we might need to implement a classification or a regression model for our data. Multiple Linear Regression (MLR) is perhaps one of the most straight forward methods to implement. It expands the application of simple linear regression to the multivariate case by linearly combining them. Due to its simplicity, this method has some drawbacks, namely an inefficient applicability in the cases of high co-linearity in the data, and when the number of features in the data set is higher than the number of samples. This is often the case of fresh fruit Vis–NIRS datasets and hence its applicability has been limited. One way of overcoming these limitations, is to use a dimensionality reduction method, such as PCA and then perform MLR on these lower dimensional components. This workflow is known as Principal Component Regression (PCR). Partial Least Square Regression (PLS) is without question the most widely used method to create calibration models to predict the most QA of fresh fruit (Tables 2-4). As opposed to PCA, the PLS algorithm takes into account the covariance between input x_i and target Y_i variables. In the same spirit as PCA, PLS also projects the data into a latent space, but this time the components are defined along the direction of maximum variance between x_i and Y_i . These components are called latent variables (also named factors by some researchers), are built in order to model the target variable, and their number is what defines the quality of the PLS model. In general, a low number of latent variables usually lead to more robust predictions, but that might not always be the case. A variant of PLS named PLS Discriminant Analysis (PLS-DA) can be used to deal with classification scenarios when the target variables Y_i are not continuous (e.g. 0, 1 for fruits without and with defects). The models mentioned so far are can be described as linear because they rely on a linear combination of multivariate solutions. Besides the easiness of implementation, they are also classically appreciated in Chemometrics because they are easy to interpret in terms of feature importance, i.e., after fitting the model to the data we can back-trace some parameters (e.g. regression coefficients) and find what wavelengths or spectral bands better contributed to the prediction. In turn, this allows inferring information about the chemical concentrations and can be used to identify biological and metabolic behaviors.

In the last couple of decades, non-linear models imported from other areas of Machine Learning have begun to permeate Chemometrics, and given its high use case in the literature, Support Vector Machines (SVM) is one of the most popular. The strategy of this model consists in searching for boundaries that separate two cluster or classes. The algorithm tries to find the best boundary between classes by maximizing a distance margin between neighbor samples. It has the advantage that it can use kernel tricks to transform the data points into another mathematical space, where these boundaries are easier to establish. SVMs were initially used for classification tasks, but have been extended to deal with regression problems as well (SVR). SVR has been used successfully for many datasets, and the most often mentioned drawback is the complexity of its optimization task. Another popular type of non-linear models that is often used for classification and regression problems is Neural Networks (NN). These represent a wide class of algorithms with many types of architectures and are derived from the field of Artificial Intelligence. In recent years, classical NN architectures such as the Multi-Layer Perceptron has

IQA	Unit	Range (nm)	Mode	Stat Method	Val	RMSEP	R ²	RPD	Ref
'Marsh' g	grapefruit								
Suc	g/kg	400–2500	R	PLS	Ι	35	0.94	3.76	[69]
Gluc	mg/g					18	0.77	1.80	
TPh	gGAE/kg					0.37	0.59	1.03	
DPPH	%					5.8	0.85	2.45	
TotCar	g/kg					2.9	0.80	1.71	
VitC	g/kg	400–700				0.20	0.95	3.33	
Chl a	g/kg					0.79	0.87	2.31	
Chl b	g/kg					2.8	0.94	3.61	
'Star Rub	y' grapefruit								
SSC	%	850-2500	R	PLS	Е	0.31	0.90	2.94	[70]
ТА	g/100 mL					0.07	0.83	2.46	
MI	SSC/TA					0.45	0.81	2.32	
BrimA	SSC-TA					0.43	0.86	2.68	
'Trovita'	orange								
RateOleoc	2	400–1000	R	PLS	Ι	0.008	0.97	_	[71]
DegOleoc						0.005	0.97	_	
'Valencia	' orange								
SSC	%	900–1800	R	PLS	Е	0.58	0.69	1.87	[72]
VitC	mg/mL	1050-2000	Ι			8.01	0.43	1.33	
SSC	%	850-2500	R	PLS	Е	0.28	0.93	3.57	[70]
TA	g/100 mL					0.02	0.93	3.88	
MI	SSC/TA					0.61	0.96	4.92	
BrimA	SSC-TA					0.006	0.96	3.96	
Firm	Ν	1000–2500	R	PLS	Е	6.22 ^a	0.85	_	[73]
Pectin	%	1000–2500				5.04 ^a	0.49	_	
VitC	mg/L	1000–2500	R	PLS	Ι	95	0.50	_	[54]
CitAc	g/L					3.8	0.56	_	
Suc	g/L					12	0.56	_	
Gluc	g/L					6.7	0.67	_	
Fruct	g/L					5.0	0.66	_	
TotSug	g/L					12	0.64	_	
RedSug	g/L					20	0.67	_	
'Shatian'	pomelo								
WC	%	400–1700	Т	PLS	Ι	0.49	0.74	_	[74]
Granul						0.096	0.97	_	
'Honey' p	oomelo								
Granul		400–1700	Т	PCA-NN	Ι	$0.97 - 0.98^{b}$	_	_	[14]
sweet len	ion								
Freeze	Binary	400–1100	Т	SIMCA	Ι	> 92% ^b	_	_	[13]

IQA	Unit	Range (nm)	Mode	Stat Method	Val	RMSEP	R ²	RPD	Ref	
Injury				PCA-NN		100% ^b	_	_		
				SVM		> 92% ^b	_	_		
^a SEP values.										

^bThis value corresponds the accuracy of the classification.

Table 2.Overview of applications of Vis-NIRS to measure the quality attributes of citrus fruit by benchtop devices. List
of symbols in Table 5.

IQA	Unit	Range (nm)	Mode	Stat Method	Val	RMSEP	R ²	RPD	Ref
'Marsh' g	rapefruit								
RindPitt		400–2500	R	PLS	Е	5E-4	0.89	—	[12]
DM	%	400-850				0.30 ^c	0.88	—	
'Ehime K	ashi28' mand	larin							
SSC	%	750–1050	Ι	PLS	Ι	0.59	0.71	—	[75]
'Imperial	' mandarin								
SSC	%	720–950	Ι	MPLS	Е	$0.6 – 0.8^{d}$	0.5-0.8	1.2-3.4	[76]
DM	%					0.6 ^e	0.9 ^e	_	
SSC	%	720-950	Ι	MPLS	Е	$0.4 – 0.5^{\mathrm{f}}$	0.6-0.9	_	[77]
						0.4–0.5 ^g	0.3-0.7	—	
						$0.8-6.8^{\mathrm{h}}$	0.0-0.8	_	
'Nanfeng	' mandarin								
TA	g/100 mL	400–1040	Т	PLS	Ι	0.09	0.41	—	[78]
SSC	%					0.65	0.85	—	
VitC	mg/100 mL					2.80	0.64	—	
TA	g/100 mL			BPNN		0.09	0.42	—	
SSC	%					0.64	0.86	—	
VitC	mg/100 mL					2.70	0.64	—	
'Nules Cl	ementine' ma	andarin							
WL	g	900–1700	Ι	PLS	Ι	2.76	0.83	2.34	[11]
DMC	%					0.92	0.92	3.57	
Sucrose	mg/g DW					24	0.77	1.4	
Glucose	mg/g DW					11	0.88	2.86	
Fructose	mg/g DW					12	0.90	3.23	
TotSug	mg/g DW					31	0.90	3.15	
RBD	binary	450–1000	Ι	PLS	Е	0.36	0.37	_	[79]
Orange									
SSC	%	600–1100	R	PLS	Ι	0.73	0.35	_	[80]
		600–950				0.60	0.66		
		600-1100B		VABPLS		0.60	0.67	_	

IQA	Unit	Range (nm)	Mode	Stat Method	Val	RMSEP	R ²	RPD	Ref	
'Gannan'	orange									
SSC	%	820–950	Ι	PLS	Ι	0.40	0.77	_	[81]	
				SVM		0.32	0.84	—		
SSC	%	600–1100	Т	PLS	Ι	0.55	0.85	2.57	[82]	
		685-983ª		SPAPLS		0.57	0.84	2.51		
Navel or	ange									
SSC	%	580-970 ^b	Т	PPSOPLS	Ι	0.43	0.79	_	[83]	
'Valencia	a Late' orange	e								
SSC	%	400–1000	R	PLS	Ι	0.33	0.92	_	[84]	
TA	mg/L					0.07	0.73	—		
BrimA	SSC-TA					0.37	0.84	—		
MI	SSC/TA					1.03	0.76	—		
'Page' ta	ngelo									
TA	c.a./100 g	400–1000	R	PLS	Ι	0.03	0.77	_	[85]	
SSC	%					0.40	0.72	—		
^a Selected wavelengths in this range. ^b Selected wavelength bands in this range. ^c Cross validation. ^d Combinations of cal. and val. Sets.										

^fCombinations of cal. and val. sets (harvest day).

^gCombinations of cal. and val. sets (orchard location).

^hCombinations of cal. and val. sets (season).

Table 3.

Overview of applications of Vis-NIRS to measure the QA of citrus fruit by portable spectrometers/systems under laboratory conditions. List of symbols in Table 5.

been increasingly substituted by more modern architectures developed for Deep Learning. The most promising of these NN are the so called Convolutional Neural Networks that have been very successful in image recognition tasks.

3.4 The quality of a calibration model

Independent of the type of model that is used for prediction or classification, the important thing is to find how well it performs on the desired data. To assess the quality of the predictions made by the calibration models, several metrics are often used. In a recent review by [38], the author makes a case for the uniformization of the report of error metrics in future publications. In what follows these recommendations are highligted. The partitioning of the data for model development is very important. The data set should always be split into two sub-sets, called train and test sets. The train set, as its name suggests, is used for the calibration model development and, once the main hyper-parameters have been established, the model is used to predict the test set and assess its performance. Model development can be done with the full train set using a cross-validation strategy or by further splitting it into calibration and tuning (or assessment or validation) sets. As a note of caution, it is important to mention that for different areas of Machine Learning the names given to these data splits can vary and that can lead to some confusion. Otherwise, the test set should be derived from a different distribution from that of the train, in which case it is named as external validation set [45]. For example, data from two

IQA	Unit	Range (nm)	Mode	Stat Method	Val	RMSEP	R ²	RPD	Ref
'Clemenvil	la' mandar	in							
TA	% acid citric	1600–2400	R	MPLS	Ι	0.15	0.56	—	[86]
SSC	%					0.77 ^a	0.50	_	
MI	SSC/TA					1.46 ^a	0.53	_	
pH						0.11 ^a	0.61	_	
MPF	Ν					7.3ª	0.45	_	
PerTh	mm					0.54 ^a	0.51	_	
TA	% acid citric	1600–2400	R	LOCAL	Ι	0.13 ^a	0.76	2.08	[87]
SSC	%					0.71 ^a	0.57	1.51	
MI	SSC/TA					1.13 ^a	0.79	2.14	
pH						0.11 ^a	0.74	1.91	
BrimA	SSC-TA					0.70 ^a	0.75	2.0	
5 cultivars	of orange								
SSC	%	450–1100	Ι	PLS	Е	0.86	0.56	_	[88]
'Newhall'	orange								
SSC	%	680–1100	Ι	PLS	I/E	1/1.1 ± .2	.5/ .3 ± .1	1.4/ 1.1 ± .1	[56]
Juice pH						.2/ .27 ± .04	.5/ .4 ± 0.1	1.4/ 1.1 ± 0.1	
'Powell Su	mmer' orai	nge							
SSC	%	500–1690	R	LOCAL	Ι	0.62	0.74	_	[89]
MPF	Ν					4.64	0.74	_	
PerTh	mm	1100–1690				0.58	0.68	_	
Juice Weight	g					18	0.68	_	
SSC	%	1600–2400	R	LOCAL	Ι	0.80	0.81	2.23	[87]
MI	SSC/TA					3.56	0.65	1.69	
BrimA	SSC-TA					0.85	0.82	2.33	
'Clemenvil	la' mandar	in and 'Powe	ll Navel	' orange					
JuiceMass	g	1600–2400	R	LOCAL	Ι	0.92 ^a	0.69	1.77	[90]
PerTh	mm					28 ^a	0.72	1.86	
SSC	%	1600–2400	R	LOCAL		0.86 ^a	0.78	2.09	[87]
pН						0.15 ^a	0.72	1.93	
TA	% acid citric					0.14 ^a	0.84	2.43	
MI	SSC/TA					2.98 ^a	0.77	2.08	
BrimA	SSC-TA					0.84 ^a	0.78	2.15	
^a Corrected to bi	as.								

Table 4.Overview of applications of Vis-NIRS to measure the quality attributes of citrus fruit by portable systems
on-tree. List of symbols in Table 5.

Colums	Rows' content
IQA : Internal Quality Attributes	 β Car: β carotene; BrimA: Brix minus Acids (SSC-k(TA)); Chl a: chlorophyll a; Chl b: chlorophyll b; CitAc: Citric Acid; DegOleoc: Degree of oleocellosis; DMC: Dry Matter Content; DPPH: 2,2-diphenyl-1- picrylnhydrazyl; DM: dry matter; Firm: Firmness; Fruct: Fructose; Gluc: Glucose; Granul: Granulation; JuiceVol: Juice Volume; MPF: Maximum Penetration Force; MI: Maturation Index; Pectin: Pectin Content; PerTh: Pericarp Thickness; RateOleoc: Rate of oleocellosis; RBD: Rind Breakdown disorder; RedSug: Reduced Sugars; RindPitt: Rind Pitting; SSC: Soluble Solid Content; Suc: Sucrose; TA: Titratable Acidity; TAO: total antioxidant capacity; TotCar: Total Carotenoids; TotSug: total sugars; TPh: Total Phenolic content; VitC: Vitamin C; WC: Water Content; WL: Weight Loss
Mode: mode of acquisition	R (Reflectance), I (interactance) or T (transmittance)
StatMethod : Statistical method employed	 iPLS: Interval PLS; LAR: Least Angle Regression; LOCAL: Patented algorithm for PLS regression from a subset of proximal calibration samples; LS-SVM: Least squares support vector machines; MLR: Multiple linear regression; MPLS: Modified PLS PCA-NN: Principal components analysis combined with artificial neural networks; PCA-BPNN: back propagation neural network (BPNN) based on principal component analysis (PCA) PLS: Partial Least Squares; PPSOPLS: piecewise particle swarm optimization (PPSO) algorithm based PLS SIMCA: Soft independent modelling of class analogy; SPAPLS: Succesive Projections algorithm coupled with PLS; SVM: Support vector machines; VABPLS: variable adaptive boosting partial least squares
Val: Model validation type	I (internal), CV (cross validation only), E (external)
RMSEP: Root Mean Square	Error of Prediction
R ² : Coefficient of determina	ation
RPD : Residual predictive de	eviation

Table 5.

List of abbreviatures used in Tables 2-4.

consecutive harvest seasons are used as train set, while the test set uses data from a third season. A similar situation can be envisaged by using a train set collected from different orchards or origins, than that used for validation. Yet, given the large amount of time invested in acquiring this type of data, multi-seasonal or multiorchards test data sets are not often found in the literature. In contrast, laboratory models with homogeneous fruit sets are abundant (Tables 2-4). Currently, the most common procedure when constructing and validating Vis-NIRS models for the various fruit OA is to separate a fraction of the available samples as train/ calibration set (usually 80%), and the remaining as test/validation set (usually 20%). Furthermore, the validation samples are typically chosen as the best possible representation of the whole set and within the variation range of the train set. This has been applied even when the models comprehend several species and/or cultivars, orchard locations and harvest years, which are mixed in the calibration and validation sets [90]. All studies included in Tables 2-4 that used this approach, were labeled as internal in the validation column. Internal validation does not ensure the success of a continuous monitoring application, which is a dynamic and open process, particularly, if one aims to use the Vis-NIRS in real world applications, being in inline grading systems or handheld devices. Once the model is applied to the test set and a final prediction is made, one can assess how well the model performs by computing several metrics. If the model was developed for regression, the most used are the root mean squared error (RMSE), bias, the slope,

the coefficient of determination (R²), and the standard deviation ratio (SDR) or ratio of performance to deviation (RPD) or residual standard deviation (RSD). If the model is developed for classification, the advised metrics are accuracy (ACC), F1 score and receiver operating characteristic (ROC) curve. For completeness, these metrics are often computed not only for the test set, but also for the calibration, and tuning sets as well. The comparison between calibration (C), tuning (CV) and test set (P) error metrics allows to understand how well the model generalizes, i.e. how the information learned by the model during training transposes to the final external validation dataset.

4. Prediction of quality attributes

Vis-NIRS combined with various chemometric methods has produced calibration models to predict simultaneously multiple QA of various citrus species and varieties, which are presented in **Tables 2–4**. These attributes range from fruit size, weight and color [90] to SSC, MI, external and internal defects, several compounds such as sugars, acids, pigments and antioxidants. As expected, these models address predominantly several varieties of orange and mandarin, but also grapefruit, lime, pomelo, sweet lemon and tangelo. The spectral ranges used cover the whole Vis-NIRS range. The majority of the Vis-NIRS calibration models were obtained from samples collected and assessed under controlled conditions in the laboratory, after fruit temperature equilibration, either with benchtop or handheld devices (**Tables 2** and **3**). Despite the large market availability of the latter, presenting different levels of portability, spectral ranges, sizes, and prices, only a few studies have focused on its application to assess the quality and ripening of oranges and mandarins on-tree (Table 4), perhaps due to the complexities involved under field conditions, and the performance deterioration of calibration models, in spite of the spectral range used [86–90]. Nevertheless, the QA assessed on-tree (Table 4) comprise fruit mass and size, color parameters [86, 89, 90], pericarp thickness, SSC, TA, firmness, MI, juice pH and mass, and BrimA index, which measures the balance between sweetness and acidity as described by [12]. Noteworthy, the majority of the calibration models exhibited $R^2 < 0.8$, despite the range used, and did not include external validation, except for [56, 88].

Grading lines equipped with Vis-NIR sensors are now commercially available from various companies, to assess both the EQA and IQA of citrus fruit [38–40]. Unfortunately, the scientific evidence about the accuracy of these systems is very scarce, due to the 'industrial secrecy'. Nevertheless, there are a few cases of partnership among the industrial sector and the research groups to know the real applicability and their performance in assessing citrus fruit QA by such equipment [91], in most cases still in the prototype stage, as reported by [92, 93]. Valero [91] evaluated the performance of a customized NIR equipment installed underneath the fruit conveyor to sort oranges and mandarins in a Spanish packinghouse. This system working in a transmittance mode in the 650–970 nm spectral range, only provided calibration models that could discriminate between low and high values of SSC for both mandarin (R = 0.76-0.86; SEP ~ 0.9 °Brix; RPD ~ 0.74) and orange (R = 0.87; SEP \sim 0.7 °Brix; RPD < 1.5). No acceptable models were obtained for TA in neither species. Miller and Zude [92] evaluated the performance of a Vis-NIRS system to assess the SSC of 'Indian River' red grapefruit and 'Honey' tangerine from Florida in a sorting inline prototype, with R^2 ranging from 0.15 to 0.67. The prototype percent correct classification averaged 85% for SSC at 10 °Brix and 79% for an 11 °Brix setpoint in the second-year of tests. Otherwise, [93] reported on the development and laboratory testing of the nondestructive citrus fruit quality monitoring

prototype system, which consisted of a light detection and ranging (LIDAR) and Vis-NIRS sensors installed on an inclined conveyor for mimicking real-time fruit size and SSC measurement respectively, during harvest. Laboratory tests in 'Valencia' orange revealed that the system was applicable for instantaneous fruit size ($R^2 = 0.91$) and SSC ($R^2 = 0.677$, SEP = 0.48 °Brix) determination.

The various Vis-NIRS calibration models presented in this chapter, show different levels of accuracy, prediction and robustness for the various attributes, SSC being the most successfully IQA assessed at all spectral ranges, independently of the devices used (**Tables 2–4**). Both juice pH and vitamin C also seem easily assessed by devices operating in the Vis-SWNIRS range devices [78, 94, 95], but TA has been shown to require wavelengths range > 1000 nm [13, 96]. Additionally, calibration models for firmness have been difficult to obtain, although there are a few exceptions reported for several orange varieties, in the reflectance mode and in the ranges 500–1690 nm [89], and 1000–2500 nm [73]. Of course, the calibration models for specific compounds, such as sugars, acids or antioxidants, require in most cases the longer NIR spectral range [12, 54, 69].

Both external and internal defects in citrus have been successfully predicted by Vis-NIRS, such is the case of the section drying in tangerine (transmittance; 780 and 960 nm) [10], oleocellosis in 'Trovita' orange (reflectance; 400–1000 nm) [71], the freeze damage in sweet lemon (half-transmittance; 400–110 nm) [13], the rind pitting in 'Marsh' grapefruit (reflectance; 400–2500) [12] or the granulation in 'Shantian' pomelo (transmittance; 400–700) [74]. However, [97] was not able to predict the external disorder RBD in 'Nules Clementine' mandarin (interactance; range 450–1000 nm).

Among the chemometrics methods used to construct the calibration models, PLS is still the main linear regression technique used, and the one to produce the best models for the widest number of QA (**Tables 2–4**). However, there are some exceptions, regarding the use of non-linear techniques, which were shown to deliver calibration models with equivalent or even better prediction and accuracy for several QA than PLS. Among these, there is the WT-LSSVR, BP-NN and LS-SMV that provided models with higher prediction capacity for SS in 'Gannan' orange [81], SSC, TA and vitamin C in Nanfeng mandarin [78] and in 'Newhall' orange [96], respectively. The LOCAL algorithm has also shown to produce better models than MPLS for firmness and juice mass in 'Powell Summer Navel' orange [89], and in 'Clemenvilla' mandarin [90]. SIMCA, SVM and particularly PCA-ANN, also allowed to assess with a total accuracy > 98% the freeze damage in sweet lemon [13] and PCA-GRNN allowed the assessment of the granulation in 'Honey' pomelo at a classification accuracy (CA) > 95% [14].

Independently of the chemometrics technique used, for the majority of the models presented in **Tables 2–4**, the train and test fruit sets were chosen from the same batch, orchards or seasons. Even when the whole data set comprised fruit from several orchards, harvest season or citrus varieties, the usual approach was to randomly choose 80% of the whole data set to construct the calibration model and 20% to validate it, as reported by [90, 97]. A truly stringent external validation is thus required to have a realistic idea of the models' performance in orchard and/or cargo batch monitoring. External validation means validation through a dataset with a different origin (spatial or temporal) relatively to the datasets used in calibration. Nevertheless, there are some clear examples of this approach, such as previously reported for mandarin [76, 79, 97–99], orange [12, 56, 70, 72, 73, 79, 88, 97], and grapefruit [12, 70]. Without the effective external validation, it is not possible to know exactly how well these models would work in real conditions due to the large variability within the trees, orchards, sites and harvest seasons. Yet, a certain degree of deterioration of the initial model prediction is expected, which

would warrant further attention. This has been reported by [56, 77, 88]. Yet, there is space and potential for improvement beyond the 'proof of concept', if one aims to use these devices on the daily routines of the orchards' management. This has been suggested by [56, 78] through model recalibration using a few fruits from the new harvest season/orchard, or by achieving a strong degree of robustness by constructing a multi-seasonal and multi-orchard model as reported by [89] for 'Newhall' orange, which will be much more advantageous when assessing the ripening of fruit on-tree.

5. Future research and perspectives

Vis-NIRS has been incorporated by a large number of companies in commercial applications to be used on inline, benchtop and handheld systems. However, there are several topics regarding the full potential and limitations of this technology that require attention and further research in order to provide the consistency warranted by the daily basis routines of the citrus supply chain when assessing fruit quality and ripening. Firstly, all researchers engaged in this area should report their results in a uniform way, particularly in what respects the obtained models' metrics [38]. This would allow a better understanding on the effective advances and contributions made by each study. Other models' metrics parameters, such as the prediction gain, may also be useful, as reported by [100]. Secondly, the calibration models' robustness must be addressed and solved through a stringent multi-year, multicultivar and multi-orchard validation, such as previously reported by [56]. The usual approach of validating calibration models with a random fraction of the total available data set, even when the models comprehend several varieties, orchard sites and harvest years does not ensure the success of a continuous monitoring application and delivers unrealistic performance metrics [76, 77]. The usual recalibration and spiking approaches used to improve the initial calibration models with a few fruits from independent data sets that will be then assessed, assume that those fruits used to recalibrate/update the model constitute a faithful representation of the new population and are common techniques in various commercial devices, for inline and benchtop systems. However, this becomes quite difficult to apply if one aims to monitor the on-tree fruit ripening evolution through time, for the fruit sampled in the first weeks cannot represent those to be measured in the last weeks of the harvest season. Thirdly, there is a large potential for models' improvement, by using the non-linear techniques of machine learning, and those of deep learning. Fourthly, there is much to understand on the effect of the rind in the assessment of the pulp IQA in citrus fruit, since the NIR radiation hardly gets to the fruit pulp, and both biochemical and optical properties have a major role to play in the spectral data acquired [47, 49, 95, 101, 102]. Fifthly, the calibration models should be able to predict attributes that are closer to the organoleptic evaluation of the fruit. It is the case of BrimA index, a better indicator of fruit sweetness that the SSC, which was satisfactorily predicted in orange, grapefruit and mandarin [12, 87]. Finally, the handheld devices must really be tested under field conditions, if one aims to assess the fruit on-tree, which is essential for the OHD decision.

6. Conclusions

The usefulness of Vis-NIRS combined with different chemometric techniques in the supply chain of citrus fruit is already quite extensive and growing, similarly to many other commodities. In this chapter the authors only addressed the classic

spectral 'point' measurements, but it is quite clear that both inline, benchtop and handheld devices are used to assess nondestructively multiple QA in various citrus species and cultivars, with a clear predominance of orange and mandarin. Among these attributes, there are both EQA and IQA, as well as defects caused by various factors, such as physiological disorders. The devices available on the market are from various brands, operate in various ranges and present a wide variety of prices. Aside from the "proof of concept" made by many studies, that the authors tried to comprise as much as possible in this chapter, there are several issues that still need to be addressed by researchers, a major one being the need for a stringent external validation of the calibration models, in order to assure robustness and to fulfill with the essential requirements to include this technology in the daily routines of these crop supply chain. This is of the utmost importance when considering the assessment of fruit ripening on-tree to determine the optimal harvest date for each orchard, or sections of the orchard. This is highly significant based on the determinant effect of producing and harvesting the fruit at its best ripening stage, thus assuring the best quality throughout the whole postharvest and shelf-life. As a concluding remark, it is very important to add that these devices are of medium and high cost, and that are not the kind of technology to 'set and forget', as reiterated by [39], which demands not only for a budget to acquire the systems, but also to maintain them, and to keep the continuous update and improvement of the calibration models, that in most cases need the selling company assistance. Thus, there must be a cost-benefit that both the producers and packinghouses have to meet through the added commercial value to citrus fruit by these systems, and the consumer willingness to pay for fresh fruit graded in terms of IQA such as sugar content, acidity and nutraceutical properties.

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Conflict of interest

There are no conflicts of interest.

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Chapter 14

Electrochemical Applications for the Antioxidant Sensing in Food Samples Such as Citrus and Its Derivatives, Soft Drinks, Supplementary Food and Nutrients

Ersin Demir, Hülya Silah and Nida Aydogdu

Abstract

Although there are many definitions of antioxidants, the most general description; antioxidants are carried a phenolic function in their structure and prevent the formation of free radicals or intercept from damage to the cell by scavenging existing radicals. Moreover, they are one of the most effective substances that contain essential nutrients for healthy individuals. The importance of these antioxidants, which have an incredible effect on the body and increase the body's resistance, is increasing day by day for healthy individuals. Numerous studies have been carried out for antioxidants with excellent properties and however new, reliable, selective, sensitive and green analytical methods are sought for their determination at trace levels in food samples. Along with the latest developments, electrochemical methods are of great interest in the world of science because they are fast, reliable, sensitive and environmentally friendly. Electrochemical methods have been frequently applied to analyze antioxidant capacity in many nutrients samples found in different forms such as solid, liquid without any pretreatment applications in the last decade. Furthermore, these methods are preferred because of the short analysis time, the ability to lower detection limits, reduction in a solvent, high sensitivity, portability, low sample consumption, wide working range, and more economical than existing other traditional analytical methods. The antioxidant sensing applications by modern electrochemical methods such as cyclic, square wave, differential pulse, and combined with stripping voltammetric techniques were used to deduce antioxidant capacity (AC) in critical nutrients. Moreover, this chapter includes a description of the classification of electrochemical methods according to the working electrode type, dynamic working range, limit of determination (LOD), limit of quantification (LOQ), sample type, and using standard analyte and so forth for each voltammetric methods. While many articles applied for the determination of antioxidant sensing by electrochemistry have gained momentum in the last two decades, we focused on the studies conducted over the last 4 years in this chapter.

Keywords: antioxidant determination, electrochemistry, voltammetric methods, potentiometry, amperometry

1. Introduction

Free radicals occur when an atom or molecule contains one or more unpaired electrons in its outermost orbitals [1]. Basically, three main factors play a role in the formation of free radicals. i) The atoms or molecules can become radical as a result of the fragmentation of covalently bonded molecules exposed to high-energy electromagnetic waves or high temperatures. ii) A molecule that does not have a radical feature experience an electron loss and radicals are formed by leaving unpaired electrons in its outer orbital. iii) A radical is formed when a molecule that does not have a radical property receives an electron from outside and has an unpaired electron in its outer orbital [1, 2]. These unshared electrons as known radicals are highly unstable, transforming them into high-energy and very efficient chemical species. The most active free radicals in biological systems are those based on oxygen and are commonly referred to as reactive oxygen species (ROS) with pathological [3]. This family group includes superoxide radical (O_2) , singlet oxygen, nitroxide (NO), hydroxyl radical (OH[']), and hydrogen peroxide (H₂O₂) which is not itself radical but causes the formation of radical [1]. Besides, we can classify the causes of free radicals in two groups as endogenous or exogenous [1, 2]. Cigarettes, air pollution, alcohol, radiation, heavy organic solvents and pesticides are among exogenous sources, while enzymes, proteins, oxidative stressors, and heavy metals are endogenous sources [1, 4].

Free radicals cause the greatest damage to human health on basic cellular components such as lipids, proteins and nucleic acids [1, 5]. Therefore, these radicals lead to immune deficiency, hypertension and even important diseases such as cancer, neurodegenerative diseases, heart disease, and atherosclerosis [1, 2]. Also, studies are revealing that radicals disturb the homeostatic balance [6]. To scavenge these drawbacks effects of radicals, which are extremely important for human health, the human body needs antioxidants obtained from the body or nutrition to fulfill biological activities such as survival and healthy life. Antioxidants can be defined as molecules that usually contain phenolic functional groups in their structure and prevent the formation of free radicals that damage the cell or by scavenging existing radicals [3]. The functional task of antioxidants is that they act as shields in the body and neutralize them by donating their electrons with the s-free radicals. Thus, radicals found in a rather unstable structure do not become a threat to human health by transforming into a more stable structure reacted with antioxidants. Moreover, many different equivalent antioxidant expressions are used in antioxidant quantification in food samples. The leading ones are the expressions of "total antioxidant capacity (TAC)", "antioxidant activity (AA)", and "antioxidant capacity (AC)". The total amount of antioxidants is expressed by measurement units such as equivalent trolox, rutin, ascorbic acid, and quercetin, etc.

Antioxidants are mainly obtained via natural and synthetic [7]. The first of these, natural antioxidants, are molecules synthesized by the organism or obtained from food sources. Natural antioxidants produced by the organism are the most important source for human health. Many factors affect the production process of this natural antioxidant. The most important of these is the age of the person. As a person gets older, the amount of natural antioxidants produced by his organism decreases day by day. For this reason, there is a greater need for the natural antioxidants found in foods for older people. The importance of healthy food sources, especially organic-based foods, is increasing day by day. Also, such nutrients should be accessible to all segments of society.

Important dietary flavonoid sources are fruits especially citrus fruits such as oranges, apples, grapes, mandarins, berries lemons, limes and their derived products as well as juices [8]. In general, citrus fruits contain pectin, sugar, carotenoid

pigments, vitamins (A, B1, and C), and; organic acids such as ascorbic acid and citric acid, minerals and a number of active phytochemicals such as flavonoids and coumarins, as naringenin, naringin, hesperidin, neohesperidin, hesperetin, rutin, narirutin and tangeretin [9]. For example; polyphenol antioxidants such as flavanols (epicatechin, catechin), phenolic acids (caffeic acid and gallic acid), anthocyanins (e.g., malvidin-3-glucoside), oligomeric and polymeric proanthocyanidins, flavonols (myricetin, quercetin, and their glycosides), and many others polyphenols exist in wine, especially in red wine [10]. Flavonoids have an important role in scavenging reactive oxygen species, which can counteract lipid oxidation, decrease peroxide formation in vivo, and improve activity of the body's antioxidant enzyme. Citrus flavonoids such as naringin, naringenin, and hesperidin have antioxidant activity [11]. Naringenin is a flavonoid, particularly a flavanone, found in citrus fruits especially oranges and grape fruits and in vegetable's such as tomatoes and their preparations. The pharmacological and biological properties of phytoestrogen naringenin and its derivatives include, anticancer, anti-inflammatory, antiulcer, antifibrotic, diastolic, antioxidant and skin protective effects [8]. Also, citrus species are a rich source of flavanone glycosides such as hesperidin and narirutin, which have anticancer, antioxidant, antiobesity and anti-inflammatory activities [12].

Secondly, the antioxidant group is synthetic, that is a molecule that is obtained as a result of chemical reactions and is generally used as food preservatives [13]. Synthetic antioxidants such as butylated hydroxytoluene (BHT), butylated hydroxyanisole (BHA), and tertiary butylhydroxyquinone (TBHQ) also extend the shelf life of foods [14]. However, natural antioxidants that can be taken from foods are less risky in terms of human health since synthetic antioxidants can have toxicity even if they are very little, they require high costs and have less capacity than natural antioxidants. Due to this reason, the investigations of foods types that can contain high levels of antioxidants in different types of endemic, organic and traditional food samples have been remarkably increased recently.

For antioxidant content and amount analyzes, oxygen radical absorbance capacity (ORAC) and radical-arrest antioxidant parameter (TRAP), ferric thiocyanate (FTC), Trolox equivalent antioxidant capacity (ABTS/TEAC), cupric ion (Cu^{2+}) reduction antioxidant capacity (CUPRAC), iron ion reducing antioxidant capacity (FRAP), DPPH radical scavenging activity determination and Folin–Ciocalteu methods are the most widely preferred as analytical methods [15–17]. Furthermore, to evaluate and characterize the antioxidant substances in food samples, various analytical methods such as high-pressure liquid chromatography (HPLC) combined with different detection, gas chromatography, micellar electrokinetic capillary chromatography, capillary electrophoresis includes different detection systems and UV-visible spectrophotometry have been used [18–20]. However, these classical methods have great shortcomings for fully validated analyzes such as long pretreatment, need for too much solvent, expensive equipment, long analysis time. They do not provide the necessary procedures for green chemistry, especially due to the use of too much solvent and too much waste in antioxidant analyses. For these reasons, scientists have turned to alternative methods for antioxidant quantification in food samples. Especially in recent years, they have focused on electrochemical techniques which are fast, inexpensive, reliable, non-pre-treatment, and environmentally friendly in the analysis of drugs, pesticides, metal ions and organic molecules such as antioxidants, vitamins and nucleic acid [21-23].

In this chapter, the applicability, sensitivity and reliable maintenance of electrochemical methods, which have attracted great attention in food and food samples, have been examined for the analysis of antioxidants. Moreover, which types of electrochemical methods are used and what advantages they provide have been investigated for the antioxidant sensing in food samples. It also describes the classification of each used in electrochemical methods by working electrode type, dynamic operating range, the limit of detection (LOD), measurement limit (LOQ), sample type, and standard analyte, etc. While many articles referenced for determining antioxidants by electrochemistry have gained momentum in the literature in the last two decades, we focused our study on the studies conducted in the last 4 years.

2. Electrochemistry

Electrochemistry is the branch of science which is investigating the physical and chemical changes coming from the interaction of the material with electrical factors such as current, potential, and electron charge. Electroanalytical chemistry is based on measuring the electrical properties of solutions containing analytes and switching to quantification using measured electrical signals a collection of electrochemical methods. Moreover, electroanalytical measurement methods are based on two basic points: potentiometric (static methods) and potentiostatic (dynamic methods). Electrode systems in both methods are immersed in the solution containing the analyte, called the electrochemical cell. Potentiostatic methods are widely used for routine analysis because they are less costly, high sensitive, and selective and have wider potential application areas than other electroanalytical methods. The basic principle of these methods is to measure the current that occurs during the oxidation or reduction of the analyte in the chemical reaction.

Electrochemical methods began with the Czech chemist Jaroslav Heyrovsky, discovering the basis of polarography in 1922 and took an important place among the analytical methods. Especially, since the 1980s, it has been possible to develop electrodes that have been modified mechanically or chemically with improved technology. In modification processes, polymers, organic ligands, inorganic clays, phthalocyanines and nanoparticles have been commonly used for the detection of electroactive substances in very small volume complex samples such as biological, environmental and human bodies. In the last twenty years, even very small quantities of substances that are electroactive have been additionally analyzed at high precision, selective by electrochemical methods by carbon-based or modified electrodes have wonderful properties. Electroanalytical methods have also an important place in quantification as well as in obtaining details such as determination, adsorption, reaction rate and equilibrium constants of the number of electrons transferred in the reduction or oxidation electrode reactions. In short, electroanalytical methods provide details on direct or indirect quantitative and qualitative analysis of electroactive species such as antioxidants, drugs, pesticides, etc.

2.1 Voltammetric application for the determination of antioxidant capacity

Voltammetry is a potentiostatic assay based on the recording of the peak current at controlled potential variation by the oxidation or reduction which enables qualitative and quantitative analysis by means in electrochemical reactions. Over the last two decades compared to other electroanalytical techniques, voltammetry has been intensely curious in all the electroanalytical methods due to their are used to analyze numerous compounds by anodic or cathodic scanning and to investigate their conceptual basis of electro-mechanism. There are four voltammetric techniques including cyclic (CV), linear (LSV), differential (DPV), and square (SWV) are commonly used to determination of antioxidant-type compounds.

Voltammetric techniques are an alternative analytical method, proved to have an excellent correlation compare with another conventional analytical process, for a while to study the AC in various food and beverage samples. They can be a benefit to characterize which species compounds have a greater contribution to the antioxidant capacity present for the real samples in terms of quantitative and qualitative by controlled the half-wave peak potential, peak current and the electron transfer number in reaction. The antioxidant capacity is related to the peak currents of oxidation species caused by hydroxyl groups (–OH) and antioxidant species contains many hydroxyl groups. They commonly give an electro-oxidation broad peak at a range of 400 mV- 600 mV depend on pH. So that, almost all antioxidant substances have electro-activity compounds and their peak current and peak potential provide quantitative and qualitative details, respectively. Further, the voltammetric techniques allow investigating the electrochemical behavior of antioxidant agents and interaction with oxygenated species.

Voltammetric methods have gained an important place among determinations of the antioxidant capacity in the last decade. Moreover, due to their great superiority, the use of complex samples such as food and beverages they have become widespread and widely found in the literature. Among these electroanalytical methods, square wave stripping, different pulse stripping, and cyclic voltammetric techniques are the most commonly preferred for the analysis of antioxidants by accuracy and precision. From past to these days, the compounds used as standard agents for the evolution of the AC by studies electrochemical methods are apigenin, ascorbic acid, caffeine, catechin, chlorogenic chrysin, p-coumarin acid, eugenol, fisetin, gallic acid, kaempferol, luteolin, morin, quercetin, rutin, t-resveratrol, Trolox and Malvidin-3-glucoside. As far as we have examined the literature, scientists have however preferred ascorbic acid, caffeic acid, gallic acid, catechin, rutin and quercetin which are often used as antioxidant standard substances due to excessive availability of these substances in food and drink. The chemical structures of some antioxidant molecules are given in **Figure 1**.



Figure 1. Molecular formulas of commonly used antioxidants.

2.1.1 Cyclic voltammetric technique

Cyclic voltammetry (CV) is usually the first experiment in the electrochemical operation of a compound in biological materials as nature samples to get in details about the electro-behaviors. In particular, to study the thermodynamics, kinetic, electron transfer, substance transfer type, and as well as quantitative determinations of oxidation or reduction processes can be carried out by cyclic voltammetric technique. In addition to taking a single measurement with CV, sequential multiple measurements can be taken. The most common applications of cyclic voltammetry are additionally electro-polymerization, electrochemical characterization, and the design of modified electroanalytical systems. Two types of cyclic voltammograms can be obtained as irreversible or reversible, depending on the chemical components of the target molecules. In reversible voltammetry, there is a difference of about 59 mV between the reduction and oxidation peak potentials (**Figure 2**).

During the past years, cyclic voltammetry has been used as an alternative to existing methods to evaluate the antioxidant sensing in natural samples such as teas, biological fluids, beverage juices plants, foods and beverage juices on different working electrodes. The most using parameter is peak current because of its proportional to the concentration of the antioxidants. Peak current heights also provide quantitative information about the amount of antioxidant capacity in food samples. The carbon-based working electrodes such as glassy carbon electrode (GCE), carbon paste electrode (CPE), screen printed carbon electrode (SPCE), and modified electrodes (Nanoparticle/GCE, Nanoparticle/CPE, Fe₃O₄/GCE) have been widely preferred in electrochemical measurements for the analysis of total antioxidant capacity (TAC). Peak current and peak potential values of standard substances such as ascorbic acid, caffeic acid, catechin, coumarin, gallic acid, morin, quercetin and rutin were commonly taken care of for the evaluation of TAC. The amount of antioxidants in food samples is generally given as equivalent gallic acid, equivalent value quercetin, etc.

Even though the CV method raises doubts about sensitivity, it also has great advantages. Quick, simple, low detection limit, cheaper and easier application are



Figure 2. Potential-excitation signal and voltammograms for the cyclic voltammetry in details.

summarized as great advantages. Interferences effect on antioxidant capacity by a nonantioxidant agent to reducing TAC and non-selective to a family of molecules between carotenoids and polyphenols unless the electrode is modified are drawbacks properties. Despite all of these disadvantages, CV attracts a great deal of attention among analytical methods, and a large number of studies deal with CV are also being undertaken. A large part of the work done up to day time to determine the antioxidant capacity by the CV method is summarized in **Table 1**. **Table 1** includes the type of working electrode, working range, the limit of determination (LOD), the limit of quantification (LOQ), measurement parameter, standard compound and food sample.

2.1.2 Square wave voltammetric technique

Square wave voltammetry (SWV) can be used to perform a faster experiment than other voltammetric techniques. Commonly when the scanning speeds of other techniques are of 1–10 mV/second or more, in the square wave voltammetry a scanning speed is used at 1 V/second. Thus, the target molecule can be analyzed more quickly by SWS. The square wave voltammetry can combine with the stripping technique. Thus, a stripping voltammetric technique was developed to determine electroactive substances at high sensitive enables in ultra-trace concentration levels. Especially, ultra-trace target substances in complex samples can be analyzed by combining the technique with the enrichment stripping process. The working principle of the stripping technique is the same as square wave voltammetry and only two new parameters are more applied as the accumulation time and the accumulation potential (**Figure 3**).

Nowadays SWV and square wave stripping voltammetry (SWSV) are frequently applied to deduce compounds such as drugs, heavy metals, pesticides and antioxidants, etc. in numerous specimen types because they have excellent analytical sensitivity and selectivity. Furthermore, SWV and its derivate combined technique can be applied for simultaneous determination of compounds which are close oxidation or reduction peak potentials like paracetamol, ascorbic acid, uric acid and dopamine. In the last decade, SWV and SWSV have been more effective in determining antioxidant substances in the complex matrix samples and are superior compared with analytical methods especially spectrophotometric to evaluate quantification and qualification. It is one of the most important electroanalytical methods for the determination of antioxidants since it is a wide working range, low detection limits, easy to apply, cheap and non-pretreatment. Furthermore, they have been successfully analyzed the phenols in food samples which is called a type of important antioxidant such as o-phenylenediamine, p-chlorophenol, paminophenol hydroquinone, pyrocatechol and phenol, etc. At the same time, various antioxidant substances such as gallate, gallic acid, quercetin and caffeine were easily studied in food or beverage samples at high precision, accuracy and selective on the carbon-based electrode. Besides, at nM concentration of antioxidant substances comparable to chromatographic techniques have been determined by modified electrodes which are increasing conductivity accurately and selectively in tea samples. Evaluation of antioxidant capacity by SWV or SWSV techniques in the last 4 years are summarized in **Table 2** according to the type of working electrode, working range, the limit of detection (LOD), quantity limit (LOQ), measurement parameter, standard composition and food sample.

2.1.3 Differential pulse voltammetric technique

Differential pulse voltammetric technique (DPV) is one of the most widely used for the analysis of both organic and inorganic species. Pulse voltammetry

Ref	[24]	[25]	[26]	[27]	[28]		[29]	[30]
Interferences	1	Catechin, gallic acid	I	L-cystine, glycine, alanine, serum albumin, uric acid, citric acid, ascorbic acid, and urea			I	I
Peak Potentials	2.1 V	+ 0.5 V	+ 0.44 V	0.47 V	0.51 V 0.54 V	0.48 V	I	-0.90 V (cathodic) -0.75 V (anodic)
Optimum pH	I	pH 7.0 (PBS)	pH 4.0 (PBS)	PH 7.6 (PBS)	pH 4 acetate buffer		l	pH 7.00 (PBS)
Samples	walnut	Black Tea Samples Camellia sinensis	pharmaceutical sample (Captopril)	Human urine samples	Apricot pomace extracts black currant pomace extracts	Grape pomace extracts.	wine solution	Venezuelan propolis
LOQ		I			1			I
LOD	1	I	89,4 µM	33.43 (±0.21) nM			I	I
Linear range		I	200– 1000 μΜ	20 μM - 200 μM	I			I
Analyte	juglone (5-hydroxy- 1,4,-naphthoquinone)	polyphenols	Rutin (RT)	sinapic acid (SA)	caffeic acid, chlorogenic acid, quercetin, gallic acid, (+)-catechin, ascorbic acid		Tannins	polyphenols and flavonoids
Electrode	Pt electrode	GCE	Graphite Paste Electrode	Nanotuned Gold Nanoparticles and Solvothermally Reduced Graphene modified GCE (GCE/EAUNPs4 /rGO/ Naf.)	glassy carbon electrode		glassy carbon disc electrode	glassy carbon electrode
Method	CV	CV	CV	CA	CV		CV	CV

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Ref	[31]	[32]	[33]	[34]	[35]	[36]	[37]	[38]
Interferences	I	Anthrarufin (H2Arf), Chrysazine (H2Cz), Anthraflavin (H2Afv)	glucose and fructose	1		I	BHA, BHT, diphenylamine, and ascorbic acid (AA)	
Peak Potentials	1	1	I	I	0.30 V	I	+0.45 V	I
Optimum pH	pH 7.0 (PBS)	pH 7.00 (Aqueous)	pH 7 (PBS)	pH 7.0 (PBS)	pH 6.0 citric acid	pH 7 (PBS)	pH 3.5 ammonium formate buffer	pH 6.0 citric acid
Samples	red wine, coffee and green tea	1	Malaysian honey	Astragalus membranaceus	Real samples	Thymus vulgaris	Salmon Samples	real food samples
LoQ	0.6 µM	$M\mu$ 10.429 \pm 1.133 μ	I	I	1	I	20.0 mM	I
LOD	Mıµ 9.1	3.129 ± 1.200 µМ	1	0.08 µМ	0.21 µM	I	7.5 mM	Мц 170.0
Linear range		0-36 µM	I	0.1–1 µМ	1.0– 48 μΜ	I	20– 100 mM	1.0– 194 µМ
Analyte	Trolox	Quinizarin (H2Qz)	polyphenols	p-nitrophenol (p-NP)	Curcumin	polyphenols, tannins, flavonoids, and sterols/ triterpènes.	Ethoxyquin (EQ)	monohydroxycinnamic acid
Electrode	carbon paste electrode	Carbon Paste Electrode (CPE)	glassy carbon electrode (GCE)	ZnO nanoflowers modified carbon paste electrode	carbon nanotube (CNT)- carboxymethylcellulose (CMC) electrode MWCNT-CMC/Au	GCE	carbon screen printed electrode (cSPE)	carbon nanotube (CNT)- carboxymethylcellulose (CMC) electrode
Method	CV	CV	CV	CV	CV	CV	CV	CV

Ref	[39]									[40]
Interferences	I									catechol (CT), dopamine (DA), caffeic acid (CA), morin (MR), hydroquinone (HQ), uric acid (DA), ferulic acid (FA)
Peak Potentials	0.24 and 0.46 V									+0.59 V
Optimum pH	pH 7.4 (PBS)									рН 3.0 (PBS)
Samples	wine and food samples									Wine sample
LOQ	8.24 µM	0.339 µM	0.371 µM	4.58 μM	0.613 µM	0.353 µM	6.16 µM	0.288 µM	0.312 µM	1
LOD	2.47 µM	0.282 µM	0.111 µМ	1.37 µM	0.184 µМ	0.106 µM	1.85 µM	0.0863 µM	0.0937 µM	0.02 µM
Linear range	30.0– 540 μΜ	10.0– 350 μΜ	1.00– 210 μΜ	30.0– 540 μΜ	10.0– 350 μΜ	1.00– 120 μΜ	30.0– 540 μΜ	0.20– 350 μΜ	1.00– 120 μΜ	0.1– 130 μM
Analyte	catechol, (CAT)	4-ethylcatechol (4-EC)	4-ethylguaiacol (4-EG)	CAT	4-EC	4-EG	CAT	4-EC	4-EG	Gallic Acid (GA)
Electrode	CPE			CPME-CNT			CPME-AB			HP-ZnO/GCE
Method	CV			I			I			C

Ref	[41]	[42]	[43]	[44]					[44]				
Interferences			I	I					I				
Peak Potentials	1		0.22 V	483 mV	445 mV	472 mV	652 mV	260 mV	132 mV	139 mV	122 mV	377 mV	201 mV
Optimum pH	pH 7 (PBS)	pH 1.2 HClO4	0.1 M H2SO4	pH 3.6 tartaric acid buffer					pH 3.6 tartaric acid buffer				
Samples	juice samples (like peaches and apple juices)		Wines and beers	grape skin and seed					grape skin and seed				
LOQ		I	14.43 µМ	I					1				
LOD	4.24 (±0.12) µМ	1	4.33 µM	I					1				
Linear range	10 µM - 1 mM	5 mM	10– 300 μΜ	0.1 mM					0.1 mM				
Analyte	caffeic acid (CA)	piperine	Caffeic acid (CA)	Catechin	Caffeic acid	Gallic acid	Oenin chloride	Rutin	Catechin	Caffeic acid	Gallic acid	Oenin chloride	Rutin
Electrode	glassy carbon electrode/poly(3,4- ethylenedioxythiophene)- gold nanoparticles- sinusoidal voltage (GC/PEDOT-AuNPs-SV)	carbon electrodes	Poly(3,4- ethylenedioxythiophene)- tyrosinase PEDOT-Tyr	glassy carbon electrode (GCE)					Single Walled Carbon Nanotubes modified Screen Printed Carbon Electrodes (SWCNT- SPCE)				
Method	CV	CV	CV	CV					CV				

Ref	; [45]	[46]	. [47]		[48]	[49]	
Interferences	uric acid, ferulic acid, vanillic acid gallic acid, and catechol	urine and blood serum samples	A-lactose, L- aspartic acid, L- glutamic acid, L- histidine		AA	AA, Fructose, Xylose, Mannose	
Peak Potentials	I	I	I	I	I	I	
Optimum pH	pH 3.6 (PBS)	I	pH 6.6 (PBS)		$1 \mathrm{M} \mathrm{H}_2 \mathrm{SO}_4$	PBS	
Samples	Active Detox, DVR-Stem Glycemo, and green tea	human urine and blood serum sample	milk sample		1	real tear samples	
LOQ	10,77 nM	I	23 µM	149 µM	1	I	
LOD	3,23 nM	0.076 µM	7 µM	45 μM	5.0 μM	0.1 µM	
Linear range	0.1– 40 μM	0.5– 50 μΜ	6– 100 μΜ	30– 500 μΜ	5-80 µM	0,3-3 µM	
Analyte	Caffeic acid (CA)	UA	UA	АА	UA	UA	
Electrode	Carbon nanofibers CNF	Graphene/Neutral Red -GCE	PEDOT (poly(3,4- ethylenedioxythiophene) /GCE		ZnO-graphene/ ITO	GOx-chitosan /Co3O4/ Au- graphene transistors (GOx-CHIT/Co3O4 modified SGGT)	
Method	CV	CA	CV		CV	CV	

	CV technique.
	юq
	capacity
	antioxidant
	of
Table 1.	Evaluation

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Figure 3.

Potential-excitation signal and voltammogram for the square wave stripping voltammetry in details.

techniques were proposed by Baker and Jenkin in 1952 as a more sensitive measurement electroanalytical method. Differential pulse voltammetry techniques can be used to determine up to 10^{-8} M concentration of the target agents. The peak current (I_p) is a function of the concentration for the electroactive species and is linear as I*p* = *f* (C). Also, it is possible to analyze substances not only quantitative analysis but also qualitative analysis with pulse technique. The peak currents are related to the concentration of the substance whereas the peak potential values are related to the selectivity. Thus, simultaneous determinations of the substances have been studied by DPV on bare or modified electrodes (**Figure 4**).

Nowadays, quite a lot of DPV studies can be found in the literature for the very sensitive detection of heavy metal, drug, pesticide, antioxidant agent and inorganic/ organic species on numerous bare and modified working electrodes. Besides, DPV is one of the most important candidates to determine the trace amount of target agents in analytical methods due to its high sensitivity and selective. Also, it can be applied to complex samples as biological and food samples such as blood and serum, beverages. Especially, DPV has an important place among antioxidant determination methods because of these advantages and the availability of low concentration.

In recent years, DPV has been used frequently in determining the total antioxidant capacity without any pretreatment of solid and liquid food samples. The complex matrix such as biological and food samples contain very dense different types of substances. For this reason, despite it is indeed very difficult to selectively and precisely determine the antioxidant capacity in some complex matrixes; DPV is the most applicable method for such species. There are also plenty of studies were published which deal with chlorogenic acid, caffeic acid, p-coumaric acid, quercetin, gallic acid and ferulic acid, etc. as illustrating the antioxidant properties were determined by DPV on bare or modified electrodes based on carbon nanomaterials. Several applications, based commonly on the used as a determination of antioxidant capacity are given in **Table 3**.

In amperometric techniques, the current produced during the reduction or oxidation of an electroactive species at a constant potential value that is applied between a working electrode and reference electrode is measured, in this way providing specific quantitative electroanalytical knowledge for the target analyte. Especially, amperometric, which is based on electrical current analysis, is commonly utilized in microchip electrophoresis applications owing to its high sensitivity, it also lets for the determination of electroanalytical active species without derivatization, accomplishing adjustable versatility and selectivity (**Table 4**).

Ganesh et al., synthesized zinc oxide nanoparticles using mechanochemical synthesis technique. New ZnO nanoparticle as hexagonal prism was investigated by scanning electron microscopy, X-ray diffraction, particle size distribution, ultraviolet– visible spectroscopy, and energy-dispersive X-ray spectroscopic methods. Electrochemical properties of the newly prepared electrode were characterized by using an amperometric method and cyclic voltammetry technique. The prepared electrode has

Ref	[50]	[51]	[52]	[8]	[53]	[54]	[55]
Interferences	inorganic ions (Na ⁺ , K ⁺ , Ca ²⁺ , Cl ⁻ , SO ₄ ²⁻ , CO ₃ ²⁻), Glucose, Fructose, Eugenol, Capscasin	Aspartic acid (ASP), Gallic acid (GAL), Sucrose (SUC) and Tartaric acid (TAC)	Vitamins (vitamin C, B2, and B6), possible ingredients in LA pharmaceutical formulations	1	1	Cu^{2*} , Pb^{2*} , Cd^{2*} , Mg^{2*} , Ca^{2*} , K^* , Cl^- , and ascorbic acid and catechol, glucose, and fructose	epinephrine, dopamine, paracetamol, guaiacol and catechol, uric acid and inorganic ions (Ca ²⁺ , Mn ²⁺ , Fe ²⁺ , Zn ²⁺ , SO ₄ ²⁻ and NO ₃ ⁻)
Peak Potentials	0.6 V	0.18–0.22 V (ox and red) 0.31–0.30 V	(ox and red) —	I	I	I	0.23 V
Optimum pH	pH 3.0 (BRB)	pH 6.0 (PBS)	pH 6 (BRB)	рН 4.00 (КНРТ)	pH 5.0 (PBS)	pH 2.0 H ₂ SO4	pH 6.0 (PBS)
Samples	walnut and pomegranate	crude natural fruits (orange, apple and onion)	dietary supplements	Citrus juice, fruits and peel	tea samples (tea:green, basil and black)	tahini halva samples	Red wine Green tea Apple juice Lemon juice
LOQ	0.21 nM		I	0,111 µМ	I	I	I
LOD	Мп 70.0	0.029 µМ 0.058 µМ	0.37 µM	44 nM	0.007 µМ	85 nM	0.026 µM
Linear range	I	0.099– Мц 090 I	2–100 µM	75 nM-0,1 mM	0.01– 100 μM	0.2 mM- 1.0 mM	0.0498– 0.794 µM
Analyte	Ellagic acid (EA)	quercetin (QRT) rutin (RT)	α-lipoic Acid	naringenin (NGN)	Quercetin (QCT)	Sesamol	Quercetin (QCT)
Electrode	glassy carbon electrode modified with graphite/ bismuth (III) oxide (Gr/ Bi ₂ O ₃ /GCE)	CPE/PAG	TPC₀₃O₄&SWCNT@CPE	pencil graphite electrode	Single-Walled Carbon Nanotube Modified Glassy Carbon Electrode (SWCNT/GCE)	Untreated boron doped diamond electrode (BDDE)	immobilization (in solution) of laccase onto the activated carboxylic groups of carboxymethyl- botryosphaeran (CMB).
Method	SWV	SWV	SWV	SWV	SWV	SWV	SWV

Ref		[56]	[57]	[10]	[58]
Interferences		NO3 ⁻ ,Na ⁺ , Cu ²⁺ , K ⁺ , 4- nitrophenol, rutin, dopamine, caffeic acid, luteolin, tetracycline, hydrogen peroxide, glucose, ascorbic acid, epinephrine, uric acid, and quercetin	glucose, sucrose, ascorbic acid, riboflavin, phenylalanine, L- tryptophan, L-tyrosine, bisphenol A, lysine, uric acid, and two metal ions such as Na + and Co2+	1	caffeic acid, p-coumaric acid, gallic acid, ferulic acid, sinapic acid, and syringic acid, K ⁺ , Na ⁺ , Ca ²⁺ , Mg ²⁺ , $Zn^{2+}, Cu^{2+}, Fe^{3+}, NO_3^{-}, CI^{-}$ and SO ₄ ²⁻
Peak Potentials		+0.20 V	I	1	0.68 V 1.15 V 1.50 V
Optimum pH		pH 7.0 (PBS)	pH 5 (BRB)	pH 3.6	0.1 M HNO3
Samples		Human serum samples and urine samples were	Human plasma	Wine samples	Food&beverage samples (vanilla- enriched instant coffee, vanilla sugar, cola soft drink)
LOQ		I	0,1 nM	I	I
LOD		1 nM	0,033 nM	I	0.4 µМ 0.38 µМ 0.15 µМ
Linear range		2-123 nM	0,1 nM - 1 mM	I	2.8 μM - 0,17 mM 3.3 μM - 0,33 mM 0.52 μM - 0,21 mM
Analyte		Silymarin	Quercetin	Polyphenols (caffeic acid, gallic acid, catechin and malvidin-3- glucoside	5-O-Caffeoylquinic acid (5-CQA) vanillin (VAN) caffeine (CAF)
Electrode	(CBPE-CMB/LCE) biosensor	CNF-ZnO modified glassy carbon electrode (CNF- ZnO-GCE)	gold nanoparticle/graphene quantum dots (AuNP/ GQD) nanozyme-modified screen-printed carbon electrode (AuNP/GQDs/ SPCE)	SWCNTs-SPCE	boron-doped diamond electrode (CPT-BDDE)
Method		SWV	SWV	SWV	SWV and AdSV

es Ref	[65]
Interferenc	1
Peak Potentials	I
Optimum pH	pH 2 (BRB)
Samples	I
LOQ	I
LOD	0.351 μg/ kg 1.13 mg/ kg
Linear range	5.66– 113.37 μg/ kg 4.76– 92.40 mg/ kg
Analyte	Cu(II) and TBHQ
Electrode	Gold disk
Method	SWASV

 Table 2.

 Evaluation of antioxidant capacity by SWV or SWSV.



Figure 4.

Potential-excitation signal and voltammogram for the differential pulse stripping voltammetry in details.

a wide working linear range between $0.1-130 \mu$ M with a detection limit of 0.02μ M. Obtained results showed that the prepared electrode has numerous active surface sites, good electronic activity, and surface area. They applied the proposed electrode to the determination of gallic acid in samples as wine successfully [40].

Kumar and coworkers successfully synthesized NiO nanoparticles from natural fruit using an efficient, simple, and low-cost technique. The obtained NiO nanoparticles were investigated with various methods such as FTIR, XRD, TEM, SEM, UV, and PL. XRD studies showed that NiO nanoparticles have cubic geometry. The band of Ni-O bond was shown at 430 cm⁻¹. Photocatalytic properties of the obtained NiO nanoparticles were applied to photodegrade the methylene blue dye. They used the prepared electrode to the determination of dopamine with the LOD of 11 μ M [93].

Koçak et al. prepared a new composite electrode using carbon nanotube and polyl-methionine onto the glassy carbon electrode. Electrochemical properties and surface structure of the prepared electrode were studied using electrochemical impedance spectroscopy and scanning electron microscopy. Electrochemical properties of gallic acid with the proposed electrode were investigated in various techniques such as differential pulse voltammetry, cyclic voltammetry and amperometry. The obtained results of electrochemical studies exhibited that the prepared electrode shows a suitable method of determination for gallic acid in pH 2.2 BR buffer solution. The prepared sensor has a wide working linear range with two linear segments between 4 nM-1.1 μ M and 1.7–20.0 μ M with LOD of 3.1 nM. They used the prepared new sensor for the detection of gallic acid in various samples as black tea, green tea and wine samples. The experimental results showed that the proposed sensor exhibit high selectivity, reproducibility, stability and catalytic effect [88].

Potentiometry is an electrochemical technique based on measuring the potential difference between two electrodes called working and reference electrodes. The working basis of the potentiometry technique is the potential difference based on the concentration of an analyte in the sample solution relative to a reference electrode (**Table 5**).

Brainina and coworkers developed a new, simple, reliable and fast potentiometric method for the determination of plant total antioxidant activity. Plant micro suspension and extracts were analyzed by the proposed method. The experimental conditions for acquiring plant extracts were selected for the highest antioxidant activity as extraction time 20 min at +80°C. The characterization of plant micro suspensions reduces the duration of plant total antioxidant activity evaluation. Comparison of the obtained results of antioxidant activity of green tea and black tea micro suspensions samples with the results of the investigations of extracts prepared by a certified method showed no difference [95] (**Tables 6** and 7).

Ref	[09]		[61]	[11]	[62]		[63]	[64]	[65]
Interferences	phenolic compounds- caffeic acid, ascorbic acid,	coumaric acid, 2,5 dihydroxybenzoic acid, chlorogenic acid, rutin, and gallic acid	Ι	p-coumaric acid, hydroquinone, trans- ferulic acid, gallic acid, glucose, and ascorbic acid	Glucose, Fructose, Galactose		Sucrose, DA, AA, Glucose, Folic acid	AA, UA, FA, Trp, Mor, GA, Glucose and FoA	catechol (CC), hydroquinone (HQ), epinephrine (EP), dopamine (DA), uric acid (UA), and ascorbic acid (AA)
Peak Potentials	0,63 V	0,53 V	I	I		I	0.3 V		0.21 V
Optimum pH	$_{ m PH}^{ m PH}$ 2 $_{ m H_2SO_4}$		pH 3 (BRB)	pH 2.65 (BRB)	pH 7 (PBS)		pH 7.0 (PBS)	pH 7.0 (PBS)	pH 7.0 (PBS)
Samples	Melissa officinallis, Rosmarinus officinals		green tea and fruit juices	wine	real samples of dopamine injection, human serum,	and urine samples	Fresh human serum samples	Coconut water, coffee, tea	Red wine samples by
ГОО	l	I	I		I	I	1.2 μM	1.910 nM	
LOD	Мц 6	Mıµ	5 nM	018 µM	1.2 μΜ 42Μ	.42 μινι	.4 μM	157 nM	002 µM
	0	0.8	2	0.	· · · ·	>	0	0	0.
Linear range	3 μM – 0. 7 μM	2 μM – 0.8 70 μM	0.22– 2 55 μΜ	0.5– 0. 100.0 µМ	1–160 µM	0 MIH UCT-T	1.2–8.2 µМ 0	0.06- 0. 500 µМ	0.01- 0. 263.96 μΜ
Analyte Linear range	Rosmarinic $3 \mu M - 0$. acid (RA) $7 \mu M$	protoca- 2 μM – 0.8 techuic acid 70 μM (PCA)	Caffeic acid 0.22- 2 (CA) 55 μM	Caffeic acid 0.5– 0. (CA) 100.0 μM	UA 1–160 μM 3	ט זאון טכו-ו אנן	UA 1.2–8.2 µМ 0	Caffeic acid 0.06- 0. (CA) 500 µМ	Caffeic acid 0.01- 0. (CA) 263.96 μM
Electrode Analyte Linear range	GCE/PoPD/Pt Rosmarinic $3 \mu M - 0$. acid (RA) $7 \mu M$	protoca- 2 μM – 0.8 techuic acid 70 μM (PCA)	ZrO2NPs-AuNPs- DES/ Caffeic acid 0.22- 2 CPE (CA) 55 µM	Fluorine doped graphene Caffeic acid 0.5– 0. oxide/GCE (CA) 100.0 μM	CuO nano-rice/ GCE UA 1–160 μM 3		Poly(DPA)/SiO ₂ @ UA 1.2–8.2 μM 0 Fe ₃ O ₄ /CPE	Carbon paste modified Caffeic acid 0.06- 0. with Bi decorated (CA) 500 μM multiwalled carbon nanotubes and cetrimonium bromide (CTAB)	Bimetallic CoFeSe ₂ Caffeic acid 0.01- 0. nanosphere in (CA) 263.96 μM functionalized carbon nanofibers CoFeSe ₂ /f- CNF

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Ref	[66]	[67]	[68]	[69]	[20]
Interferences	catechol (CT), gallic acid (GA), ascorbic acid (AA), hydroquinone (HQ), and uric acid (UA)	dopamine (DA), catechol (CC), ascorbic acid (AA), uric acid (UA) and epinephrine (EP)	Cl., Br., SO4 ²⁻ , NO3 ⁻ , H ₂ PO ₄ ⁻ , Na ⁺ , K ⁺ , Mg ²⁺ and Al ³⁺ , glucose, L- serine, uric acid, urea, oxalic acid, glycine, alamine, L-cysteine, L- tyrosine, L-glutamic acid, and guanidine acid	Na ⁺ , K ⁺ , SO ₄ ²⁻ , Cl ⁻ , 40- fold glutamic acid, glycine, glucose, sucrose, urea, ascorbic acid, uric acid, and equal concentration of baicalein, luteolin, and vanillic acid	anions, cations and other polyphenols such as SO_{4}^{2-} , NO_{3}^{-} , CI^{-} , Fe^{3+} , Fe^{2+} , Zn^{2+} , Ni^{2+} , Cu^{2+} , Mg^{2+} , Ca^{2+} , K^{+} , Na^{+} , Li^{+}
Peak Potentials	I	I	I	1	A 12.0
Optimum pH	pH 7.0 (PBS)	PH 7.0 (PBS)	pH 6.0 (PBS)	pH 3.0 (PBS)	pH 4 (BRB)
Samples	coffee, green tea, red wine	red wine samples	caffeic acid tablets samples	human serum and urine samples	orange juices samples
LOQ	0.0068 µM	I		1	0.60 µM
LOD	0.002 µM	0.004 µМ	0.0003 µМ	5,2 nM	0.18 µМ
Linear range	0.1– 17.2 μΜ	0.05– 43 µМ	0.001 -10 µМ	0.2–7 µМ 7–50 µМ	0.50– 12.00 µМ
Analyte	Caffeic acid (CA)	Caffeic acid (CA)	Caffeic acid (CA)	caffeic acid (CA), chlorogenic acid (CGA), and gallic acid (GA)	p-coumaric acid (CA)
Electrode	Carbon/iron-based active catalyst f-MWCNTs/a-NaFeO2	N-doped carbon quantum dots/hexagonal porous copper oxide decorated multiwall carbon nanotubes N-CQD/HP- Cu ₂ O/ MWCNT/GCE	Ce-TiO₂/carbon nanotube composite Ce-TiO₂/CNTs	MOF-818 metal-organic framework-reduced graphene oxide/ multiwalled carbon nanotubes composite MOF-818/ RGO/MWCNTs/GCE	Fe ₃ O ₄ @ZIF-4 nano- hybrid on a glassy carbon electrode (GCE) (Fe ₃ O ₄ /GCE, ZIF- 4/ GCE)
Method	DPV	DPV	DPV	DPV	DPV

ĺ									
Ref		[71]	[72]	[25]	[73]	[74]			
Interferences	ions, citric acid, glucose, catechin and quercetin	1	tannic acid (TA)	Catechin, gallic acid	caffeic and ascorbic acid	I			
Peak Potentials		0.268 (±0.014) V	I	+ 0.5 V	I	$0.519 \pm 0.002 \text{ V}$ (Green tea)	0.528 ± 0.002 V (Black tea) 0.526 ± 0.001 V	(Mint tea)	$\begin{array}{l} 0.533 \pm 0.002 \ V\\ (Hibiscus tea)\\ 0.508 \pm 0.001 \ V\\ (Rosemary tea)\\ 0.571 \pm 0.005 \ V\\ (Phytotherapic)\\ 0.532 \pm 0.001 \ V\\ (Wine 1)\\ 0.525 \pm 0.002 \ V\\ (Wine 2)\end{array}$
Optimum pH		pH 7.4	pH 2.0 (BRB)	pH 5.5 (PBS)	pH 5.8; 7; 8 (PBS)	0.1 M HClO ⁴			
Samples		food supplements	Orange juice	Black Tea Samples Camellia sinensis	White wine Green tea Apple juice	Tea (Green, Black, Mint, Hibiscus, Rosemary),	wine and phytotherapics	1	
LOQ		I	1	I	70–310 μM	0.01 µМ	0.02 µМ	0.003 µM	0.02 µМ
LOD		0.03 mg/L	1.96 nM	I	23–103 μM	0.004 µM	0.005 µM	0.001 μM	0.005 µМ
Linear range		I	1	I	0.1–2 mM	0.1–5 µM	0.1–5 µM	$0.1-5 \mu M$	0.1–15 µM
Analyte		Melatonin	quercetin (QU)	polyphenols	gallic acid	Caffeic Acid (CA)	Gallic Acid (GA)	Catechin	Quercetin (QCT)
Electrode		graphene modified screen-printed electrode	multi-walled carbon nanotubes modified carbon paste electrode (MWCNTs/CPE).		screen printed carbon electrode	alumina-modified glassy carbon electrode GCE			
Method		DPV	DPV	DPV	DPV	DPV			

Ref	[75]		[76]		rbic [77] ne U), nd 3P)	such [78] 1,), Mg ²	uric [79] CC), HQ), Ni ²⁺ ,	ioi
Interferences	I		0.05 V		uric acid (UA), asco acid (AA), dopami (DA), hydroquinoi (HQ), catechol (CT epinephrine (EP), a norepinephrine (NI	AA, Bioflavonoids (s as quercetin, rutir naringenin, morin linorganic ions (Ca ²⁺ , *, K ⁺ , Zn ⁺ , Cu ²⁺ , Cl SO ₄ ²⁻)	ascorbic acid (AA), acid (UA), catechol (acid (UA), catechol (and hydroquinone (F K*, Na*, Mg ^{2*} , Cu ^{2*} ,	NO ³⁻ , CI ⁻ , sulfate i (SO ₄ ²⁻ , and glucose
Peak Potentials	0.05 V	0.35 V	0.445 and 0.550 V	(oxidation peaks)	0.34 V	I	+0.06 V (GA)	+0.14 V (CA)
Optimum pH	pH 7.4 (PBS)		pH 5 (BRB)		pH 7.0 (PBS)	pH 7.5 (PBS)	pH 7.0 (PBS)	
Samples	mimic human serum sample of fetal bovine	serum	mango (pulp, peel, and seed)		spiked meat samples (Chicken, Beef)	Fortified Fruit Juice Samples (lemon juice, orange rind, and peppermint extract that contain HES)	Grape juice samples	
LOQ	I	1	I		1	1	1	
LOD	63 µM	Мц 0.6	I		16.35 (±0.46) nM	50.0 nM	0.58 μM (1) 0.82 μM (2)	0.143 μM (1) 0.30 μM (2)
Linear range	0.32 to 3.4 mM	0.065 to 1.5 mM	19.92– 98.04 ppm	2.34– 472.18 μΜ	0.075- 460.15 µМ	0.1– 7.0 μΜ 7.0– 100.0 μΜ	5-1187 μM (1) 5-128 μM (2)	5-2082 μM (1) 5-128 μM (2)
Analyte	ascorbic acid (AA)	uric acid (UA)	Gallic Acid (GA)	Trolox	propyl gallate	Hesperidin (HES)	Gallic Acid (GA)	Caffeic Acid (CA)
Electrode	nanoporous gold electrodes (NPG)		glassy carbon electrode (GCE)		CoSe2@rGO modified SPCE	Nano-Graphene-platelets (nGp) - Brilliant green (Bg)/Modified carbon paste electrode (nGp- Bg/ MCPE)	5-amino-2-mercapto- 1,3,4-thiadiazole (p- AMT) on mitrogen-doped carbon sphere (N-CS)	modified glassy carbon (GC) electrode p- AMT@N-CS/GC (1) p-AMT@N-CS/GC (simultaneous addition)
Method	DPV		DPV		DPV	DPV	DPV	

Ref	[80]	[81]	[82]				[83]	
Interferences	metal ions (such as K ⁺ , Na ⁺ , Li ⁺ , Cu ²⁺ , Ca ²⁺ , Mg ² ⁺ , Fe ²⁺ , Zn ²⁺ , Fe ³⁺), rutin, 6-methoxy flavone, gallic acid, caffeic acid, biomolecules (viz. caffeine, ascorbic acid, citric acid and glucose)	Metals ions (K+, Cl–, Na +, Fe3+), ascorbic acid and quercetin	I	I			K^+ , Na^+ , Mg^{2+} , Ca^{2+} , NO_3^- , CI^- , and SO_4^{-2-} and glucose, rhamnose, and glucose, rhamnose,	sucrose as well as ascorbic acid, phenolic aldehydes (vanillin, syringaldehyde)
Peak Potentials	I	0.61 V	0.102–0.155 V	0.561-0.571 V	0.561-0.592 V	0.134–0.155 V	0.48 V	0.63 V
Optimum pH	pH 3 (BRB)	pH 2.0 (PBS)	Ethanol	3:2 ethanol/ water	4:1 ethanol/ water	Water	pH 2.0 (BRB)	
Samples	real tea samples (such as green, rosehip fruit, Turkish and Indian black tea)	Red and White Wine	pharmaceutical	"Quercetin"			Cognac XO Brandy VS Brandy 5-Star	
LOQ	43 nM	1	Ι				0.41 µM	0.37 µМ
LOD	13 nM	1.52 µM	3.2 μM	3 µM	4.4 μM	0.46 μM	0.12 µM	Мц 11.0
Linear range	0.1–1.5 µМ	0,1–1 µM	9–102 μM	10– 100 µМ	13–95 µM	0.15– 60 μM	0.75–10 10– 100 μΜ	0.75–7.5 7.5– 100 µM
Analyte	catechin	Gallic Acid (GA)	Quercetin	(QRT)			Gallic acid (GA)	Ellagic Acids (EA)
Electrode	SWCNT- Subphthalocyanine (CS) Hybrid Material modified GCE electrode (CS/GCE)	Cobalt oxide nanoparticles-modified carbon-paste electrodes (CoO-NPs-CPE)	graphite/chemically	modified silica ceramic electrode (SMICI/C)			glassy carbon electrode modified with polyaminobenzene	sultonic acid functionalized single- walled carbon nanotubes (f-SWNT) and poly (pyrocatechol violet) (polyPCV/f-SWNT/ GCE)
Method	DPV	DPV	DPV				DPV	

Ref	[84]	[85]	[86]	[87]	[88]	[8]	[68]
Interferences	Na ⁺ ,K ⁺ , Mg ²⁺ , Zn ²⁺ , ascorbic acid, glucose, starch, tyrosine and tartazine	AA, GLU, Na ⁺ , Mg ²⁺ , K ⁺ , Ca ²⁻ , Cl ⁻ , SO ⁴⁻	I	I	Na ⁺ , K ⁺ , Ca ²⁺ Mg ²⁺ , Zn ²⁺ , Cu ²⁺ , Ni ²⁺ , ascorbic acid, theophylline, caffeine, cysteine, glucose, fructose, sucrose, and glycine	I	ascorbic acid, caffeine, citric acid, l-cysteine, glycine, glucose, Na ⁺ , Mg ² * Ca ²⁺ , SO ₄ ²⁻ , NO ³⁻ and Cl ⁻ gallic acid, 6- methoxyflavon
Peak Potentials	I	1	I	I		I	
Optimum pH	pH 6.0 (PBS)	pH 7.0 (PBS)	pH 7.4 (PBS)	pH 3.20 Tartaric Acid Solutions	pH 2.2 (BRB)	рН 4.00 (КНРТ)	pH 2.0 (BRB)
Samples	Natural food supplement	fruit juices(Grape juice, Tomato juice, Orange juice)	Greigia sphacelata fruit (Chupón or Quiscal)	Wine	green tea, black tea, and red wine samples	citrus juice	Tea samples
LOQ	92 nM	0.93 µM	120 µM	1	I	102 nM	66 nM
LOD	27 nM	0.30 µM	43.8 μM	I	3.1 nM	30,6 nM	20 nM
Linear range	0,2 - 1 μΜ 1.5 - 4.5 μΜ	10.0– 50.0 µМ 0.9– 10.0 µМ	50 μM to 600 μM		0.004– 1.1 μΜ 1.7–20 μΜ	78,6 nM - 0,182 mM	0.25-3 µМ
Analyte	Curcumin	Naringenin (NR)	Trolox	Polyphenols	Gallic acid (GA)	naringenin (NGN)	Quercetin (QCT)
Electrode	Sodium dodecyl sulfate modified carbon composite paste electrode	implemented functionalized-MWCNT/ Nileblue- composite on carbon paste electrode (fMWNCT/NB/ MCPE)	vitreous carbon electrode	Screen Printed Carbon Electrodes	Poly(L-Methionine)/ Carbon Nanotube Glassy Carbon Electrode (PLM/ MWCNT/GCE)	pencil graphite electrode	3D SWCNTs-coumarin hybrid modified glassy carbon electrode (3DSWCNTs- coumarin/ GCE)
Method	DPV	DPV	DPV	DPV	DPV	DPV	DPSV

Ref	[06	91]	92]
Interferences	$\begin{array}{l} Fe^{3+},Cu^{2+},K^{*},Na^{*},\\ Ga^{2+},Cl^{-},SO_{4}^{2-} and\\ glucose, and 100-fold of\\ Mg^{2+} \end{array}$	morin and quercetin	Methotrexate (MTX), [folinic acid (FCA), tetrahydrofolic acid (THF), pyridoxine (PYR), and 5- methyltetrahydrofolate (5- THF)
Peak Potentials	0.40 V	0.25 V (ox) 0.096 V (red)	1
Optimum pH	0.10 M HCl	PH 3.0 (PBS)	PH 6.2 (PBS)
Samples	fresh chili pepper samples (Meiren chili pepper, Chaotian green chili pepper, Chaotian red chili pepper, Xiaomi green chili pepper, and Xiaomi red chili pepper)	black tea, coffee and synthetic drink of tea	drug tablets and human urine samples
год	0.15 µM	I	
LOD	0.05 µM	0.01 µM	2.02 ng/ mL
Linear range	0.16 - 16.37 µM	0.03 to 0.24 μM 0.25 to 2.0 μM	2.2- 30.8 ng/ mL
Analyte	Capsaicinoids	Rutin	folic acid (FA)
Electrode	unmodified screen- printed carbon electrodes (SPCEs)	screen-printed carbon electrode modified with single-walled carbon nanotubes (SWCNTs)) and Prussian blue (PB) coated with chitosan	molecularly imprinted poly (p-aminobenzene sulphonic acid) on carbon nanodots coated pencil graphite electrode (FA- imp/CNDs/PGE)
Method	DPAdSV	DpAdSV	DPCV

Table 3. Evaluation of antioxidant capacity by DPV or DPSV.

Ref	[40]	[93]	[88]	[60]		[94]			[62]	
Interferences	catechol (CT), dopamine (DA), caffeic acid (CA), morin (MR), hydroquinone (HQ), uric acid (UA), ascorbic acid (AA), ferulic acid (FA)	1	Na ', K*, Ca ²⁺ Mg ²⁺ , Zn ²⁺ , Cu ²⁺ , Ni ²⁺ , ascorbic acid, theophylline, caffeine, cysteine, glucose, fructose, sucrose, and glycine	I		I		Ι	Glucose, Fructose, Galactose	
Peak Potentials	+0.48 V	0.41 V	0.5 V	I		I		I	I	I
Optimum pH	pH 3.0 (PBS)	pH 7.2 (PBS)	pH 2.2 (BRB)	$_{ m H^2}^{ m PH~2}$ $_{ m H_2SO_4}$					pH 7 (PBS)	
Samples	Wine sample	Limonia acidissima natural fruit juice	green tea, black tea and red wine samples	Melissa officinallis, Rosmarinus	officinals	Chestnut shell waste extract/TPC			real samples of dopamine injection,	human serum and urine samples
LOQ	I	I	I	l	l	I	I	I	I	I
LOD	0.02 µM	11 µМ	0.5 nM	0.5 µM	0.6 µM	0.29 μM	0.14 µM	Μμ 60.0	0.83 µM	0.083 μM
Linear range	0.1– 130 µМ		0.002– 0.1 μΜ 0.2– 12 μΜ	1 μM – 55 μM	1 μM – 60 μM	1– 150 μΜ	1–50 μM	1–20 µM	0.83– 253 μΜ	0.083– 428.8 µМ
Analyte	Gallic Acid (GA)	Dopamine (DA)	Gallic acid (GA)	Rosmarinic acid (RA)	protoca- techuic acid (PCA)	gallic acid	Caffeic acid	Rosmarinic acid	UA	DA
Electrode	HP-ZnO/GCE	Nickel oxide nanoparticles modified glassy carbon electrode (NiO NPs/GCE)	Poly(L-Methionine)/ Carbon Nanotube Glassy Carbon Electrode (PLM/ MWCNT/GCE)	GCE/PoPD/Pt		Graphite/Lacc-PDA			CuO nano-rice/ GCE	
Method	Amperometry (AMP)	Amperometry (AMP)	amperometry (AMP)	Amperometry		Chronoamperometry (CA)			Chronoamperometry (CA)	

Table 4. Evaluation of antioxidant capacity by Amperometric technique.

Potentiometry – Potentiometry GCE	Antioxidant			2 2	Samples	Optimum pH	Peak Potentials	Interferences	Ret
Potentiometry GCE		I			black and green tea microsuspensions	pH 7.2 (PBS)	I	I	[95]
	chicoric acid	I	I	I	Echinacea flowers	pH 7.4 (PBS)	I	Ι	[96]
Potentiometry POM immobilization on the surface of a glassy carbon electrode	ice of a Polyoxometalates (POMs)	I			I	0.1 M HClO4	I	I	[67]

Table 5. Evaluation of antioxidant capacity by potentiometric technique.

	Ionic liquid-rGO-titania-											
NST	nafion-GCE	capsaicin	0.03–10 µМ	I 0.0032 µМ		F (Chun	Korean hot peppe Igyang pepper) sc	ır olution	pH 1.0 (BRB)	0.75 V	I	[86]
(AS1	gold disk electrode	2-tert-butylphenol (2-TBF)	9.12– 80.83 µg cm ⁻	-3 0.67 μg /L	2.22 µg cm	-3 mine	eral and synthetic	c oils	0,16 M H ₂ SO4	I	I	[66]
Table 6. Evaluation of	° antioxidant capacity by line	ear sweep voltammetr	v (LSV).									
Method	E	lectrode A	ınalyte Liı	near range	LOD	LOQ	Samples	Optimu pH	E E	Peak otentials	Interferences	Ref
EI electro index	ochemical CPEs (ele	(carbon paste ectrodes)	TAC 0.10)5-0.500 μM	40,4 nM	0,105 μM	olive oil samples	pH 7 (PBS	S)	I	I	[100]
PC peak	current		8.C	32 x 10 ⁻² -).500 μΜ	30,5 nM 8 ¹	0,2 nM						
Redox m	icrosensor Redox r	neasurements	gallic 0	.2–2 mM	49 μM 1	.48 μM	White wine	pH 5.8	8	Ι	Ι	[23]
			acid 0).1–2 mM	109 µM 3	31 μM	Serum	PH 7				
			0.	1-1.5 mM	74 μM 2	123 µM		pH 8				

> Table 7. Evaluation of antioxidant capacity by other techniques.

3. Conclusion

Electrochemistry is a powerful and versatile analytical technique for the determination of numerous substances such as drugs, pesticides, inorganic, antioxidanttype compounds and electroactive compounds by rapidly possible applications in a lot of fields. Electroanalytical methods besides providing details on quantitative and qualitative of analyte that offer validation parameters such as sensitivity, accuracy and precision, selective and linear working range. Moreover, it is superior to determine the target analyte by electroanalytical methods lack of interferences effect especially in a complex matrix such as biological and food samples contain countless substances. The improvement of simultaneous determination of analytes considerably has been carried out to be applied in biological and environmental systems by the sensitive and selective electrochemistry methods. Because of this, the use of many areas of electrochemistry is widespread.

Nowadays, electrochemical methods, especially voltammetry from medicine to the determination of antioxidants, have made an important place especially in the world of science. Not only analytical chemists but also biology, food engineering and all people who are engaged in food have been used electrochemical methods to determine the antioxidant capacity in plants, tea, beverages, carbonated beverages and solid food samples, etc. Compounds such as ascorbic acid, caffeic, catechin, ascorbic acid, quercetin, gallic acid and coumarin have been widely used as reference standard agents to an evaluation of antioxidant capacity by electrochemical methods have been carried out until today. Due to advances in electronics and computer science have provided significant benefits in terms of electrochemical instrumentation such as accuracy, sensitivity and easy application, the electro-analysis of antioxidant compounds is successfully applied by stripping voltammetric techniques at nM concentration level. The purpose of this review is to show that electroanalytical methods for commonly used antioxidant types may be the best analytical method for the quantitative and qualitative analyte and that they can successfully compete with more conventional methods especially spectrometric methods. Consequently, voltammetric techniques supply that even at low concentrations, the antioxidant capacities of food samples can be determined to be very fast, simple, non-pretreatment and highly sensitive compared to conventional analytical methods. The review presented that the antioxidant capacity of various food samples can be carried out by voltammetric techniques in the estimation in real samples.

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Citrus is an extensively produced fruit crop and is cultivated predominantly in tropical and subtropical regions of the world. The *Citrus* genus consists of a variable number of species due to the admixture of wide morphological diversity, intra- and interspecific sexual compatibility, apomixis, and spontaneous mutations. Citrus fruits are highly nutritious and beneficial for health due to the presence of bioactive compounds that have antioxidant, antitumor, anti-inflammatory, and blood clot-inhibiting characteristics. This book describes the citrus plant and its nutrients, nutritional value, and nutraceutical applications, as well as related biotic and abiotic challenges in its cultivation. Chapters cover such topics as citrus genealogy, production, and crop management; milestones achieved in citrus improvement; importance of weather conditions in cropping systems; effects of changing climate on citrus; and much more.

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