ANTIMICROBIAL ACTIVITY OF NATURAL SUBSTANCES



Edited by

Tomasz M. Karpiński

and

Artur Adamczak



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Tomasz M. Karpiński

Poznań University of Medical Sciences Poznań, Poland

and

Artur Adamczak

Institute of Natural Fibres and Medicinal Plants,
Plewiska, Poland

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Dr hab. Tomasz M. Karpiński, Assoc. Prof.

Department of Genetics and Pharmaceutical Microbiology Poznań University of Medical Sciences Święcickiego 4, 60-781 Poznań, Poland

Dr. Artur Adamczak

Department of Botany, Breeding and Agricultural Technology, Institute of Natural Fibres and Medicinal Plants, Kolejowa 2, 62-064 Plewiska, Poland

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Antibacterial and antifungal influence

of a melanin producer Pseudonadsoniella

brunnea culture fluid

T. Kondratiuk, T. Beregova, L. Ostapchenko

Educational and Scientific Centre 'Institute of Biology and Medicine', Taras Shevchenko National University

of Kiev, Volodymyrska str. 64/13, 01601 Kyiv, Ukraine

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ABSTRACT

Fungicidal and bactericidal effects of melanin producer, black yeast

Pseudonadsoniella brunnea (Meripilaceae, Agaricomycotina) culture fluid on test cultures of

pathogenic fungi Gibberella baccata (anamorph: Fusarium lateritium), Gibberella intricans

(anamorph: Fusarium gibbosum), Gibberella pulicaris (anamorph: Fusarium sambucinum),

as well as Fusarium incarnatum, Fusarium solani, Fusarium poae, Rectifusarium

ventricosum and phytopathogenic bacteria Agrobacterium tumefaciens, Clavibacter

michiganensis subsp. michiganensis, Pectobacterium carotovorum subsp. carotovorum,

Pseudomonas syringae pv. syringae, Xanthomonas campestris pv. campestris is for the first

time established within this study.

Keywords: Pathogenic fungi; Phytopathogenic bacteria; Bioactive compounds; Antagonism;

Biocontrol.

1. INTRODUCTION

The development of biotechnology, which is based on the potential of microorganisms

in obtaining biologically active substances (BAS), which can be widely used in various fields of human activity, is one of the main areas of modern world science.

The fungi of the genus Fusarium Link are the most common pathogens and harmful fungal diseases of crops. Today, many species of the genus Fuzarium considered to be the members of the genus Gibberella Sacc. after the teleomorphic (sexual) stage. So Fusarium verticilloides (Sacc.) Nirenberg. (synonym Fusarium moniliforme J. Sheld.), now considered to be synonym to Gibberella fujikuroi (Sawada) Wollenw. complex. Fungi of the genus Gibberella are often pathogen of corn cobs disease and other crops (rice, corn, sugar reed, soybeans etc.) in the world, especially in areas with high humidity and in years with high rainfall (rain) (to more than 60% of corn harvest may be damaged by Gibberella in these areas) [1-6]. For most Fusarium species, the sexual cycle does not predominate in the field [7]. Fungi of the genus Gibberella (anamorphs of the genus Fusarium) can produce mycotoxins (the fumonisins), which cause a carcinogenic effect on animals and humans. The presence of these toxins in plants is subjected to strict control [3, 5, 8]. The bacteria of the genera Pseudomonas [9], Pectobacterium [10], Xanthomonas [11], Clavibacter [12], and Agrobacterium [13] are the most common pathogens and harmful diseases of crops too. Species of bacteria *Pseudomonas syringae* is actually represented by over 50 different pathovar strains, which is a set of bacterial strains with similar characteristics differentiated by their distinctive pathogenicity toward one or more plant hosts [9, 14]. The pathogen Pseudomonas syringae pv. syringae takes its name from the host from which it was first isolated, but strains that have been proved to be pathogenic to lilac also infect more than 44 plant species and there are strains with the same determinative characteristics that do not attack lilac [15]. In a comparative analysis of rpoD sequences from a comprehensive range of strains from the *P. syringae* complex, found that strains from hosts associated with particular pathovars are distributed widely in the *P. syringae* complex [16]. *Pseudomonas syringae* pv. panici is a phytopathogenic bacterium causing brown stripe disease in economically important crops worldwide [17]. The strains of Pseudomonas syringae that are pathogenic causing disease on their hosts through the release of toxins and cell wall degrading enzymes [18].

Bacteria belonging to *Pectobacterium carotovorum* cause soft rot disease in diverse vegetables and crops worldwide. These bacteria produce several different plant cell-wall degrading enzymes such as pectinase (PCWDEs), polygalacturonase and cellulase. Since *Pectobacterium* species can produce PCWDEs, soft rot disease can occur in the field as well as during transportation and storage. *P. carotovorum* subsp. *carotovorum* has a wide host range including potato, carrot, lettuce, onion and Chinese cabbage [19-21]. In Morocco,

approximately 95% of the *P. carotovorum* isolates from potato plants with tuber soft rot are *P. carotovorum* subsp. *carotovorum* [22]. The plant pathogen *Clavibacter michiganensis* subsp. *michiganensis* is a gram-positive bacterium responsible for wilt and canker disease of tomato [12]. *Agrobacterium tumefaciens* causes crown gall disease on various plant species by introducing its T-DNA into the genome. Therefore, *Agrobacterium* has been extensively studied both as a pathogen [13].

Plant diseases need to be controlled to maintain the quality and abundance of food, feed, and fiber produced by growers around the world. Different approaches may be used to prevent, mitigate or control plant diseases. Beyond good agronomic and horticultural practices, growers often rely heavily on chemical fertilizers and pesticides. Conventional biogicides are widely used to manage fungal and disease of plants, with two major consequences. On the one hand, fungicide overuse threatens the human health and causes ecological concerns. On the other hand, this practice has led to the emergence of pesticide-resistant microorganisms in the environment [23-26].

Today, there are strict regulations on chemical pesticide use, and there is political pressure to remove the most hazardous chemicals from the market. Additionally, the spread of plant diseases in natural ecosystems may preclude successful application of chemicals, because of the scale to which such applications might have to be applied. Consequently, some pest management researchers have focused their efforts on developing alternative inputs to synthetic chemicals for controlling pests and diseases. Among these alternatives are those referred to as biological controls [23].

The microorganisms of different taxonomic and physiological groups and their metabolic products and the drugs created on the base on these biological substances are now widely used in many countries to protect crops from pathogenic microorganisms. This can significantly reduce the negative impact on the environment, including agrocoenosis which inflicted by chemicals (pesticides). The use of biological agents against pathogenic fungi and bacteria not only provides plant protection from diseases, but may stimulate their growth and development, enhance seed germination, increased productivity. The most common method of biocontrol of phytopathogenic microorganisms is search of antagonist microorganisms. The methods of biocontrol of phytopathogenic microorganisms are environmentally safe, cost-effective and do not disturb ecological [27-40]. For example, *Pectobacterium carotovorum* and *Pectobacterium atrosepticum* are dreadful causal agents of potato soft rot. Actually, there are no efficient bactericides used to protect potato against *Pectobacterium* spp.

Biological control using actinobacteria could be an interesting approach to manage this

disease [41].

Thus, research aimed at finding BAS microorganisms to create on their basis of

biological agents that protect plants from pathogens, is relevant direction. Special attention of

researchers deserves microorganisms producers of BAS habitats of which are associated with

extreme conditions. The stringent physical and chemical factors causing significant adaptive

changes in microorganisms, accompanied by increased synthesis of a number of metabolites

(potential BAS). For example, the metabolites of such Antarctic microscopic fungi found to

be potent sources of antimicrobial and antifungal activity [42].

Microscopic fungi and bacteria with distinct physiological and biochemical

characteristics, pointing the adaptation to adverse environmental conditions (synthesis and

accumulation of lipids, expression of antagonistic properties in relation to other

microorganisms) were found during a preliminary study of samples of moss, soil, lichen

obtained from Ukrainian Antarctic expeditions [43]. The Antarctic black yeast-like

fungus Pseudonadsoniella brunnea T.O. Kondratyuk et S.Y. Kondr. (Meripilaceae,

Agaricomycotina) [44] synthesizes and excretes into a culture fluid dark pigment melanin.

The first data on antioxidant, antibacterial, fungistatic wound healing properties of the gel

containing 0.05% melanin ("Melanin-gel"), which was synthesized by P. brunnea are

obtained. Application of the "Melanin-gel" on wound area enhanced wound cleaning from

dead tissue and reduced eschar, stimulated the early growth of granulation tissue, and

improved epithelialization of the wound [57].

The high fungicidal effect of melanin producer Pseudonadsoniella brunnea culture

fluid on test cultures of pathogenic fungi Fusarium oxysporum Scherht. em. Snyder & Hansen

and Gibberella fujikuroi (anamorph: Fusarium verticilloides) for the first time found in our

previous studies [45, 46].

The purpose of this study was to find out the nature of influence of the culture fluid of

Pseudonadsoniella brunnea on pathogenic fungi and bacteria.

2. MATERIAL AND METHODS

The pure cultures of pathogenic fungus from the collection of microscopic fungi of

Taras Shevchenko National University of Kiev (international acronym of collection – FCKU)

[47], and the phytopathogenic bacteria, as well as the culture fluid of Antarctic black yeast-

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like fungus, melanin producers *Pseudonadsoniella brunnea* 470 FCKU [44] were used for this study.

The following fungi, i.e.: *Gibberella baccata* (Wallr.) Sacc. (anamorph: *Fusarium lateritium* Nees) 332 FCKU, *Gibberella intricans* Wollenw. (anamorph: *Fusarium gibbosum* Appel & Wollenw.) 147 FCKU, *Gibberella pulicaris* (Kunze) Sacc. (anamorph: *Fusarium sambucinum* Fuckel) 336 FCKU, *Fusarium incarnatum* (Desm.) Sacc. 330 FCKU, *Fusarium solani* (Mart.) Sacc. 329 FCKU, *F. solani* 331 FCKU, *F. solani* 334 FCKU, *Fusarium poae* (Peck) Wollenw. 339 FCKU, *F. poae* 340 FCKU, and *Rectifusarium ventricosum* (Appel & Wollenw.) L. Lombard & Crous 333 FCKU were used as test-cultures during our study. The fungi *Gibberella baccata* 332 FCKU, *Gibberella intricans* 147 FCKU, *Gibberella pulicaris* 336 FCKU were studied in anamorph stage. So hereafter in the discussion we will use names of anamorph stage of fungi mentioned.

The following phytopathogenic bacteria, i.e.: *Pseudomonas syringae* van Hall 590 FCKU, *Pectobacterium carotovorum* subsp. *carotovorum* 591 FCKU, *Xanthomonas campestris* 592 FCKU, *Clavibacter michiganensis* subsp. *michiganensis* 593 FCKU, *Agrobacterium tumefaciens* 594 FCKU were used as test-cultures during our study too. Thesebacteria were obtained from the D.K. Zabolotniy Institute of Microbiology and Virology, National Academy of Sciences of Ukraine, Kyiv and were grown on potatodextrose agar (PDA) medium. Mentioned above bacterium cultures and some other bacterial isolates are included to the FCKU collection [47].

The standard potato-dextrose agar (PDA) culture medium was used for cultivation of phytopathogenic fungus. In the experimental process the zones of inhibition of fungal and bacterial pathogens (in mm) were assessed using the methods of well diffusion assay (the method of diffusion in agar) [48]. In the method of the well diffusion assay after stabilization of the PDA nutrient agar the medium surface was coated with 0.5 ml of spore suspension of each of the ten fungal pathogens and 0.5 ml of cell suspension of each of the five bacterial pathogens. Spores were washed-off from the initial medium with sterile water and placed in a sterile glass test tube. The suspension of test culture of fungi with density 1×10^6 conidia/ml was used. The density of the bacterial cell suspension (5×10^5 cell/ml) - 0.5 per McF was determined using a densitometer (microbiological analyzer) Vitek-2 (Bio Merieux, France). Then, a small well (10 mm in diameter) in the center of agar in petri dish was created. This well was then poured with 100 μ l of each examined of *Pseudonadsoniella brunnea* culture fluid treatment.

The zone of inhibition around the area of the well was measured after incubation at

28°C. Term of cultivation was 30 days at a temperature of 28°C for the fungal test-cultures

and three days for bacterial test-cultures. The number of the experiments per treatment was 3

and the experiment was repeated twice.

Performance of *Pseudonadsoniella brunnea* culture fluid on test culture of fungus and

bacteria was compared with the effect of the biocide of known class of quaternary ammonium

compounds benzalkonium chloride (trade name Katamin AB) at a concentration of 3% for the

active substance. The cultivation of *Pseudonadsoniella brunnea* was performed on a standard

liquid culture medium the Maltese-extract broth (MEB, HiMedia Laboratories, India). The

impact of Pseudonadsoniella brunnea culture fluid and benzalkonium chloride on fungal and

bacterial cultures was determined by the diameter of the zones with absence of growth of test

cultures of fungi that formed around the hole, where the test compound was added in culture.

Cultures of phytopathogenic microorganisms without additions of Pseudonadsoniella

brunnea culture fluid and biocide mentioned were served as control cultures.

Trinocular microscope Primo Star of Carl Zeiss and related morphometric computer

program AxioVision 4.8 (Carl Zeiss) were used for study of morphological features of studied

fungi. Calculation of arithmetic mean and standard deviation using Statistica 12 was used for

analyzing morphometric parameters of Gibberella fujikuroi, and the diameter of the zones of

growth absence.

3. RESULTS AND DISCUSSION

3.1. Influence of *Pseudonadsoniella brunnea* culture fluid on test culture of fungi

Within our study the Pseudonadsoniella brunnea culture fluid found to have

fungicidal effect on the studied test cultures of pathogenic fungi Gibberella baccata

(anamorph: Fusarium lateritium 332 FCKU), Gibberella intricans (anamorph: Fusarium

gibbosum 147 FCKU), Gibberella pulicaris (anamorph: Fusarium sambucinum 336 FCKU),

Fusarium incarnatum 330 FCKU, Fusarium solani (strains 329 FCKU, 331 FCKU and 334

FCKU), Fusarium poae (strains 339 FCKU and 340 FCKU), and Rectifusarium ventricosum

335 FCKU. Zones of growth inhibition of test cultures under the influence of the

Pseudonadsoniella brunnea culture fluid found to be stable and did not change throughout the

duration of the experiment. It was also established that the diameter of the zones without

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growth of the test cultures of pathogenic fungi, exposed to fungicidal activity of *Pseudonadsoniella brunnea* culture fluid, is similar to the diameter of the sterile zones formed under the influence of biocide benzalkonium chloride (Table 1, Fig. 1).

Table 1. Antifungal effect of *Pseudonadsoniella brunnea* culture fluid and benzalkonium chloride on the test cultures of pathogenic fungi

	The diameter (mm) of growth inhibition zone			
The test cultures	(including well diameter)*			
of pathogenic fungi	Pseudonadsoniella brunnea	Benzalkonium chloride, 3%		
	culture fluid	for the active substance		
Fusarium lateritium 332 FCKU	59.8±0.2	65.1±0.1		
Fusarium gibbosum 147 FCKU	45.6±0.25	48.2±0.15		
Fusarium sambucinum 336 FCKU	48.5±0.3	49.7±0.2		
Fusarium incarnatum 330 FCKU	49.8±0.25	55.6±0.2		
Fusarium solani 329 FCKU	39.5±0.5	57.8±0.25		
Fusarium solani 331FCKU	56.3±0.25	57.1±0.1		
Fusarium solani 334 FCKU	46.7±0.2	43.9±0.1		
Fusarium poae 339 FCKU	29.1± 0.15	45.6±0.2		
Fusarium poae 340 FCKU	31.2±0.2	46.8±0.1		
Rectifusarium ventricosum 335 FCKU	51.4±0.3	63.3±0.15		

Note: *- diameter of well is 10 mm; \pm Standard error of mean

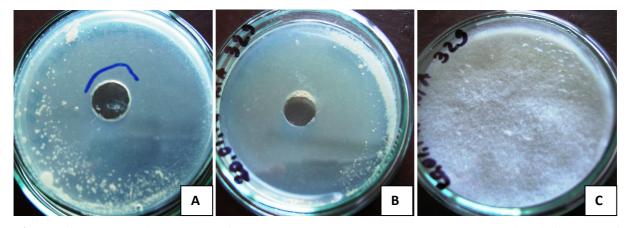


Figure 1. Zones of absence of *Fusarium solani* 329 FCKU growth under influence of *Pseudonadsoniella brunnea* culture fluid (A) and benzalkonium chloride (B). C = control culture. A – the reverse side of the Petri dish

Thus all test cultures of the pathogenic fungi investigated appeared to show high sensitivity to influence of *Pseudonadsoniella brunnea* culture fluid and biocide benzalkonium chloride. The growth of pathogenic fungal cultures studied found to be often characterized mainly by individual colonies of different diameters under influence of tested substances (Fig. 2).

The color of the mycelium of *Fusarium poae* 339 FCKU, *F. poae* 340 FCKU cultures found to be also changed in comparison with mycelium color in control cultures of strains studied. That is especially well distinct at the reverse side of the Petri dish (Figs. 1 and 2).

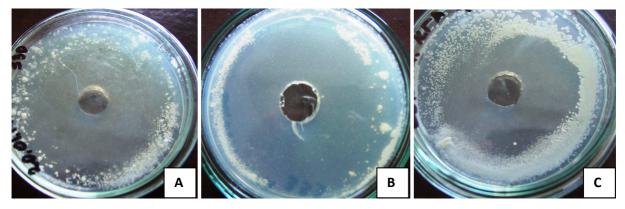


Figure 2. Effect of *Pseudonadsoniella brunnea* cultures fluid on the test culture: the absence of growth zone around the hole, where the *Pseudonadsoniella brunnea* culture fluid was added, growth by individual colonies. *Fusarium incarnatum* 330 FCKU (A), *Rectifusarium ventricosum* 335 FCKU (B), *Fusarium poae* 340 FCKU (C). B – the reverse side of the Petri dish

3.1. Influence of *Pseudonadsoniella brunnea* culture fluid on test culture of phytopathogenic bacteria

Within our study the *Pseudonadsoniella brunnea* culture fluid found to have biocidal effect on the test cultures of the phytopathogenic bacteria studied, except *Pectobacterium* carotovorum subsp. carotovorum. The impact of *Pseudonadsoniella brunnea* culture fluid on *Pectobacterium carotovorum* subsp. carotovorum 591 FCKU was evaluated as bacteriostatic: diameter of the growth inhibition zone after one day was similar to the zone of growth inhibition of the other bacterial test cultures (it was 39.3 ± 0.15). However zone of the growth inhibition of *Pectobacterium carotovorum* subsp. carotovorum decreased to 15.7 ± 0.1 at the third day (Table 2).

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Table 2. Antifungal effect of *Pseudonadsoniella brunnea* culture fluid and benzalkonium chloride on the test cultures of phytopathogenic bacteria

	The diameter (mm) of growth inhibition zone			
The test cultures of	(including well diameter)*			
phytopathogenic bacteria	Pseudonadsoniella	Benzalkonium chloride,		
	brunnea culture fluid	3% for the active substance		
Agrobacterium tumefaciens 594 FCKU	39.3±0.5	41.5±0.2		
Clavibacter michiganensis subsp. michiganensis 593 FCKU	38.8±0.4	40.2±0.1		
Pectobacterium carotovorum subsp. carotovorum 591 FCKU	15.7±0.1	39.7±0.15		
Pseudomonas syringae pv. syringae 590 FCKU	38.5±0.5	39.9±0.2		
Xanthomonas campestris pv. campestris 592 FCKU	Growth absent	41.1±0.3		

Note: *- diameter of well is 10 mm; \pm Standard error of mean

The test cultures of phytopathogenic bacterium *Xanthomonas campestris* pv. *campestris* 592 FCKU found to be the most sensitive to influence of *Pseudonadsoniella brunnea* culture fluid. The growth of this bacterium on the Petri dishes was not observed within our study (Fig. 3).

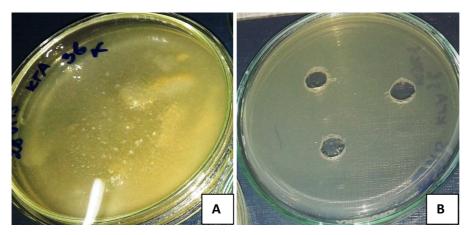


Figure 3. Effect of *Pseudonadsoniella brunnea* culture fluid on the test culture of phytopathogenic bacterium *Xanthomonas campestris* pv. *campestris* 592 FCKU. A – control, B – absence of growth (the reverse side of the Petri dish)

Thus, the different species and strains of test cultures of pathogenic fungi and phytopathogenic bacteria found to show different sensitivity to the effect of the compounds studied, as evidenced by data on the diameter of the zones of growth delay (Tables 1, 2) similarly to our previous study of the resistance of Fusarium oxysporum (strains 150 FCKU and 328 FCKU) [45] and Gibberella fujikuroi (anamorph: Fusarium verticillioides; strains 234 FCKU, 333 FCKU, 338 FCKU and 434 FCKU) to impact of compounds studied [46]. Our data on the diameter of growth inhibition zones of test cultures of pathogenic fungi were compared also with an antagonistic effect of fungi of the genus Trichoderma on pathogenic fungi of the genera Gibberella and Fusarium after data of other authors [32]. From this comparison the effect of Pseudonadsoniella brunnea culture fluid on pathogenic fungi of the genera Gibberella, Fusarium and Rectifusarium ventricosum can be assessed as highly active. It should be especially emphasized that increased synthesis of pigments by Fusarium poae 339 FCKU, F. poae 340 FCKU, similarly to those in previously investigated by us Gibberella fujikuroi strains 234 FCKU and 434 FCKU influenced by Pseudonadsoniella brunnea culture fluid was observed. It is known that under the influence of various factors, including stress, the fungi of the of genera Gibberella and Fusarium can synthesize important biologicallyactive pigments of interest in connection with a wide range of biological activity (antibacterial, antifungal, phytotoxic, insecticidal, and cytotoxic etc.) [49, 50].

According to our data obtailed within this study the impact of *Pseudonadsoniella brunnea* culture fluid on the test cultures of the phytopathogenic bacteria is evaluated as antibacterial one.

Thus, fungicidal and bactericidal effects of melanin producer, black yeast Pseudonadsoniella brunnea culture fluid on test cultures of pathogenic fungi Gibberella baccata (anamorph: Fusarium lateritium), Gibberella intricans (anamorph: Fusarium gibbosum), Gibberella pulicaris (anamorph: Fusarium sambucinum), Fusarium incarnatum, Fusarium solani, Fusarium poae, Rectifusarium ventricosum and phytopathogenic bacteria Agrobacterium tumefaciens, Clavibacter michiganensis subsp. michiganensis, Pectobacterium carotovorum subsp. carotovorum, Pseudomonas syringae pv. syringae, Xanthomonas campestris pv. campestris is for the first time established within this study. This line of research is rather promising from our point of view.

Because one way of controlling phytopathogens is through the use of antagonistic microorganisms. The usage of metabolic substances of microorganisms which can inhibit the growth of other microorganisms (including pathogenic microscopic fungi and bacteria) as

biological methods of pest management of agricultural production will result in decreasing a

negative impact on the environment that caused by biocidal chemicals.

The yeast-like fungus Pseudonadsoniella brunnea used by us within this study

synthesizes and excretes into a culture fluid dark pigment melanin, which is abioactive

compound. Earlier with the usage of the methods of the molecular phylogeny it was

shown that Pseudonadsoniella brunnea shows the highest similarity to the members of the

family Meripilaceae (Poliporales, Basidiomycota) [44]. The polyphenolcarbon complex of

the Pseudonadsoniella brunnea culture fluid is a more than 90% of its.

It is also known that many polymeric and monomeric compounds, including

polyphenols, tyrosine, indole, etc., play a significant role in the formation of melanin by

various fungi [51-53]. It is known that polyphenolic compounds exhibit a strong biological

effect, such as restorative, antioxidant, anti-inflammatory and wound healing. The effect of

natural polyphenols derived from different parts of plant species on the growth of fungi and

bacteria (especially on growth of bacteria of the Pseudomonas syringae complex) has been

studied by many researchers [54, 55].

Our previous studies shown that the melanin, which is produced by Antarctic black

yeast-like fungi *Pseudonadsoniella brunnea*, with oral administration has cyto-protective,

stress-protective, radio-protective, antioxidant, anti-inflammatory, immunomodulatory and

antitumor properties [56].

The first data on antioxidant, antibacterial, fungistatic wound healing properties of the

gel containing 0.05% melanin ("Melanin-gel"), which was synthesized by P. brunnea are

obtained by us earlier too. Biocidal effect of 0.05% "Melanin-gel" was revealed on test

cultures of bacteria Staphylococcus aureus and Pseudomonas aeruginosa. In a test culture of

Candida yeasts melanin gel produced the fungistatic effect (there were areas of growth

retardation, where a decrease in the intensity of yeast growth was observed) [57].

Thus, the use of the culture fluid of the yeast-like fungus *Pseudonadsoniella brunnea*,

which synthesizes and excretes dark pigment melanin into a culture, against pathogenic fungi

and bacteria is effective and represents promising results as for antifungal and antibacterial

protection.

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Antimicrobial activity of papain

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Arun Kumar Srivastava¹, Vinay Kumar Singh²

¹ Department of Zoology, Shri Guru Goraksha Nath Degree College, Maharajganj-273151, U.P. India

² Malacology Laboratory, Department of Zoology, DDU Gorakhpur University, Gorakhpur - 273009, U.P. India

*Corresponding Author: Dr. Arun Kumar Srivastava; Phone: +91-9792250710 (Mobile);

E-mail: aksgkp5@gmail.com

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ABSTRACT

Microbial contamination is one of the main problems that may affect the shelf life of food and may also cause consumer illness. Therefore, many chemicals are used as preservatives to increase the safety and shelf life of food products. Enzymes are biocatalysts that increase the rate of otherwise slow reactions by decreasing the activation energy, without undergoing any net change in their structures at the end of a reaction. Papain enzyme belongs to the papain superfamily, as a proteolytic enzyme. Papain is of crucial importance in many vital biological processes in all living organisms responsible for breaking down proteins. Besides, enzymes in the latex are also involved in protection of the plant against predator attack. The presence of bacteriolytic action in the lattices of Carica papaya confirms the fact that bacteriolytic and proteolytic enzymes act in unison to degrade undesirable proteins. It preferentially cleaves peptide bonds involving basic amino acids, particularly arginine, lysine and residues following phenylalanine. Papain is obtained by cutting the skin of the unripe papaya and then collecting and drying the latex which flows from the cut. The greener the fruit, more active is the papain. The unique structure of papain gives its functionality that helps to understand how this proteolytic enzyme works and it's useful for a variety of purposes.

Keywords: Cystein protiases; Peptide; Antibacterial; *Carica papaya*; Proteolytic.

1. INTRODUCTION

Microbial contamination is one of the main problems that may affect the shelf life of food and may also cause consumer illness. Therefore, many chemicals are used as preservatives to increase the safety and shelf life of food products [1]. As a result of the increased awareness of the consumer about the deleterious effects of chemical preservatives and the increasing preference for natural components, researchers have focused on the generation of natural additives that demonstrate antimicrobial significance to be used in the food industry [2]. Enzymes are biocatalysts that increase the rate of otherwise slow reactions by decreasing the reactions activation energy, without undergoing any net change in their structures at the end of a reaction [3]. They are mostly protein in nature and mediate all synthesis and degradation reactions carried out by living organisms [3]. Papain is an endolytic plant cysteine protease enzyme which is isolated from papaya (Carica papaya L.) latex [4]. Papain enzyme belongs to the papain superfamily, as a proteolytic enzyme, papain is of crucial importance in many vital biological processes in all living organisms [5]. Papain shows extensive proteolytic activity towards proteins, short chain peptides, amino acid esters and amide links and is applied extensively in the fields of food and medicine [6]. Papain is also used in topical formulations as a proteolytic debriding agent for the treatment of open, extensive wounds and burnings. It is also employed as an enhancer for cutaneous permeation of active compounds, chemical peeling and as a progressive depilatory agent [7]. Gurudatta et al. [8] reported that papain preferentially cleaves peptide bonds involving basic amino acids, particularly arginine, lysine and residues following phenylalanine. The unique structure of papain gives its functionality that helps to understand how this proteolytic enzyme works and it's useful for a variety of purposes [9]. Gartika et al. [10] reported that papain is bactericidal, bacteriostatic, anti-inflammatory and debridement material and shows a broad proteolytic activity against the protein, short chain peptides, amino acid ester and amid. This chapter addresses mainly structural features of enzyme, the biological importance and anti-microbial action of papain.

2. PROPERTIES AND STRUCTURE

The globular protein, the papain is a single chain protein with molecular weight of 23,406 DA and consists of 212 amino acid with four disulfide bridges and catalytically important residues in the following positions Gln19, Cys25, His158 and His159 [11]. It is very stable even at elevated temperatures [6]. Papain is unusually defiant to high concentrations of denaturing agents, such as, 8M urea or organic solvent like 70% EtOH [5]. Amanu [12] reported the optimum pH for activity of papain is in the range of 3.0-9.0 which varies with different substrate. Edwin et al. [13] reported that cysteine proteases in papain superfamily are usually consisting of two well-defined domains which provide an excellent system for studies in understanding the folding-unfolding behavior of proteins. The protein is stabilized by three disulfide bridges in which the molecule is folded along these bridges creating a strong interaction among the side chains which contributes to the stability of the enzyme [11, 14]. Its three-dimensional structure consists of two distinct structural domains with a cleft between them. This cleft contains the active site, which contains a catalytic diad that has been similar to the catalytic triad of chymotrypsin [15]. Ezekiel and Florence [6] reported that Histidine-159. Aspartate-158 was thought to play a role analogous to the role of aspartate in the serine protease catalytic triad, but that has since then been disproved. Papain molecule has an all a domain and an anti parallel b-sheet domain [16]. Amanu [12] reported the enzymatic activity of papain may be influenced by environmental conditions such as temperature, light, oxygen, humidity and packing. This enzyme is more stable and active in pH 5.0-7.0. Atalla et al. [17] reported the stability of the enzyme has been investigated at different temperatures and results have confirmed the decrease in its activity with the temperature increase. Catalytic activity of papain involves hydrolysis of proteins with broad specificity for peptide bonds, but preference for an amino acid bearing a large hydrophobic side chain at the P2 position while does not accept Val in P1 [18]. The enzyme has been reported to be generally more stable in hydrophobic solvents and at lower water contents and can catalyze reactions under a variety of conditions in organic solvents with its substrate specificity little changed from that in aqueous media [19]. In general, native proteins have a hydrophobic core and charged and/or polar group on the surface. The hydrophobic core helps to stabilize the tertiary structure of the protein by hydrophobic interaction while the outer polar surfaces preferentially interact with the exterior aqueous medium [6, 20].

Figure 1. Structure of papain.

3. MODE OF ACTION

The mechanism in which the function of papain is made possible is through the cysteine-25 portion of the triad in the active site that attacks the carbonyl carbon in the backbone of the peptide chain freeing the amino terminal portion. As this occurs throughout the peptide chains of the protein, the protein breaks apart. The mechanism by which it breaks peptide bonds involves deprotonation of Cys-25 by His-159. Asparagine-175 helps to orient the imidazole ring of His-159 to allow this deprotonation to take place. Although far apart within the chain, these three amino acids are in close proximity due to the folding structure. It is though these three amino acids working together in the active site that provides this enzyme with its unique functions. Cys-25 then performs a nucleophilic attack on the carbonyl carbon of a peptide backbone [11, 21]. In the active site of papain, Cys -25 and His -159 are thought to be catalytically active as a thiolate-imidazolium ion pair. Papain can be efficiently inhibited by peptidyl or non-peptidyl N-nitrosoanilines [22]. The inactivation is due to the formation of a stable S-NO bond in the active site (S nitroso- Cys25) of papain [23]. Recently Srivastava and Singh [4] noted that feeding of bait formulation of papain with attractant (starch or serine) have sufficient molluscicidal activity against L. acuminata. It significantly reduced the fecundity of the snail and inhibited the AChE activity in the nervous tissue simultaneously. It seems that after sublethal treatment of papain caused the decrease in the level of serotonin and inhibits prostaglandins synthesis by inhibiting 5-lipoxygenase and leukotriene directly or indirectly CDCs. Possibly, the active molluscicidal component papain affect the CDCs and reduce the release of ovulation hormone, resulting a decrease in the fecundity of treated snail.

4. PAPAIN SUPER FAMILY

Cysteine proteases of the papain super family are widely distributed in nature [24]. They can be found in both prokaryotes and eukaryotes e.g. bacteria, parasites, plant, invertebrates and vertebrates [25]. The papain family contains peptidases with a wide variety of activities, including endopeptidases with broad specificity (such as papain), endopeptidases with very narrow specificity (such as glycyl endopeptidases), aminopeptidases, a dipeptidylpeptidase, and peptidases with both endopeptidase and exopeptidase activities (such as cathepsins B and H) [26]. Dubey et al. [24] reported that enzymes of papain family are found in a wide variety of life forms: baculovirus, eubacteria like *Porphyromonas* and *Lactococcus*, yeast, and probably all protozoa, plants, and animals. Xiang et al. [27] reported that lysosomal cysteine proteases, also known as cysteine cathepsins (Cats), include Cat B, Cat H, Cat S, Cat K, Cat O/2, Cat F, Cat W and Cat U and also belong to the papain family sharing similar protein structure and mechanism of action. However, slight structural differences make these enzymes distinct with respect to their substrate specificity and regulation. Marksmann et al. [28] reported that cathepsins are synthesized as 30-50 kDa precursor proteins, which are glycosylated and phosphorylated in the Golgi apparatus. They are processed in the lysosomes to their active forms by one or more proteolytic cleavage. The optimum activity of cathepsins is pH 5.0-6.5, although they can hydrolyze large substrates also at neutral pH [29]. Dubay et al. [24] reported that the pH dependent activity of cathepsins is rather complex and depends not only on the microenvironment and the nature of the conformation of the substrate, but also on the presence or absence of stabilizing factors. Most of these papain-like enzymes are relatively small proteins with mass values in the range 20-35 kDa [30]. Disturbance of the normal balance of enzymatic activity of lysosomal cysteine proteases may lead to pathological conditions, and these proteases have been found to be involved in many such cases [31]. The participation of these enzymes in various diseases seems to be restricted to their proteolytic function outside the lysosomes, after secretion from lysosomes or after translocation into different intracellular granules [32]. The resulting uncontrolled proteolysis is a result of an imbalance between catalytically active proteases and their natural inhibitors, and can be observed in e.g. inflammation and tumor growth, although these processes are very complex [33]. Cysteine proteases of the papain family have been reported in bacteria as well [34]. How et al. [35] reported that proteolytic enzymes produced by Porphyromonas gingivalis are important virulence factors of this periodontopathogen. In the periodontal

disease proteolytic enzymes are produced in large quantities. It has been shown that these proteases can directly or indirectly degrade constituents of the periodontal tissues, destroy host defense elements, dysregulate coagulation and complement kallikerinkinin cascades [36, 37]. Recently, proteases belonging to two catalytic classes and produced by *P. gingivalis* have been identified. One enzyme is described as an Arg-X specific proteinase and another is Lys-X specific [38]. After transcription, the synthesis of the enzymes as inactive precursors, which are subject to several steps of post-translation modification, is the next regulatory mechanism for the papain like cysteine proteases [39]. Bravo et al. [40] reported that in the case of the lysosomal enzymes, the signal peptides are removed when the molecules pass into the lumen of the endoplasmic reticulum, and glycosylation, phosphorylation and formation of disulfides then take place in the golgi complex. He also reported that in the lysosomes, the proenzymes are dephosphorylated and converted to the active enzymes by limited hydrolysis. The N terminal proregions of the enzymes, which are removed during this final maturation step, act as a potent reversible inhibitor against the mature enzymes [41]. The propeptides of the plant enzymes act as their inhibitors [42]. The crystal structure procaricain have shown that the structure of the mature enzyme is already formed in its zymogens and the propeptides prevent the enzymatic activity by blocking the active site cleft using the inhibitory mechanism [24].

5. LOCALIZATION OF PLANT PAPAIN-LIKE CYSTEINE PROTEASES

Cornell and Stelmasiak [43] reported that in plants, they are primarily found in the latex and fruits of plants. In the latter, they are located in the vacuoles, which are plant counterpart of lysosomes, but are also extracellular as in the latices like papaya, figs and in arthopods such as lobsters. Dubey et al. [24] reported that cysteine proteases of plants play a major role in intracellular and extra cellular processes such as development and ripening of fruits, as nutritional reserve, degradation of storage protein in germinating seeds, activation of proenzymes and degradation of defective proteins. Otegui et al. [44] reported that cysteine proteases play an important role in seed germination and have been observed during maturation of storage proteins in *Cannavalia ensiformis, Riccinus communis, Glycine max*. Papain is an endolytic plant cysteine protease enzyme which is isolated from papaya (*Carica papaya* L.) latex. Papain is obtained by cutting the skin of the unripe papaya and then collecting and drying the latex which flows from the cut and the greener the fruit, more active

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is the papain [6]. It preferentially cleaves peptide bonds involving basic amino acids, particularly arginine, lysine and residues following phenylalanine [45]. The unique structure of papain gives its functionality that helps to understand how this proteolytic enzyme works

and it's useful for a variety of purposes [8].

6. ANTI-MICROBIAL ACTIVITY OF PAPAIN

Schmelcher et al. [46] reported that the lysis of micro-organisms by lysozyme and

related enzymes and concluded that the lytic action of lysozyme on bacteria can be ascribed

simply to the dissolution of the rigid cell-wall structures. When the degradation of the wall

occurs in dilute media, the underlying structures of a lysozyme-sensitive bacterial cell will

collapse and the liberation of the cytoplasmic components into the medium will result in the

lysis of the bacterial suspension. Weibull [47] showed that dissolution of the wall of Bacillus

megaterium in the presence of osmotically protective quantities of sucrose was accompanied

by liberation of bacterial protoplasts and only partial optical clearing of the cell suspension.

Although protoplast formation may account for incomplete lysis of bacterial suspensions

when treated with lysozyme under appropriate conditions, it is evident that the walls of

bacteria differ quantitatively in the 'amount' of lysozyme substrate present and such a factor

as this may contribute to the partial optical clearing in dilute media. The cell-wall amino acids

and sugars are not detectable as the free substances in the dialyzable fractions of the digests.

The available evidence thus suggests that lysozyme is splitting the glycosidic bonds of the

cell-wall amino sugars, liberating the disaccharide of acetylglucosamine and acetyl 'muramic

acid' as the simplest product, together with more complex fragments which differ

quantitatively rather than qualitatively in their chemical composition.

Gartika et al. [10] reported that the antibacterial activity of papain against

Streptococcus mutans ATCC 25175 and concluded that this enzyme shows a broad

proteolytic activity against the protein, short chain peptides, amino acid ester and amid,

including bacterial cell wall. The purpose of this study is to produce a proper papain

concentration to inhibit the growth of or kill *Streptococcus mutans*. The type of research is an

experimental laboratory by determining the minimum inhibitory concentration (MIC) and

minimal bactericidal concentration (MBC) with a dilution method, and measured using a

microplate reader papain's minimum inhibitory concentration (MIC) papain against

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Streptococcus mutans was 7.5% and the minimum bactericidal concentration (MBC) was 15%. Papain has antibacterial activity to Streptococcus mutans.

Bharwajd et al. [48] compared the antimicrobial activity of 2% chlorhexidine (100%), extract of *Morinda citrifolia* (86.02%), aloe vera gel (78.9%), papain gel (67.3%) and calcium hydroxide (64.3%) against *Enterococcus faecalis*. Phankhongsap et al. [49] compared the effectiveness of the antimicrobial between papain with mangosteen pericarp extract and papain with propolis extract against mixture *Streptococcus gordonii* and *Enterococcus faecalis* with the inhibiton zonw size11.25±0.66 and 10.42±0.72 mm, respectively. Minimum inhibitory concentration of the two materials were 25 mg/ml, while the MBC were 50 mg/ml.

Kumar et al. [50] tested papaya leaf extracts against human pathogenic microbes. Bacteria such as *Bacillus subtilis*, *Clostridium tetanus*, *Escherichia coli*, *Proteus vulgaris*, *Stapylococcus aureusand* fungi such as *Aspergillus conicus*, *spergillus flavus*, *Aspergillus niger*, *Aspergillus sulphureus* and *Rhizopus* by agar well diffusion method. All the leaf extracts of *Carica papaya* L. exhibited greater activity towards bacteria and fungi. The extract demonstrated higher activities against all the bacteria and fungi tested, with the highest activity (acetone extract of 13 mm zone of inhibition) demonstrated against *Staphylococcus aureus* and (ethanol extract of 18 mm zone of inhibition) demonstrated against *Aspergillus flavus*. *Carica papaya* may be used for the treatment of gastroenteritis, uretritis, otitis media, dengue fever, typhoid fever and wound infections.

Krishna et al. [51] reviewed on nutritional, medicinal and pharmacological properties of papaya and find that the seed of papaya has antimicrobial activity against *Trichomonas vaginalis* trophozoites. The report suggests the use of papaya seed in urinogenital disorder like trichomoniasis with care to avoid toxicity. The seed and pulp of papaya was shown to be bacteriostatic against several enteropathogens such as *Bacillus subtilis, Enterobacter cloacae, Escherichia coli, Salmonella typhi, Staphylococcus aureus, Proteus vulgaris, Pseudomonas aeruginosa and Klebsiella pneumoniae by the agar cup plate method 24. Purified extracts from ripe and unripe fruits also produces very significant antibacterial activity on <i>S. aureus, Bacillus cereus, E. coli, P. aeruginosa and Shigella flexneri.* The aqueous extract of fruit exhibited antimicrobial activity and promoted significant wound healing in diabetic rats. The seeds of irrespective stage of fruit maturity have bacteriostatic activity on Gram positive and Gram negative organisms, which could be useful in treating chronic skin ulcers. The papaya seed macerate has a clinical potential on conjugal R plasmid transfer from *Salmonella typhimurium* to *Escherichia coli*, on in vitro and in the digestive tract of genotobiotic mice. Herbal formulations containing papaya leaves and root or leaves alone as one of

the constituent has antibacterial activity against *Salmonella typhi*, *S. paratyphi* and *S. typhimurium*; however, water, acetone and ethanol extract of papaya leaves showed no microbicidal activity.

Anibijuwon and Udeze [52] reported the antimicrobial activity of Carica papaya (pawpaw leaf) on some pathogenic organisms of clinical origin from South-Western Nigeria and investigated for antibacterial activity against some human pathogenic bacteria using the agar diffusion method. The aqueous extracts of the root extracts did not show significant activity, but the organic extracts had significant activity with the methanol extracts demonstrating the highest activity against the test bacteria. The root extracts demonstrated higher activities against all the Gram-positive bacteria than the gram-negative bacteria tested, with the highest activity (14 mm zone of inhibition) demonstrated against Pseudomonas aeruginosa while the aqueous leaf extract showed pronounced inhibition demonstrating higher activities against the test bacteria than the organic solvents. The extracts demonstrated higher activities against all the Gram-positive bacteria than the Gram-negative bacteria tested, with the highest activity (4.2 mm zone of inhibition) demonstrated against *Pseudomonas* aeruginosa. Increase in temperature enhanced the activity of the extracts, while alkaline pH decreased the activity. The Minimum Inhibitory Concentration (MIC) and Minimum Bactericidal Concentration (MBC) of the root extracts ranged between 50-200 mg/ml. Preliminary phytochemical analyses showed that the extracts contain alkaloids, tannins, saponins, glycosides and phenols. Carica papaya may be used for the treatment of gastroenteritis, uretritis, otitis media and wound infections.

Kumar et al. [53] reported the antifungal medicinal properties of *Carica papaya*. The effects of different concentrations of alcoholic extract of *Carica papaya* (root, shoot and seed) on the radial growth of plant against the pathogenic fungi viz. *Aspergillus niger*, *Aspergillus flavus*, *Candida albicans* and *Microsporum fulvum*. That with the increase in concentrations the rate of growth inhibition also increases. Observation further shows that like root extract growth is also inhibited in the presence of shoot and seed alcoholic extract under culture medium. Further shows that the growth of these fungi inhibits more in presence of higher concentrations as compared to lower concentrations of extract.

Islam et al. [54] experimented the evaluation of antibacterial activities of latex of Caricaceae (*Carica papaya* L.) He has reported that latex was evaluated against one Grampositive bacterium *Bacillus subtilis* and three Gram-negative pathogenic bacterial strains as *Escherichia coli*, *Agrobacterium sp* and *Rhizobium sp*. Ciprofloxacin was used as a control for investigating the bacterial species. Antibacterial activity was expressed in terms of the

radius of zone inhibition. Latex of this plant was tested in seven doses (1, 2, 5, 7, 10, 15 and 20 mg/disk) and it was found that the antibacterial activity was dose dependent and a significant difference was also observed in case of different bacterial stains. The results demonstrated noticeable inhibition of the bacterial growth against the tested organisms. In case of *Agrobacterium* sp. 20 mg of latex showed the average of 20.66 mm zone of inhibition and for *E. coli* this value was 16 mm for the same concentration of latex. The rest of the two bacterial species showed comparative resistance to papaya latex.

Coello et al. [55] assessed anti-protozoan activity of crude $Carica\ papaya$ seed extract gainst $Trypanosoma\ cruzi$ and in order to determine the $in\ vivo$ activity against the protozoan $Trypanosoma\ cruzi$, two doses (50 and 75 mg/kg) of a chloroform extract of $Carica\ papaya$ seeds were evaluated compared with a control group of allopurinol. A significant reduction (p < 0.05) in the number of blood trypomastigotes was observed in animals treated with the evaluated doses of the $C.\ papaya$ extract in comparison with the positive control group (allopurinol 8.5 mg/kg). Parasitemia in animals treated with the fatty acids mixture was also significantly reduced (p < 0.05), compared to negative control animals. These results demonstrate that the fatty acids identified in the seed extracts of $C.\ papaya$ (from ripe fruit) are able to reduce the number of parasites from both parasite stages, blood trypomastigote and amastigote (intracellular stage).

Coello et al. [56] evaluated the anti-protozoan activity of the chloroform extract of *Carica papaya* seeds during the sub-acute and chronic phase of infection of *Trypanosoma cruzi*, doses of 50 and 75 mg/kg, including a mixture of their main components (oleic, palmitic, and stearic acids). Subsequently, doses of 50 and 75 mg/kg in mice during the chronic phase of infection (100 dpi) were also evaluated. It was found that chloroform extract was able to reduce the amastigote nests numbers during the subacute phase in 55.5 and 69.7% (p > 0.05) as well as in 56.45% in animals treated with the mixture of fatty acids. Moreover, the experimental groups treated with 50 and 75 mg/kg during the chronic phase of the infection showed a significant reduction of 46.8 and 53.13% respectively (p < 0.05). It is recommended to carry out more studies to determine if higher doses of chloroformic extract or its administration in combination with other anti-chiasmic drugs allows a better response over the intracellular stage of T. cruzi in infected animal models and determine if the chloroform extract of C. papaya could be considered as an alternative for treatment during the indeterminate and chronic phase of the infection.

Kovendan et al. [57] reported the anti-malarial activity of the ethanol leaf extract of *Carica papaya*, blood stages of CQ-sensitive and CQ resistant strains against *Plasmodium*

falciparum as target species. The larvae and pupae values of 1st to 4th instars values of

 $LC_{50} = 3.65\%$, 4.28%, 5.41%, 6.70%, and 7.50%, respectively. The LC_{90} to good 9.1%,

11.75%, 13.53%, 16.36%, and 16.92%, respectively. These four concentrations (25, 50,100

and 150 µg/ml) of ethanol leaf extracts exhibited promising inhibitory activity against the CQ

sensitive strain with (IC $_{50}$) values 40.75%, 36.54%, 25.30%, and 18.0% and in CQ resistant

50.23%, 32.50%, 21.45%, and 23.12% against *P. falciparum*.

Quintal et al. [58] reported the antifungal activity in ethanolic extracts of Carica

papaya L. cv. maradol leaves and seeds antifungal effectiveness was determined by

challenging the extracts (LE, SRE, SUE) from the best extraction treatment against three

phyto-pathogenic fungi: Rhizopus stolonifer, Fusarium spp. and Colletotrichum

gloeosporioides. The leaf extract exhibited the broadest action spectrum. The MIC 50 for the

leaf extract was 0.625 mg/ml for Fusarium spp. and 10 mg/ml for C. gloeosporioides, both

equal to approximately 20% mycelial growth inhibition. Ethanolic extracts from Carica

papaya L. cv. maradol leaves are a potential source of secondary metabolites with antifungal

properties.

7. CONCLUSION

In conclusion, plant-based antimicrobials have enormous therapeutic and preferential

potential. They can serve the desired purpose with lesser side effects that are often associated

with synthetic antimicrobials. Papain is found naturally in papaya which is a versatile plant

having number of uses, enzymatic properties and antimicrobial activity demonstrated in this

chapter. Antimicrobial activity of papain is an indication that there is possibility of sourcing

alternative antibiotic substances in these plants for the development of newer antibacterial

agents.

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3

Continuous research for natural drugs with potential non-resistance antimicrobial activity and reduced adverse effects

Rasha Y. Elbayaa^{1,2}, Ibrahim A. Abdelwahab³

¹ Department of Pharmaceutical Chemistry, Faculty of Pharmacy, University of Alexandria, Alexandria

21521, Egypt

² Department of Analytical & Pharmaceutical Chemistry, Faculty of Pharmacy & Drug Manufacturing, Pharos

University in Alexandria, 21311, Egypt

³ Department of Microbiology and Immunology, Faculty of Pharmacy & Drug Manufacturing, Pharos

University in Alexandria, 21311, Egypt

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ABSTRACT

Resistant-microbes are found in people, animals, food, and the environment (in water, soil and air). Poor infection control, inadequate sanitary conditions and inappropriate food-handling encourage the spread of antimicrobial resistance also Antibiotic resistance is present worldwide so Patients with infections caused by drug-resistant bacteria are at increased risk of worse clinical outcomes causing death, in addition, this consumes more health-care resources than patients infected with non-resistant strains of the same bacteria. Nature is a generous source of a number of compounds with potential application for the treatment of several diseases including the infectious diseases. The presently investigated natural products derived from local botanical are promising candidates that could be used against multi drug resistant pathogens with high potency and less side effects.

Keywords: Resistant-microbes; Natural products; Structure-activity relationship.

1. INTRODUCTION

Antimicrobial resistance has been attributed as a challenging problem worldwide. It

happens when microorganisms (such as bacteria, fungi, viruses, and parasites) change when

they are exposed to antimicrobial drugs (such as antibiotics, antifungals, antivirals,

antimalarials, and anthelmintics). Microorganisms that develop antimicrobial resistance are

sometimes referred to as "superbugs". As a result, the medicines become ineffective and

infections persist in the body, increasing the risk of spread to others [1].

New resistance mechanisms are emerging and spreading globally, threatening our

ability to treat common infectious diseases, resulting in prolonged illness, disability, and

death. Also, Antimicrobial resistance increases the cost of health care with lengthier stays in

hospitals and more intensive care required. In addition, antimicrobial resistance is putting the

gains of the Millennium Development Goals at risk and endangers achievement of the

Sustainable Development Goals [1].

Antimicrobial resistance occurs naturally over time, usually through genetic changes.

However, the misuse and overuse of antimicrobials is accelerating this process. In many

places, antibiotics are overused and misused in people and animals, and often given without

professional oversight. Antimicrobial resistant-microbes are found in people, animals, food,

and the environment (in water, soil and air). They can spread between people and animals,

and from person to person. Poor infection control, inadequate sanitary conditions and

inappropriate food-handling encourage the spread of antimicrobial resistance [1].

2. RESISTANCE AMONG DIFFERENT PATHOGENS

2.1. Resistance in Klebsiella pneumoniae

The common intestinal bacteria that can cause life-threatening infections – to a last

resort treatment (carbapenem antibiotics) have spread to all regions of the world.

K. pneumoniae is a major cause of hospital-acquired infections such as pneumonia,

bloodstream infections, and infections in newborns and intensive-care unit patients. In some

countries, because of resistance, carbapenem antibiotics do not work in more than half of

people treated for *K. pneumoniae* infections [1].

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2.2. Resistance in *Escherichia coli*

Resistance in E. coli to one of the most widely used medicines for the treatment of

urinary tract infections (fluoroquinolone antibiotics) is very widespread. There are countries

in many parts of the world where this treatment is now ineffective in more than half of

patients. Colistin is the last resort treatment for life-threatening infections caused by

Enterobacteriaceae which are resistant to carbapenems. Resistance to colistin has recently

been detected in several countries and regions, making infections caused by such bacteria

untreatable [1].

2.3. Resistance in tuberculosis (TB)

WHO estimates that, in 2014, there were about 480 000 new cases of multidrug-

resistant tuberculosis (MDR-TB), a form of tuberculosis that is resistant to the 2 most

powerful anti-TB drugs. Only about a quarter of these (123 000 cases) were detected and

reported. MDR-TB requires treatment courses that are much longer and less effective than

those for non-resistant TB. Globally, only half of MDR-TB patients were successfully treated

in 2014.

Among new TB cases in 2014, an estimated 3.3% were multidrug-resistant. The

proportion is higher among people previously treated for TB, at 20%. Extensively drug-

resistant tuberculosis (XDR-TB), a form of tuberculosis that is resistant to at least 4 of the

core anti-TB drugs, has been identified in 105 countries. An estimated 9.7% of people with

MDR-TB have XDR-TB [1].

2.4. Resistance in malaria

As of July 2016, resistance to the first-line treatment for P. falciparum malaria

(artemisinin-based combination therapies, also known as ACTs) has been confirmed in 5

countries of the Greater Mekong subregion (Cambodia, the Lao People's Democratic

Republic, Myanmar, Thailand and Viet Nam). In most places, patients with artemisinin-

resistant infections recover fully after treatment, provided that they are treated with an ACT

containing an effective partner drug. However, along the Cambodia-Thailand border,

P. falciparum has become resistant to almost all available antimalarial medicines, making

treatment more challenging and requiring close monitoring. There is a real risk that multidrug

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resistance will soon emerge in other parts of the subregion as well. The spread of resistant

strains to other parts of the world could pose a major public health challenge and jeopardize

important recent gains in malaria control [1].

2.5. Resistance in human immunodeficiency virus (HIV)

In 2010, an estimated 7% of people starting antiretroviral therapy (ART) in developing

countries had drug-resistant HIV. In developed countries, the same figure was 10–20%. Some

countries have recently reported levels at or above 15% amongst those starting HIV treatment,

and up to 40% among people re-starting treatment. This requires urgent attention.

Increasing levels of resistance have important economic implications as second and

third-line regimens are 3 times and 18 times more expensive, respectively, than first-line

drugs.

Since September 2015, WHO has recommended that everyone living with HIV

start on antiretroviral treatment. Greater use of ART is expected to further increase ART

resistance in all regions of the world. To maximize the long-term effectiveness of

first-line ART regimens, and to ensure that people are taking the most effective regimen, it is

essential to continue monitoring resistance and to minimize its further emergence and

spread [1].

2.6. Resistance in influenza

Antiviral drugs are important for treatment of epidemic and pandemic influenza. So

far, virtually all influenza A viruses circulating in humans were resistant to one category of

antiviral drugs - M2 Inhibitors (amantadine and rimantadine). However, the frequency of

resistance to the neuraminidase inhibitor oseltamivir remains low (1-2%). Antiviral

susceptibility is constantly monitored through the WHO Global Influenza Surveillance and

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Response System [1].

3. MAJOR BACTERIAL RESISTANCE STRATEGIES

There are four major bacterial resistance strategies:

3.1. Prevention of the antimicrobial from reaching its target by reducing its ability to

penetrate into the cell

Antimicrobial compounds almost always require access into the bacterial cell to reach

their target site where they can interfere with the normal function of the bacterial organism.

Porin channels are the passageways by which these antibiotics would normally cross the

bacterial outer membrane. Some bacteria protect themselves by prohibiting these

antimicrobial compounds from entering past their cell walls. For example, a variety of Gram-

negative bacteria reduce the uptake of certain antibiotics, such as aminoglycosides and beta

lactams, by modifying the cell membrane porin channel frequency, size, and selectivity.

Prohibiting entry in this manner will prevent these antimicrobials from reaching their intended

targets that, for aminoglycosides and beta lactams, are the ribosomes and the penicillin-

binding proteins (PBPs), respectively [2].

This strategy has been observed in:

• Pseudomonas aeruginosa e.g. against imipenem (a beta-lactam antibiotic);

• Enterobacter aerogenes and Klebsiella pneumoniae against imipenem;

• Glycopeptide intermediate-resistant Staphylococcus aureus so-called "GISA" strains

with thickened cell wall trapping vancomycin/teicoplanin;

• Many Gram-negative bacteria against aminoglycosides;

• Many Gram-negative bacteria against quinolones.

3.2. Inactivation of antimicrobial agents via modification or degradation

Another means by which bacteria preserve themselves is by destroying the active

component of the antimicrobial agent. A classic example is the hydrolytic deactivation of the

beta-lactam ring in penicillins and cephalosporins by the bacterial enzyme called beta

lactamase. The inactivated penicilloic acid will then be ineffective in binding to PBPs

(penicillin binding proteins), thereby protecting the process of cell wall synthesis. This

strategy has also been observed in:

• Enterobacteriaceae against chloramphenicol (acetylation)

• Gram negative and Gram positive bacteria against aminoglycosides (phosphorylation,

adenylation, and acetylation) [3].

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3.3. Expulsion of the antimicrobial agents from the cell via general or specific efflux

pumps

To be effective, antimicrobial agents must also be present at a sufficiently high

concentration within the bacterial cell. Some bacteria possess membrane proteins that act as

an export or efflux pump for certain antimicrobials, extruding the antibiotic out of the cell as

fast as it can enter. This results in low intracellular concentrations that are insufficient to elicit

an effect. Some efflux pumps selectively extrude specific antibiotics such as macrolides,

lincosamides, streptogramins and tetracyclines, whereas others (referred to as multiple drug

resistance pumps) expel a variety of structurally diverse anti-infectives with different modes

of action e.g. the qac genes which pump out chlorhexidine, propamidine and quaternary

ammonium agents [4]. This strategy has also been observed in:

• E. coli and other Enterobacteriaceae against tetracyclines;

• Various members of the *Enterobacteriaceae* against chloramphenicol;

• Staphylococci against macrolides and streptogramins;

• Staphylococcus aureus and Streptococcus pneumoniae against fluoroquinolones.

3.4. Modification of the antimicrobial target within the bacteria

Some resistant bacteria evade antimicrobials by reprogramming or camouflaging

critical target sites to avoid recognition. Therefore, in spite of the presence of an intact and

active antimicrobial compound, no subsequent binding or inhibition will take place. This

strategy has been observed in:

• Staphylococci against methicillin and other beta-lactams (changes or acquisition of

different PBPs that do not sufficiently bind beta-lactams to inhibit cell wall synthesis);

• Gram-positive cocci: erythromycin-resistant methylase is encoded by erm genes and

causes structural changes to rRNA which prevent macrolide binding and allow

synthesis of bacterial proteins to continue;

• Enterococci against vancomycin (alteration in cell wall precursor components to

decrease binding of vancomycin);

• Mycobacterium spp. against streptomycin (modification of ribosomal proteins or of

16s rRNA);

• Various microbes which develop mutations in RNA polymerase resulting in resistance

to the rifamycins e.g. Staphylococcus spp;

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• members of *Enterobacteriaceae* with mutations in DNA gyrase resulting in resistance to quinolones [5].

4. MECHANISMS OF RESISTANCE AGAINST DIFFERENT ANTIMICROBIAL CLASSES

Table 1. Mechanisms of resistance against different antimicrobial classes [6]

Antimicrobial	Mechanism	Specific means to achieve	т
class	of resistance	resistance	Examples
		Destruction of beta-lactam rings	Resistance
		by beta-lactamase enzymes.	of staphylococci
		With the beta-lactam ring	to penicillin;
	Enzymatic	destroyed, the antibiotic will	Resistance
	destruction	no longer have the ability to	of Enterobacteriaceae
Beta-lactams		bind to PBP (Penicillin-binding	to penicllins,
Examples:		protein), and interfere with cell	cephalosporins,
penicillin,		wall synthesis	and aztreonam
ampicillin,		Changes in penicillin binding	
mezlocillin,		proteins. Mutational changes in	Resistance
peperacillin,	Altered	original PBPs or acquisition of	of staphylococci
cefazolin,	target	different PBPs will lead to inability	to methicillin
cefotaxime,		of the antibiotic to bind to the PBP	and oxacillin
ceftazidime,		and inhibit cell wall synthesis	
aztreonam,		Porin channel formation is	Desire
imipenem		decreased. Since this is where beta-	Resistance of Enterobacter
	Decreased	lactams cross the outer membrane	
		to reach the PBP of Gram-negative	aerogenes, Klebsiella
	uptake	bacteria, a change in the number or	pneumoniae and Pseudomonas
		character of these channels can	
		reduce betalactam uptake	aeruginosa to imipenem
		Alteration in the molecular	
Glycopeptides	Altered target	structure of cell wall precursor	Resistance
Example:		components decreases binding of	of enterococci
vancomycin		vancomycin so that cell wall	to vancomycin
		synthesis is able to continue	

Antimicrobial	Mechanism	Specific means to achieve	E
class	of resistance	resistance	Examples
Aminoglyosides Examples: gentamicin, tobramycin, amikacin, netilmicin, streptomycin, kanamycin	Enzymatic modification	Modifying enzymes alter various sites on the aminoglycoside molecule so that the ability of this drug to bind the ribosome and halt protein synthesis is greatly diminished or lost entirely	Resistance of many Gram-positive and Gram negative bacteria to aminoglycosides
	Decreased uptake	Change in number or character of porin channels (through which aminoglycosides cross the outer membrane to reach the ribosomes of gram-negative bacteria) so that aminoglycoside uptake is diminished	Resistance of a variety of Gram-negative bacteria to aminoglycosides
	Altered target	Modification of ribosomal proteins or of 16s rRNA. This reduces the ability of aminoglycoside to successfully bind and inhibit protein synthesis	Resistance of <i>Mycobacterium</i> spp. to streptomycin
Quinolones Examples: ciprofloxacin, levofloxacin, norfloxacin, lomefloxacin	Decreased uptake	Alterations in the outer membrane diminishes uptake of drug and/or activation of an "efflux" pump that removes quinolones before intracellular concentration is sufficient for inhibiting DNA metabolism	Resistance of Gram negative and staphylococci (efflux mechanism only) to various quinolones
	Altered target	Changes in DNA gyrase subunits decrease the ability of quinolones to bind this enzyme and interfere with DNA processes	Gram negative and Gram positive resistance to various quinolones

5. TEST METHODS TO DETECT ANTIMICROBIAL RESISTANCE

There are several antimicrobial susceptibility testing methods available today, and

each one has their respective advantages and disadvantages. They all have one and the same

goal, which is to provide a reliable prediction of whether an infection caused by a bacterial

isolate will respond therapeutically to a particular antibiotict reatment. Some examples of

antibiotic sensitivity tesing methods are:

5.1. Dilution methods

The Broth dilution method involves subjecting the isolate to a series of concentrations

of antimicrobial agents in a broth environment. Microdilution testing uses about 0.05 to

0.1 ml total broth volume and can be conveniently performed in a microtiter format.

Macrodilution testing uses broth volumes at about 1.0 ml in standard test tubes. For both of

these broth dilution methods, the lowest concentration at which the isolate is completely

inhibited (as evidenced by the absence of visible bacterial growth) is recorded as the minimal

inhibitory concentration or MIC. The MIC is thus the minumum concentration of the

antibiotic that will inhibit this particular isolate. The test is only valid if the positive control

shows growth and the negative control shows no growth. A procedure similar to broth

dilution is agar dilution. Agar dilution method follows the principle of establishing the lowest

concentration of the serially diluted antibiotic concentration at which bacterial growth is still

inhibited [7].

5.2. Disk diffusion method

Because of convenience, efficiency and cost, the disk diffusion method is probably the

most widely used method for determining antimicrobial resistance in private veterinary

clinics. A growth medium, usually Mueller-Hinton agar, is first evenly seeded throughout the

plate with the isolate of interest that has been diluted at a standard concentration

(approximately 1 to 2 x 10⁸ colony forming units per ml). Commercially prepared disks, each

of which are pre-impregnated with a standard concentration of a particular antibiotic, are then

evenly dispensed and lightly pressed onto the agar surface. The test antibiotic immediately

begins to diffuse outward from the disks, creating a gradient of antibiotic concentration in the

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agar such that the highest concentration is found close to the disk with decreasing

concentrations further away from the disk. After an overnight incubation, the bacterial growth

around each disc is observed. If the test isolate is susceptible to a particular antibiotic, a clear

area of "no growth" will be observed around that particular disk.

The zone around an antibiotic disk that has no growth is referred to as the zone of

inhibition since this approximates the minimum antibiotic concentration sufficient to prevent

growth of the test isolate. This zone is then measured in mm and compared to a standard

interpretation chart used to categorize the isolate as susceptible, intermediately susceptible or

resistant. MIC measurement cannot be determined from this qualitative test, which simply

classifies the isolate as susceptible, intermediate or resistant [8].

5.3. E-test

E-test (AB Biodisk, Solna, Sweden) is a commercially available test that utilizes a

plastic test strip impregnated with a gradually decreasing concentration of a particular

antibiotic. The strip also displays a numerical scale that corresponds to the antibiotic

concentration contained therein. This method provides for a convenient quantitative test of

antibiotic resistance of a clinical isolate. However, a separate strip is needed for each

antibiotic, and therefore the cost of this method can be high [9].

5.4. Automated antimicrobial susceptibility testing systems

Several commercial systems have been developed that provide conveniently prepared

and formatted microdilution panels as well as instrumentation and automated reading of

plates. These methods are intended to reduce technical errors and lengthy preparation times.

Most automated antimicrobial susceptibility testing systems provide automated inoculation,

reading and interpretation. These systems have the advantage of being rapid (some results can

be generated within hours) and convenient, but one major limitation for most laboratories is

the cost entailed in initial purchase, operation and maintenance of the machinery. Some

examples of these include: Vitek System (bioMerieux, France), Walk-Away System (Dade

International, Sacramento, Calif.), Sensititre ARIS (Trek Diagnostic Systems, East Grinstead,

UK), Avantage Test System (Abbott Laboratories, Irving, Texas), Micronaut (Merlin,

Bornheim-Hesel, Germany), Phoenix (BD Biosciences, Maryland) [10].

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5.5. Mechanism-specific tests

Resistance may also be established through tests that directly detect the presence of a

particular resistance mechanism. For example, beta lactamase detection can be accomplished

using an assay such as the chromogenic cephalosporinase test (Cefinase disk by BD

Microbiology Systems, Cockeysville, MD and BBL DrySlide Nitrocefin, Becton Dickinson,

Sparks, MD) and detection for chloramphenicol modifying enzyme chloramphenicol

acetyltransferase (CAT) may utilize commercial colorimetric assays such as a CAT reagent

kit (Remel, Lenexa, Kansas) [11].

5.6. Genotypic methods

Since resistance traits are genetically encoded, we can sometimes test for the specific

genes that confer antibiotic resistance. However, although nucleic acid-based detections

systems are generally rapid and sensitive, it is important to remember that the presence of a

resistance gene does not necessarily equate to treatment failure, because resistance is also

dependent on the mode and level of expression of these genes. Some of the most common

molecular techniques utilized for antimicrobial resistance detection are as follows:

5.6.1. Polymerase chain reaction (PCR)

This is one of the most commonly used molecular techniques for detecting certain

DNA sequences of interest. This involves several cycles of denaturation of sample DNA,

annealing of specific primers to the target sequence (if present), and the extension of this

sequence as facilitated by a thermostable polymerase leading to replication of a duplicate

DNA sequence, in an exponential manner, to a point which will be visibly detectable by gel

electrophoresis with the aid of a DNA-intercalating chemical which fluoresces under UV

light [12].

5.6.2. DNA hybridization

This is based on the fact that the DNA pyrimidines (cytosine and thymidine)

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specifically pair up with purines (guanine and adenine; or uracil for RNA). Therefore, a

labeled probe with a known specific sequence can pair up with opened or denatured DNA

from the test sample, as long as their sequences complement each other. If this

"hybridization" occurs, the probe labels this with a detectable radioactive isotope, antigenic

substrate, enzyme or chemiluminescent compound. Whereas if no target sequence is present

or the isolate does not have the specific gene of interest, no attachment of probes will

occur [13].

5.6.3. Modifications of PCR and DNA hybridization

With these basic principles, several modifications have been introduced which further

improvement of the sensitivity and specificity of these standard procedures. Examples of such

development were the use of 5'-fluorescence-labeled oligonucleotides, the development of

molecular beacons, development of DNA arrays and DNA chips, among many others [14].

6. NATURAL PRODUCTS ACTING AS ANTIMICROBIALS

Nature is a generous source of a number of compounds with potential application for

the treatment of several diseases including the infectious diseases. The presently investigated

natural products derived from local botanical are promising candidates that could be used

against MDR pathogens. Nevertheless, there is still a vast flora that once systemically

explored could provide additional antimicrobial leads and drugs.

The mechanisms of action of the natural products include the degradation of the cell

wall [16], damaging the cytoplasmic membrane, cytoplasm coagulation [17], damaging the

membrane proteins, increased permeability leading to leakage of the cell contents,

reducing the proton motive force [18], reducing the intracellular ATP pool via decreased

ATP synthesis and augmented hydrolysis that is separate from the increased membrane

permeability and reducing the membrane potential via increased membrane permeability

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[19].

Herein we will discuss the following antimicrobials:

Antibacterial

• Antifungal

Antiviral

Antiprotozoal

6.1. Antibacterial agents

The cinnamon, clove, pimento, thyme, oregano, and rosemary plants had strong inhibitory effect against several bacterial pathogens. It has been also reported that essential oils extracted from some medicinal plants had the antibacterial effects against all the five tested food borne pathogens due to presence of phenolic compounds such as carvacrol, eugenol and thymol [20]. However found that benzoic acids, benzaldehydes and cinnamic acid were able to inhibit the growth of *Listeria monocytogenes* [21].

The antimicrobial activity of garlic, ginger, clove, black pepper and green chilli analyzed on the human pathogenic bacteria viz. *Bacillus sphaericus, Enterobacter aerogenes, E. coli, P. aeruginosa, S. aureus, S. epidermidis, S. typhi* and *Shigella flexneri* and stated that amongst all the tested spices, aqueous garlic extracts was sensitive against all the bacterial pathogens [22]. Similarly, effect of clove extracts on the production of verotoxin by enterohemorrhagic *Escherichia coli* O157:H7 was investigated [23]. Furthermore it was evident from the study that the verotoxin production was inhibited by clove extract. However the effectiveness of cardamom, anise, basil, coriander, rosemary, parsley, dill and angelica essential oil for controlling the growth and survival of pathogenic and saprophytic microorganisms. The results of their study showed that essential oils extracted oregano, basil and coriander plants have inhibitory effect against *Pseudomonas aeruginosa, S. aureus* and *Yersinia enterocolitica* [24].

The effect of oregano essential oils on the behavior of *Salmonella typhimurium* in sterile and naturally contaminated beef fillets stored under aerobic and modified atmospheres also they have concluded that the addition of oregano essential oils checked the reduction in initial population of the tested bacterial pathogens [25]. However the bacterial growth may be inhibited by the ample application of essential oils or their use at high concentrations and their mode of action results in decline of the bacterial cells [26].

The antibacterial activity of essential oils extracted from thyme and mint leaves against the *Staphylococcus aureus*, *Salmonella typhimurium* and *Vibrio parahaemolyticus* and the result showed that all the plants have antibacterial activity against the tested pathogens but the effect of thyme leaves extract was more pronounced compared to other plants [27]. Moreover cinnamon, oregano, clove, pomegranate peel, and grape seed were found effective against *S. enterica* at room temperature, but the clove extracts possess highest antibacterial activity, thyme, sage, myrtle, laurel, and orange essential oils have a potential to inhibit and inactivate four microorganisms in agar and milk medium at different concentrations also the

inhibitory effects of essential oils increased with increasing concentration so it is suggested to

investigate higher essential oils concentrations than were those used in research, and to study

the effects over a longer time period in milk and other available milk products to access the

potential of plant species essential oils as preservatives [28].

6.1.1. Main antibacterial phytochemicals

Plant-derived compounds of therapeutic value are mostly secondary plant metabolites

traditionally used for medicinal purposes. They have a wide activity range, according to the

species, the topography and climate of the country of origin, and may contain different

categories of active principles [29]. Variations in the chemical composition modify their

antimicrobial activity. Some main categories of phytochemicals extracted from medicinal

plants are examined to evaluate their pharmacological activity.

6.1.1.1. *Flavonoids*

Flavonoids, previously called bioflavonoids and included in aromatic compounds, are

phenolic structures ubiquitous in photosynthesizing cells and are commonly found in fruit,

vegetables, nuts, seeds, stems, flowers, tea, wine, propolis and honey. For centuries,

preparations containing these compounds as the principal physiologically active constituents

have been used to treat human diseases. The basic structural feature of flavonoid compounds

is the 2-phenyl-benzopyrane or flavane nucleus, consisting of two benzene rings linked

through a heterocyclic pyrane ring.

In total, there are 14 classes of flavonoids, differentiated on the basis of the chemical

nature and position of substituents on the different rings. The antibacterial properties of

flavonoids are thought to come from the ability to form complexes with both extracellular and

soluble proteins, as well as with bacterial membrane [30].

Kuete demonstrated that among the flavonoids hydroxylating the prenyl groups of

stipulin, the compounds obtained, angusticornin B and bartericin A, had a superior

antimicrobial activity [31]. Thus, the prenyl group plays an important role in the activity of

chalcones. Recently two flavonoids (6-hydroxy-7-methoxyluteolin and the xanthone

8-carboxymethyl-1,5,6-trihydroxy-3-methoxyxanthone) extracted from the leaves of *Leiothrix*

spiralis, a South American plant belonging to the Eriocaulaceae family, showed a promising

activity on Escherichia coli and Pseudomonas aeruginosa [32]. Some flavonoids also

revealed activity against *M. tuberculosis* [33].

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A synergy has been demonstrated between active flavonoids as well as between flavonoids and existing chemotherapeutics, even if the reports of activity in the field of antibacterial flavonoid research are widely conflicting, probably owing to interand intra-assay variation in susceptibility testing [34]. Future optimization of these compounds through structural alteration may allow the development of a pharmacologically acceptable antimicrobial agent or group of agents. Existing structure-activity data suggest that it might be possible, for example, to prepare a potent antibacterial flavanone by synthesizing a compound with halogenation of the B ring as well as lavandulyl or geranyl substitution of the A ring. Also, it is worth noting that by elucidating flavonoid biosynthetic pathways it would be possible to produce structural analogs of active flavonoids through genetic manipulation. Numerous research groups have sought to elucidate the antibacterial mechanisms of action of selected flavonoids; the activity of quercetin has been at least partially attributed to the inhibition of DNA gyrase, whereas sophoraflavone G and (-)-epigallocatechin gallate inhibit cytoplasmic membrane function, and licochalcones A and C inhibit energy metabolism.

6.1.1.2. *Alkaloids*

Alkaloids are heterocyclic nitrogen compounds characterized by different antimicrobial activities. The analysis of the leaf extracts of *Gymnema montanum* and of ethanol extract of *Tabernaemontana catharinensis* root bark revealed an antimicrobial activity [35] in the first case due to an activity depending upon the chemical composition of the extracts and membrane permeability of the microbes, and in the second case linked to indole alkaloids responsible for the observed antibacterial and antidermatophytic activity. Diterpene alkaloids, commonly isolated from the plants of the Ranunculaceae group, had antimicrobial properties [36]. Berberine, an isoquinoline alkaloid, present in roots and stem-bark of *Berberis* species, is a hydrophobic cation widely used in traditional medicine owing to its activity against bacteria, fungi, protozoa and viruses [37]. It accumulates in cells driven by the membrane potential and is an excellent DNA intercalator active on several microorganisms with a target on RNA polymerase, gyrase and topoisomerase IV and on nucleic acid [38].

6.1.1.3. Terpenes

Terpenes compounds are also referred to as isoprenoids and their derivatives containing additional elements, usually oxygen, are called terpenoids. The antibacterial activity of some monoterpenes (C10), diterpenoids, sesquiterpenes (C15), triterpenoids and their derivatives was recently reviewed. The results obtained illustrate the strong structure-

function influence of the antibacterial potential of terpenes. Diterpenoids, such as

sesquiterpenes, isolated from different plants exhibited bactericidal activity against Gram-

positive bacteria and inhibited the growth of M. tuberculosis [39]. The mechanism of action

of terpenoids is not fully understood, but is speculated to involve membrane disruption by the

lipophilic compounds [40].

6.1.1.4. Phenolics & polyphenols

Phenolic compounds are widely distributed in plants, where they protect the plants

from microbial infections. They have potential anti-oxidative properties but are also potent

anti-infectives. They are a large group of aromatic compounds, consisting of flavones,

flavanoids and flavanols containing one carbonyl group, quinones with two carbonyl groups,

tannins, polymeric phenolic substances, and coumarins, phenolic compounds with fused

benzene and pyrone groups [41].

6.1.1.5. Flavones and their derivatives

Flavones and their derivatives represent an antibacterial therapeutic possibility to

disrupt bacterial envelopes. The catechins are included among the flavan-3-ols or flavanols,

present in different plants, particularly in tea-plant Camelia sinensis, where they form

complexes with the bacterial cell wall and are active on intestinal microorganisms [42].

Biological assays indicated the inactivation of specific bacterial enzymes by several of these

compounds. Moreover significant synergy was also observed between theaflavin and

epicatechin against important nosocomial Gram-negative pathogens [43].

6.1.1.6. Quinones

Quinones (aromatic rings with two ketone substitutions), ubiquitous in nature, are

another significant group of secondary metabolites with potential antimicrobial properties.

They provide a source of stable free radicals and irreversibly complex with nucleophilic

amino acids in microbial proteins determining loss of their function. Anthraquinones in

particular had a large spectrum of antibacterial (also antimycobacterial) activity, based on

inactivation and loss of function of bacterial proteins, such as adhesins, cell wall polypeptides

and membrane-bound enzymes, consequently leading to the death of the pathogens [41].

6.1.1.7. Tannins

Tannins are a group of polymeric phenolic substances found in almost every plant part

characterized by antibacterial activity owing to inactivation of bacterial adhesins, enzymes,

cell envelope and transport proteins. Recently, gallotannin-rich plant extracts demonstrated

inhibitory activities on different bacteria attributable to their strong affinity for iron and to the

inactivation of membrane-bound proteins.

Hydrolysable and condensed tannins, derived from flavanols, and called

proanthocyanidins, exert antimicrobial activity by antiperoxidation properties inhibiting in

particular the growth of uropathogenic E. coli. Anthocyanidin synthesis occurs in plants on

the cytoplasmic leaflet of the endoplasmic reticulum and then accumulates in the large central

vacuole; in many plants, anthocyanidins might occur in oligomeric form and in this case they

are called proanthocyanidins. Depending on the type of bond between the oligomer-forming

anthocyanidin molecules, two general types (A and B) of proanthocyanidins are

distinguished. In less common A-type proanthocyanidins, two bonds are formed between

 2β -7 and 4β -8 carbon of oligomer-forming molecules; in B-type, only one 4β -8 bond is

formed. The beneficial effects of anthocyanins on human health have been known at least

from the 16th century, when blackberry juice was used in the treatment of mouth and eye

infections. However, only few studies have focused on the antimicrobial activity of these

compounds. Recently, Cisowska et al. described the anthocyanin profile of action of different

fruits, mainly berries, but also red grapes and, by consequence, red wine, also containing

stilbenoid resveratrol, indicating a superior activity against Gram-positive bacteria [44].

6.1.1.8. Coumarins

One known coumarin, scopoletin, and two chalcones were isolated as antitubercular

constituents of the whole plant Fatoua pilosa.

Also, spices and aromatic plants have an antimicrobial effectiveness that depends on

the kind of plant, its composition and concentration of essential oils, often rich in

monoterpens and sesquiterpenes. Studies analyzing the antimicrobial activity of essential oil

of Allium sphaerocephalon inflorescenses revealed the accordance with the popular use of

plants belonging to the Allium genus in traditional medicine, indicating the importance of

aroma precursors (cysteine sulfoxides) for a potent biologic activity [45].

6.1.2. Plant extracts with efflux pump inhibitory activity

Multidrug resistance due to the expression by bacteria of an efflux pump is an

increasing clinical problem. Therefore an interesting approach to the therapy of many

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infections would be one based on the identification of molecules interfering with the process of efflux.

In 1998, it was shown that plant-derived compounds are active against Gram-positive bacteria, in particular *Staphylococcus aureus*; successively numerous phytochemicals were shown to act as potential efflux pump inhibitors (EPIs) with antimicrobials for Gram-positive bacteria [46]. Gram-negative bacteria have innate multidrug resistance to many antimicrobial compounds owing to the presence of efflux pumps, in particular, the AcrAB-TolC efflux system, and some authors suggested that plants may not produce molecules active on these organisms [47].

However, the chemical diversity between plants and microorganisms represents an ecological possibility to identify EPIs from natural sources. Reviewing the literature concerning bacterial resistance modulators from natural plants, Stavri et al. described different bacterial EPIs, such as the plant alkaloid reserpine, berberine and methoxylated flavones and isoflavones, that revealed putative interfering activity on efflux [48]. Moreover, the level of accumulation of berberine in the cells was increased in presence of 5'-methoxyhydnocarpin, a multidrug pump inhibitor, reported as a minor component of chaulmoogra oil, used in traditional therapy for leprosy [49]. Recent data indicate that the AcrAB-TolC (in Enterobacteriaceae) and MexAB-OprM (in P. aeruginosa) efflux pumps are involved in the resistance of Gram-negative bacteria to most of the natural products [50]. In the presence of the EPI phenylalanine arginine β-naphthylamide (PAβN), the activities of some natural products belonging to the phenolics, in particular to the naphthoquinones (plumbagin), and flavonoids (4-hydroxyloncharpin), showed a significant increase in activity, whereas terpenoids are not active, probably due to difficulty in passing through the bacterial membrane barrier. The natural products exhibiting the best antibacterial activities have the same pharmacophore; plumbagin, which revealed significant antibacterial activity in the absence of an EPI, is the minimal scaffold required for activity. The other functional groups may modulate the susceptibility of the molecule to bacterial resistance mechanisms. Moreover, extracts from plants, and in particular an extract of an essential oil from a Corsican plant, Helicrysum italicum, containing geraniol, was able to synergize with chloramphenicol against different Gram-negative bacteria [51]. Garvey et al. indicated that extracts of different plants that are used as herbal medicinal products contain inhibitors of efflux in Gram-negative bacteria [52]. The most active compound, falcarindiol, extracted from Levisticum officinale, revealed a synergistic activity with ciprofloxacin. By adding EPI PAβN to *Dichrostachys* glomerata extracts, an increase of the activity on E. coli, Klebsiella pneumoniae and

Providencia stuartii resistant strains was shown; moreover, a synergistic effect was noted by associating *D. glomerata* extracts with some antibiotics.

Since the antimicrobial effectiveness of flavonoids comes from the ability to form complexes with both extracellular and soluble proteins and bacterial membranes, penetration of, and maintaining its position in, a microorganism is a critical point. Thus, the presence of EPIs is essential for flavonoid antimicrobial activity. Recently Fowler et al. used the natural flavonoid scaffold to synthesize non-natural flavanone molecules with functional groups responsible for activity against bacteria and fungi with minimal toxicity to human cells [53].

6.1.3. Plant extracts with bacterial quorum sensing inhibitory activity

It is now well recognized that populations of bacteria from many Gram-positive and Gram-negative species cooperate and communicate to perform diverse social behaviors, including swarming, toxin production and biofilm formation. Communication among bacterial cells involves the production and detection of diffusible signal molecules and has become commonly known as quorum sensing (QS), a density-dependent system that regulates the bacterial expression of specific genes, whose products modify the local host environment favoring the invasion and persistence of the pathogen [54]. The discovery that many pathogenic bacteria employ QS to regulate their virulence makes this system interesting as a target for antimicrobial therapy. Therefore, the ability to interfere with QS interrupting bacterial communication opens new therapeutic prospects. The ideal QS inhibitor (QSI) would be a low-molecular-mass molecule able to reduce the expression of QS-controlled genes; in order to avoid toxic side effects; the inhibitor should exhibit a high degree of specificity for the target QS-related molecule. Finally, the QSI agent should be chemically stable and resistant to the metabolic and disposal processes of the host organism. The study of a strategy to interfere with bacterial QS is the classical pharmacological approach to receptor antagonism. In particular, halogenated furanones, a class of natural products isolated from the marine red algae *Delisea pulchra*, have an effect on bacterial QS. Zang et al. showed that the mechanism of action is the modification and inactivation of LuxS (S-ribosylhomocysteine lyase), the enzyme which produces autoinducer-2, that mediates interspecies QS among many bacteria, but is absent in humans [55]. Moreover, a number of plant extracts and natural compounds inhibiting P. aeruginosa QS have been identified by Rasmussen et al., including bean sprout, chamomile, carrot, garlic, habanero (Capsicum chinensis), propolis, water lily, yellow pepper, and two products of *Penicillium* fungi, patulin and penicillic acid [56]. The

authors further investigated the effects of garlic extract, which contains at least three different QS inhibitors and was able to inhibit QS in a concentration-dependent manner and with a structure–activity relationship hypothesizing competitive binding. In fact, GeneChip® analysis revealed that garlic extract had a profound effect on QS-regulated virulence genes, significantly reduced *P. aeruginosa* biofilm tolerance to tobramycin and lowered the pathogenicity of *P. aeruginosa* in a *Caenorhabditis elegans* nematode model. The phytoalexin resveratrol (3,5,4'-trihydroxystilbene), an antifungal agent found in grapes and other plants, has direct antibacterial activity against *Neisseria gonorrhoeae* and *Neisseria meningitides*. However, we observed that resveratrol can inhibit *P. aeruginosa* QS *in vitro*. Also, solenopsin A, a venom alkaloid from the fire ant *Solenopsis invicta*, has been shown to

P. aeruginosa QS, probably by targeting the C₄-HSL-dependent *rhl* system [57]. Solenopsin A reduced biofilm production in *P. aeruginosa* in a dose-dependent manner, indicating a QS signaling suppression mechanism. An inhibition of QS-controlled virulence factors, such as *LasA* protease, *LasB* elastase, pyoverdin and biofilm production, in the same microorganism by extracts from different south Florida (USA) plants was also reported. Recently, some traditional Chinese medicine herbs, in particular *Areca catechu*, are a rich source of compounds which exhibit anti-QS properties. Several QSI of natural origin, in particular the isothiocyanate iberin from horseradish, and ajoene, a sulfur rich molecule from garlic that inhibits *P. aeruginosa* genes controlled by QS, were identified [58]. Both ajoene and horseradish juice extract, in combination with tobramycin, have a synergistic antibacterial efficacy. A natural nonpeptide compound isolated from the bark of *Hamamelis virginiana*, hamamelitannin (2',5-di-*O*-galloyl-D-hamamelose), was found to inhibit QS in *S. aureus* and *S. epidermidis*, inhibiting the production of RNAIII and δ hemolysin *in vitro* [59].

6.1.4. Plant extracts with biofilm inhibitory activity

Biofilms are the default mode-of-life for many bacterial species and biofilm-based infections cause harm to millions of humans annually. The difficulty of eradicating biofilm bacteria with classic systemic antibiotic treatments is a prime concern of medicine. In particular, the ability of staphylococci to adhere on both eukaryotic cells and abiotic surfaces and to form biofilm are important virulence factors in chronic infections associated with implanted biomaterials, which are particularly difficult to eradicate. Recently, Artini et al., assessing four compounds (derived from aerial and root parts of *Krameria lappacea*, *Aesculus*

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be able to interfere with

hippocastanum, Chelidonium majus and Macleya cordata) that contained several alkaloids and flavonoids, revealed a potentially interesting activity on staphylococci, clinically significant microorganisms also for the emergence of methicillin-resistant variants [60]. Two compounds in particular, proAc (proanthocyanidin A2-phosphatidylcholine) isolated from A. hippocastanum and CH (chelerythrine) purified from Macleya cordata, exhibited an inhibition of 'de novo' biofilm formation without bactericidal activity. The treatment of bacteria with these alkaloids downregulates some important proteins belonging to different pathways. In particular, proAc acts on the iron-binding protein (determining the impairment of the uptake of iron, an essential micronutrient for microorganisms), blocking the switch process from planktonic to sessile state of bacteria and ablating autolysin (penicillin-binding protein), thus inhibiting biofilm formation. The treatment with sanguinarine and CH acts on some bacterial proteins involved in heat shock response, surface exposed lipids and methoxy– mycolic acid synthase, until protein synthesis disappearance. Both sanguinarine and CH also act on some elements of the bacterial cytoskeleton, structural compart recognized as a potential target for antimicrobial therapy; therefore, inhibitors of cytoskeletal proteins may function as lead compounds for the development of novel antimicrobials. Hamamelitannin, a polyphenol extracted from the bark of Hamamelis virginiana belonging to the family of tannins, significantly reduces biofilm metabolic activity of different microorganisms [61].

Carvacrol, a monoterpenic phenol natural biocide, had an effect on dual-species biofilms formed by S. aureus and Salmonella enterica serovar typhimurium [62]. Nonbiocidal concentrations of this molecule disrupted normal development of biofilm, preventing the build up of protein mass and arresting at the microcolony stage. This component, together with thymol, is the principal phenolic component that determined the antimicrobial activity of oregano oil on staphylococci. These molecules, characterized by a hydrophobic nature, interact with the lipid bilayer of cytoplasmic membranes causing considerable effects on its structural and functional properties and loss of integrity of bacterial cell. Moreover, these compounds may diffuse through the polysaccharide matrix of the biofilm thus destabilizing it. A compound (1-deoxynoijirimycin) purified from Morus alba inhibited biofilm formation of S. mutans, a major causal organism of dental caries, reducing bacterial extracellular polysaccharide secretion [63]. Similarly a new naphthalene compound from Trachyspermum ammi seeds exhibited the same effect indicating great potential as a therapeutic agent against caries [64]. Moreover, another novel strategy to reduce development of dental caries may be the use of plant lectins, proteins that recognize the glycoconjugates present on the surface of S. mutans; in particular glucose/mannose-specific lectin altered the adhesion of bacteria on

saliva-coated surfaces [65]. Also, *Propionibacterium acnes*, microorganism responsible for acne vulgaris and able to form biofilm, resulted susceptible to plant extracts containing icariin, resveratrol and salidroside, compounds able to reduce biofilm formation [66].

For the treatment of urinary tract infections, *Melia dubia*, a plant from Meliaceae family present in the Indian subcontinent, has been used in folk medicine. Ravichandiran et al., examining the antivirulent potentiality of this plant, evaluated the principles antagonizing the quorum sensing systems of uropathogenic *E. coli* [67]. They found few compounds which can curtail the bacterial biofilm formation and virulence factor by controlling their quorum sensing

6.2. Antifungal agents

Human fungal infections, particularly in immunocompromised persons (AIDS, cancer and transplant patients), are a very challenging problem because the therapeutic options are hampered by serious drawbacks, such as the development of drug resistance and toxic side effects [68]. Thus, there is a clear demand for new therapeutic approaches based on molecules found in plants that may be used directly or considered as a model for developing better molecules. Before 2009, more than 600 plants have been reported for their antifungal properties, but few were examined for the active molecules [69]. Recently the use of the natural product tea tree oil in antifungal therapy has been proposed. This compound appears to be effective in vitro against multidrug-resistant Candida and in vivo against mucosal candidiasis [70]. Moreover, it has also been documented that terpinen-4-ol rather than 1,8-cineole is the most likely mediator of tea tree oil activity or, at least, a main contributor to anti-Candida activity. The genus Paeonia is one of the most important sources of drugs in traditional Chinese medicine. Picerno et al. observed that its extracts and some of their compounds inhibited C. albicans growth [71]. The antifungal properties of essential oils obtained from different aromatic plants, in particular from Mentha suaveolens, whose main microbicidal components were pulegone and piperitone oxide [72]. A strong antifungal activity of essential oils obtained from other plants was demonstrated [73]. In particular, in Bidens tripartite L. roots, the main components are α -pinene, β -bisabolene, p-cymene, hexanal and linalool; in Coriandrum sativum extracts, the effect is fungicidal and responsible for a marked reduction of germ tube formation [74]. From several parts (flower, leaf and stem containing different compounds) of Aloysia triphylla, Gypsophila bicolor, Lavandula viridis, Erigeron acris and annuus, and also from star anise (Illicium verum) an activity, linked to

trans-anethole, was observed [75]. Coumarin and phytoalexins, which are hydroxylated

derivatives of coumarins, revealed a certain antifungal activity. The antifungal activity of dill

(Anethum graveolens) oil results from its ability to disrupt the permeability barrier of the

plasma membrane and from mitochondrial dysfunction-induced reactive oxygen species

accumulation in Aspergillus flavus [76].

Promising activity against C. albicans biofilm formation was displayed by eugenol

and cinnamaldehyde, molecules belonging to the phenolic group of essential oil compounds,

which also showed synergy with fluconazole in vitro [77].

The essential oils of different Curcuma spp., containing caryophyllene as major

compound, displayed varying degrees of antimicrobial activity, in particular against

Cryptococcus neoformans. A protective effect of an oral natural phytonutrient was observed

in recurrent vulvovaginal candidiasis, and promising alternatives were revealed by several

terpenic derivatives for the topic treatment of oral candidiasis and denture stomatitis [78].

Some antidermatophitic compounds that have been long used as Chinese medicines to treat

various ailments such as dermatomycosis, were obtained from extracts of Fructus psoraleae

and Folium eucalypti globuli and also from Achillea millefolium extracts. Moreover,

flavonoids isolated from mango (Mangifera indica) leaves revealed antifungal activity on

different species, in particular Aspergillus sp., and schinol and a new biphenyl compound

were active on Paracoccidioides brasiliensis. Moreover, metronidazole showed a potentiation

of its antifungal effect when combined with plant extracts, as did fluconazole with other

phytocomponents. Using genetic and biochemical approaches, Xu et al. showed the antifungal

activity of a plant-derived acetylenic acid by interfering with the fatty acid homeostasis

pathway [79].

6.3. Antiviral agents

6.3.1. Oregon grape (Mahonia aquifollium/nervosa)

This antiviral herb's active ingredient is an alkaloid called berberine, a bright yellow

substance that's most prevalent in its roots. The roots are best harvested during the fall and

winter months when the plant is in a state of dormancy. At this time, all of the plant's energy

goes into the roots making the potency of the medicine at its highest. It's also starting to be

used as an alternative to goldenseal (*Hydrastis canadensis*), which is one of the antiviral herbs

that has been over harvested and has similar medicinal properties. Once you've harvested the

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roots, remove the brown outer bark and keep the yellow inner bark and wood. Scrape the

yellow inner bark and wood into shavings and make a tincture with the shavings using grain

alcohol. Oregon grape tincture can be used to address the following viruses: cytomegalovirus

(CMV), human papillomavirus (HPV), influenza, common cold (rhinovirus) [80].

6.3.2. St. John's Wort (Hypericum perforatum)

Hypericin is the active ingredient in St. John's Wort which is most prevalent in its

flowers. The flowers bloom during mid to late summer and that's when it is the best time to

harvest. People that infuse it in olive oil as well. While it has a multitude of medicinal

properties, it's an incredibly strong antiviral which shouldn't be underestimated. St. John's

Wort can b used to address the following viruses: HSV 1 and 2, HIV, Hepatitis C, MCMV,

Sindbus virus [81].

6.3.3. Ginger (Zingiber officinale)

Ginger root, like garlic, is best taken raw and it's typically my second "go to" herb

when either a cold or flu sneaks up on me. While it's often used for upset stomachs and

motion sickness, its 12 antiviral compounds make it incredibly effective against viruses as

well. Oftentimes I will shave portions of the root off onto my vegetables or just eat the

shavings raw. The root is best harvested during the fall and winter months when it's most

potent. Ginger can be used to address the following viruses: HRSV, common cold, flu and

most viral infections of the lungs [82].

6.3.4. Astragalus root (Astragalus membranaceus)

This is of the most popular antiviral herb in Chinese medicine because of it's ability to

strengthen the immune system. It's a popular treatment for people who have undergone

chemotherapy because of its ability to help the body quickly recover from such a severe

amount of stress. Roots are best harvested in the fall and winter months and you should make

it into a tincture similar to the method used for Oregon grape. Astragulus roots can be used to

address the following viruses: HIV, influenza, common cold and most viruses in general

because of how strong it makes your immune system [83].

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6.3.5. *Oregano* (*Origanum vulgare*)

Oregano is a member of the mint family which is high in volatile oils. The active

ingredient in these volatile oils is called carvatrol and its best extracted using an olive oil

infusion (let the oregano infuse for 6 weeks submerged in the olive oil). Oregano has also

been vastly underrated as a medicine since store-bought oregano oil is typically less than half

as potent as the oil you make yourself with wildcrafted oregano. The wildcrafted variety is

considered by many to be the best treatment for both strep throat and the flu. While the oil has

many benefits, it also kills beneficial bacteria, especially in your digestive system. Either

don't use for a prolonged period of time or supplement dosages with foods that contain

probiotics such as yogurt or sauerkraut. It's best to harvest when it's about 4 inches high

before it goes to seed. Oregano can be used to address the following viruses: HSV 1, shingles,

influenza, strep throat [84].

6.3.6. Usnea (Usnea australis)

While it's actually a lichen and not an herbaceous plant, I decided to include it on my

list of antiviral herbs because it's one of my cure-all favorites. Also known as Old Man's

Beard, it can often be found on fallen branches of deciduous trees here in the Pacific

Northwest. I often gather my usnea after wind storms. Its active ingredient is usnic acid and

its best consumed in tincture form. People with autoimmune disorders should avoid usnea

since it will encourage the immune system to attack healthy cells. Usnea can be used to

address the following viruses: influenza, common cold, HSV 1 and 2, Epstein-Barr, Junin

virus, Tacaribe virus and polyomavirus [85].

6.3.7. Lemon balm (Melissa officinalis)

Lemon balm is another member of the mint family whose volatile oils contain antiviral

compounds. It's best to harvest it between summer and mid fall since that's when the oils are

most abundant. You can make it into a tincture as an antiviral herb but I personally prefer it as

a tea or an infusion. It loses its volatile oils quickly so it's best to steep the leaves fresh rather

than dry them out. It's also a great herb for calming the central nervous system. Lemon balm

can be used to address the following viruses: HSV 1 and 2, influenza, common cold [86].

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6.3.8. Lomatium (Lomatium dissectum)

Lomatium is also known as desert parsley and is largely an underrated antiviral herb.

The root was used by Native Americans to treat a host of different viruses but was mainly

used for upper respiratory infections. It actually started to be written about in great detail

around 1918 after the outbreak of the Spanish flu where it saved a great deal of lives. Today

it's used for almost all types of bad respiratory infections, bad fevers and pneumonia. It's

active ingredient is an oleoresin stored in it's roots. It's a big and powerful medicine that is

best taken in tincture form. Take in small dosages since some people develop rashes when

taken in large dosages. Harvest the root either in the fall/winter when the plant is dormant or

in the early spring before the plant goes to flower. The root should be harvested when the

plant is between 4-10 years of age because that's when they have their highest medicinal

concentrations. It's considered to be one of the better influenza herbs because not only is it

antiviral, it's also respiratory clearing, meaning it prevents secondary infections which are

common occurrences with the flu virus. Lomatium can be used to address the following

viruses: cytomegalovirus, HIV, Epstein-Barr, influenza, common cold and practically any

respiratory virus [87].

6.3.9. White sage (Salvia apiana)

White sage is yet another mint family member. Its active ingredients are the volatile

oils camphor and eucalyptol. The best time to harvest sage is in the mid to late summer once

the flowers have gone to seed. It is best used in tincture form as an antiviral herb since many

of the active ingredients aren't water soluble. White sage can be used to address the following

viruses: common cold, influenza [88].

6.4. Antiprotozoal agents

Several natural compounds have been identified for the treatment of leishmaniasis and

research on plants and their metabolites can contribute to overcoming the drug resistance of

Leishmania parasites. Among the plant species evaluated here, N. falcifolia presented the best

results regarding anti-leishmanial activity, with the ethanolic leaf extract displaying an LD₅₀

of 138.5 μ g/ml and 65.6 \pm 5.4% growth inhibition of the promastigate forms of L. (V.)

braziliensis at the highest concentration tested, 320 µg/ml. Extracts of H. gardneriana (aerial

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parts) and *C. podantha* (leaves), which also demonstrated reasonable potency, presented an LD_{50} of 237 and 271 μ g/ml, respectively. No growth inhibition was obtained at drug concentrations lower that 40 μ g/ml. The medium containing DMSO did not affect the growth of the protozoa [89].

Ethanolic extracts of *C. podantha* and *M. arenosa* (aerial parts) inhibited the growth of epimastigote forms of *T. cruzi* even at very low concentrations (10 μ g/ml), presenting 90.4 \pm 11.52 and 88.9 \pm 2.20% growth inhibition of this protozoan, respectively. On the other hand, extracts of *H. gardneriana*, *N. falcifolia*, and *P. elegans* (leaves) showed similar activities only when a concentration of 1000 μ g/ml was used. The medium containing 1.0% DMSO did not affect the growth of the protozoa. Benznidazole used as the positive control against *T. cruzi* at 10 μ g/ml, showed 80% growth inhibitions [90].

The best results in terms of molluscicidal activity were obtained with the ethanolic extract of M. arenosa, which induced 100 and 60% snail mortality at concentrations of 200 and 150 μ g/ml, respectively, with an LD₅₀ of 143 μ g/ml. The N. falcifolia extract was 100% lethal to the snails at the concentration of 200 μ g/ml, but mortality was not obtained at lower concentrations. Control assays with DMSO showed no effect on the snails. Niclosamide at 5 μ g/ml was used as positive control against B. glabrata and showed 100% lethality [90].

Although the literature indicates that ideal concentrations of plant extracts are below $100 \,\mu\text{g/ml}$ for molluscicidal activity, the results obtained for *M. arenosa*, LD₅₀ of 143 $\mu\text{g/ml}$, justify the continuation of its study. This plant is native to the area and the extract was obtained from regenerating parts of the plant, factors that can be considered of importance [91].

The genus *Nectandra* is well represented in the Brazilian flora, with several species presenting many benefits to man. They have been used in popular medicine for the relief of pain, arthritis, rheumatism and diarrhea, and also as antifungals. Pharmacological studies have demonstrated the antitumoral activity of *N. rigida* Nees, the antimalarial activity of *N. cuspidata* Nees and the vascular and antimalarial activities of *N. salicifolia* Nees. In our study, *N. falcifolia* leaves presented good results regarding their antiprotozoal activity against promastigote forms of *L. (V.) braziliensis* [91].

Some species of the genus *Helicteres* have been used in folk medicine, such as *H. isora* L. (as an expectorant, demulcent, astringent, antigalactagogue, and for the relief of the flu, against empyema, stomach affections, and diabetes), *H. angustifolia* (analgesic, anti-inflammatory and anti-bacterial effects), *H. ovata* Lam. (depurative, emollient and antisyphilitic effects), and *H. sacarolha* Juss. (depurative and in syphilitic inflammations).

Pharmacological studies have demonstrated the antidiabetic and hypolipidemic activities of *H. isora* L. Also *H. gardneriana* (aerial parts) also displayed good antiprotozoal activity against promastigote forms of *L.* (*V.*) *braziliensis* [92].

Among the species of the genus *Cayaponia* that have been used popularly, *C. tayuya* (Vell.) Cogn. and *C. espelina* Cogn. (anti-snake venom, tonic, diuretic, anti-asthmatic, antisyphilitic, and purgative effects, and to combat epilepsy, diarrhea and bronchitis), *C. cabocla* M. (purgative and depurative effects in cutaneous diseases and as an emmenagogue) and *C. pilosa* Cogn. (emmenagogue, antisyphilitic and purgative effects). Our data demonstrated that *C. podantha* (leaves) presents important antiprotozoal activity against epimastigote forms of *T. cruzi* and promastigote forms of *L. (V.) braziliensis* [93].

Some species of the genus *Melochia* have been used in folk medicine, such as *M. corchorifolia* L. (dysentery, abdominal swellings and water-snake bites), *M. umbellata* (Houtt.) Stapf (deobstruent) and *M. pyramidata* L. (bronchitis and cough). The extract obtained from the aerial parts of *M. arenosa* demonstrated molluscicidal effects and activity against *T. cruzi* epimastigotes [94].

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Antimicrobial activity of certain secondary metabolites derived from family Scrophulariaceae

4

Fadia S. Youssef, Mohamed L. Ashour

Pharmacognosy Department, Faculty of Pharmacy, Ain Shams University, Cairo, Egypt

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ABSTRACT

The figwort family (Scrophulariaceae) is a big family comprising about 87 genera with nearly 4800 species. Members of this family are highly popular for their folk medicinal uses in addition to their phytotherapeutic importance. Scrophulariaceae represents an appealing, unique and diverse resource that furnishes a vast array of bioactive secondary metabolites including alkaloids, phenylpropanoids, iridoid glucosides, and terpenoids. Additionally, various species belonging to the Scrophulariaceae contain substantial amounts of flavolignans, polyphenols and phenolic acids. Undoubtedly, these phytoconstituents are responsible for their wide medicinal values as anti-inflammatory, antimicrobial, antinociceptive, antioxidant and cytotoxic properties. Buddleja, Eremophila, Leucophyllum, Myoporum, Scrophularia and Verbascum are among the highly valuable medicinal genera in the figwort family. In this chapter we are shading light on the antimicrobial potency of various species belonging to the aforementioned genera with special emphasis on the effects of their derived secondary metabolites and interpretation of their previously reported mode of action. Moreover, in silico molecular modeling study of the major constituents isolated from these genera of interest on various key enzymes responsible for the incidence, resistance and progression of infectious diseases will be carried out in an attempt to verify the probable

mechanism of action for the previously reported antimicrobial members belonging to

Scrophulariaceae.

Keywords: Antimicrobial activity; Buddleja; Eremophila; Leucophyllum; Verbascum;

Myoporum; Scrophularia; Molecular modeling; Phytoconstituents; Scrophulariaceae.

1. INTRODUCTION TO MICROBIAL INFECTIONS AND ITS

HAZARDOUS EFFECT

Infection can be defined as an invasive attack of the living organism by disease

producing agents as well as their multiplication within the organism's body with the

subsequent interaction of the immune system of the host organism towards them and their

toxins. Infectious diseases, also known as transmissible or communicable diseases, can be

classified anatomically into skin, odontogenic, respiratory tract, urinary tract infections as

well as vaginal and intra amniotic infections. The main symptoms of infectious diseases

comprise aches, fatigue and appetite loss with concomitant loss of weight as well as fever,

chills and night sweets. Additionally, other symptoms which are greatly specific to the

affected organ including cough, runny nose and skin rashes may appear in addition to the

existence of asymptomatic infections [1].

Despite of the enormous progress in the medicinal strategies for the curing of many

health problems, infectious diseases due to bacteria, fungi and viruses still constitute a major

impendence to public health in the 21st century. This is ultimately obvious in developing

countries attributing to the lack of medicine in addition to the appearance of many resistant

strains to commonly used antibiotics. Accordingly, novel classes of antibiotics are constantly

required to overcome the disturbing side effects of synthetic antimicrobial agents. Hence,

attention has been given to the beneficial therapeutic potential of herbal medicine setting an

example of cheap, substantially safe remedy offering a mine that could be used as

antimicrobial candidates [2].

The development of bacterial resistance combating the commonly used antibiotics has

been seriously reducing the cure rate. The probable mechanisms of bacterial resistance that

were previously reported include the inactivation of the antimicrobial agent directly through

the changing of the important functional groups in the drug as acetylation, methylation and

the opening of the beta-lactam ring of penicillin. Moreover, the modification of the targeted

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site of action, the variation in the metabolic pathways that the drug prohibited, inhibition of drug uptake by the affected tissues and reduction of the intracellular amount of the drug *via* its exportation out of the infected cell by ABC transporters are also among the prominent mechanisms of bacterial resistance [3].

2. PREVIOUSLY REPORTED MECHANISMS OF ACTIONS FOR POPULAR ANTIMICROBIALS

The usage of antimicrobial agent is widely accepted to prohibit the spread of infection. It can be classified according to their purpose into antiseptics that are effective in the removal of microorganisms from living tissue and/or skin. However, disinfectants are used for the destruction of the microorganisms existing on non-living things in addition to the antibiotics that are given as a prophylactic rather than as a cure. Besides, hand washing, wearing gowns together with face masks, the adherence to healthy lifestyle with a balanced diet and regular exercise may help to reduce the risk of incidence of bacterial infections. It is worthy to mention that the prolonged utilization of these agents causes bacterial mutations with consequent appearance of bacterial resistance [4].

There are a lot of mechanisms by which the antimicrobial agents exert their effects; this includes inhibition of protein, nucleic acids as well as cell wall synthesis, prohibition of the function of cell membrane, in addition to interfering with other metabolic processes (Fig. 1) [5]. The existence of cell wall in the bacterial cells that is crucial to their survival make them more susceptible for being attacked by the antimicrobial agents that targeted cell wall synthesis inhibition as penicillins and cephalosporins comparable to human and animal cells that lack cell wall. On the contrary, polymixin B and colistin that affect the cell membrane structure causing its damage and the leakage of vital substances necessary for the cell's survival is nonselective and also adversely affect human and animal cells. However, antimicrobial agents that inhibit protein are designed to selectively attack certain bacterial enzymes and structures that are required for their multiplication and growth with concomitant interference with their metabolism as 30S or 50S subunits of the intracellular ribosomes as aminoglycosides chloramphenicol and tetracyclines. Meanwhile, it goes without saying that nucleic acids, DNA and RNA, are of great necessity to the bacterial division and survival so selective binding with the bacterial nucleic acids could effectively lead to their death as quinolones. Additionally, many antibiotics hits certain metabolic paths that are essential for

the bacterial survival as inhibition of dihydrofolate reductase and dihydropteroate synthase enzymes resulting in disruption of folic acid synthesis with consequence destruction of DNA synthesis as sulfonamides and trimethoprim [6].

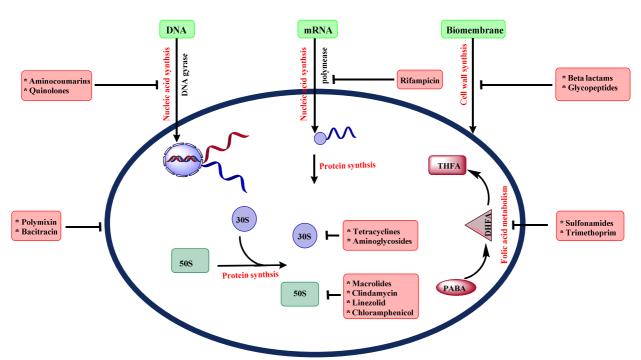


Figure 1. Major microbial targets for antimicrobial agents

Apart from the well-defined targets of the commonly used antimicrobial agents, antimicrobial enzymes that constitute a prominent part from the host immune system implicated in its defense against pathogenic microbes are nowadays adopted as an advanced method for combating infectious disease. These enzymes include hydrolyzing ones that prohibit bacterial growth either by invading the crucial components of the cell wall or by destroying the substances that adheres the cells together and to surrounding surfaces. Besides, proteolytic enzymes exemplified by subtilisins and lysostaphin are highly effective in hydrolyzing the adhesive proteins that are mandatory in the formation of bacterial biofilms. Polysaccharide-hydrolyzing enzymes comprising α -amylase, dispersin B, chitinases, β -glucanases and alginate lyases are found to be effective in both inhibiting biofilm formation as well as destroying the formed films in a vast array of microorganisms *via* cleavage of various glycoside linkages. Additionally, oxidative enzymes mainly cellobiose dehydrogenase, glucose oxidase and superoxide dismutase are known to elicit an anti-infective

manner via the liberation of a huge amount of the destructive hydrogen peroxide that causes

cytotoxicity in the infectious agent. Among the interesting host enzymes that exhibited potent

antifungal properties are quorum-quenching and urease enzymes. The formers include AHL-

lactonase, AHL-acylase and paraoxonases that showed a great interference to bacterial cell-

to-cell communication (quorum sensing) that undoubtedly decrease the ability of the

organisms to release virulence components causing eradication of the organisms. However the

latter is highly efficacious in case of urinary tract infections inhibiting the formation of hard

coatings around the bacteria that protects it [7-9].

3. REPORTED ANTIMICROBIAL ACTIVITY OF SOME IMPORTANT

GENERA BELONGING TO FAMILY SCROPHULARIACEAE AND

THEIR ISOLATED PHYTOCONSTITUENTS

3.1. Buddleja

Genus Buddleia (Buddleja) belonging to the family Scrophulariaceae comprises about

100 species that are native to Africa, Asia, North and South America [10]. Traditionally,

Buddleia species were highly recommended to be used as a topical antiseptic as well as a

diuretic owing to the different classes of compounds predominating in the genus [11].

Recently many of the Buddleia species have been investigated for their antimicrobial

properties and many of which are found to be highly effective either in the form of crude

extract or isolated compounds.

The stems and the leaves of B. saligna were evaluated for their antimicrobial activity

using isolates of 10 bacteria species comprising five Gram-positive and five Gram-negative

strains. The tested samples showed substantial activity against the Gram-positive and some

Gram-negative strains that further consolidates the fact that Gram-negative bacteria are more

resistant relative to the Gram positive ones. The methanol extract of its leaves that showed

higher antibacterial potency comparable to the stems was found to exert its antimicrobial

effect against all the tested bacterial strains except for Serratia marcescens and Pseudomonas

aeruginosa. Bacillus cereus, Streptococcus pyrogens and Pseudomonas aeruginosa only were

susceptible to the stem effect [12].

Additionally, the ethanol extracts of *B. globosa* leaves exhibited a potent antibacterial

activity against Staphylococcus aureus and Escherichia coli mainly attributing to the presence

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of verbascoside that showed a minimal inhibitory concentration of 1 mM [13]. Verbascoside (1) was also isolated from *B. cordata* leaves methanol extract and was monitored for its antibacterial property against *Staphylococcus aureus* utilizing killing kinetics together with incorporation of precursor methods. Results clarified the lethal effect of verbascoside on *S. aureus via* interfering with protein synthesis and inhibition of leucine incorporation [14]. Additionally, *B. globosa* stem bark showed antifungal activity against *Trichophyton rubrum*, *Trichophyton interdigitale*, and *Epidermophyton floccosum*. Buddlejone (2), maytenone (3), buddledin A (4) and buddledin B (5) and deoxybuddlejone (6) were found to be responsible for the antifungal activity with buddledins A and B showed the greatest activity, with MIC values of 43 µM and 51 µM, respectively [15].

Similarly, the antifungal activity of the chloroform extracts of *B. cordata* and *B. davidii* stem bark against the soil fungiwas explained owing to the predominance of the sesquiterpene buddledin A [16].

Moreover, the acetone/water (4:1) crude extracts of *B. saligna* leaves and stem showed pronounced activity by bioautography against *Escherichia coli*, *Staphylococcus aureus*, and *Mycobacterium aurum*. Oleanolic acid (7) isolated from the *n*-hexane soluble fraction of the crude extracts exhibited bactericidal activity against *Mycobacterium microti*, *Mycobacterium avium* and *Mycobacterium scrofulaceum* at loading doses of 2.5 μg/spot for the first and 1.25 μg/spot for the last two ones. Its mechanism of action could be explained in virtue of suppression of DNA polymerase [17]. For *B. brasiliensis* leaves methanol extract, it showed antibacterial properties against *Aeromonas hydrophila*, *Bacillus subtilis*, *Pseudomonas aeruginosa* and *Staphylococcus aureus* [18].

B. perfoliata Kunth, used traditionally for the alleviation of digestive disorder, exhibited *in vitro* anti-*H. pylori* activity with minimum inhibitory concentration (MIC) of aqueous and methanol extracts of its aerial parts equals to 500, and 62.5 μg/ml, respectively [19]. The crude extract of the stem bark of *B.cordata* exhibited a marked anti-mycobacterial activity owing to the presence of various phytoconstituents comprising 2[4'-hydroxyphenyl]-ethyl lignocerate (8) that revealed a substantial antibacterial activity against *Mycobacterium tuberculosis* with MIC value of 64 micrograms/ml [20].

The leaf extracts of *B. salviifolia* as well as its isolated compounds namely, 4'-hydroxyphenyl ethyl vanillate (9), verbascoside (1) and quercetin (10) showed a broad spectrum of antibacterial activity. Its ethyl acetate fraction revealed a good activity against *Bacillus subtilis* and *Staphylococcus aureus* wheraeas, the hexane and dichloromethane fractions showed the highest activity towards *Candida albicans* [21]. Besides, the volatile oil

of *B. asiatica* leaves that was found to be enriched with β -caryophyllene oxide (11), citronellol (12), and β -caryophyllene (13) revealed a highly potent antifungal, antibacterial and anthelmintic activity [22].

3.2. Eremophila

Members of the genus *Eremophila*, commonly known as Fuchsia bush or Emu bush, represent perennial shrubs containing approximately 214 species with an endemic existence in the arid and semi-arid areas in Australia. They have been employed in the folk medicine for the cure of many health disorders including respiratory, gastro-intestinal tract and skin infections. Additionally many biological activities have been assigned to the genus antiinfective, immunomodulatory, and anti-inflammatory as well as antiproliferative activities. This could be due to its richness with flavonoids, lignans, phenylpropanoids and terpenoid [23]. Evaluation of the antimicrobial potency of various *Eremophila* extracts as well as their isolated compounds revealed their antimicrobial efficacy particularly against Gram-positive bacteria [24]. E. duttonii extract showed a high antimicrobial potency against Gram-positive bacteria exemplified by Enterococcus faecalis, Bacillus cereus, Staphylococcus aureus and Streptococcus pyogenes [25]. Additionally, E. alternifolia and E. duttonii ethanol extracts of inhibited clinical isolates of methicillin-resistant Staphylococcus aureus (MRSA) and vancomycin-resistant Enterococcus (VRE) [26]. Moreover, various food-borne pathogensas Clostridium spp. namely, C. perfringens, C. sporogenes in addition to Listeria monocytogenes were greatly susceptible to the lethal effect of E. duttonii extract. This pronounced activity could be to a great extent relied upon the existence of various sterols and terpenes as revealed by bioautography [27]. Recently, extracts for both E. alternifolia and E. duttonii have shown inhibition zones of 8.8 mm and 9.6 mm, respectively for Listeria monocytogenes [28]. Serrulat-14-en-7,8,20-triol (14) and serrulat-14-en-3,7,8,20-tetraol (15) isolated from its nhexane fraction revealed potency against Staphylococcus aureus, Staphylococcus epidermidis and Streptococcus pneumonia [29].

Regarding the diethyl ether extract of *E. neglecta*, 8,19-dihydroxyserrulat-14-ene (**16**) and 8-hydroxyserrulat-14-en-19-oic acid (**17**) were isolated and showed antimicrobial potency against *Staphylococcus aureus*, *Streptococcus pyogenes*, and *Streptococcus pneumoniae* in addition to exertion of activity against the Gram-negative bacteria *Moraxella catarrhalis* showing MIC values of 3.1, 6.2 µg/ml for the two compounds, respectively [30, 31]. *E. serrulata* leaves extracts offered O-naphthoquinone (**18**), and 20-acetoxy-8-

hydroxyserrulat-14-en-19-oic acid (19) that showed activity towards *Streptococcus pyogenes*, and *Streptococcus pneumonia* [32]. However the ethyl acetate fraction obtained from *E. sturtii* leaves together with its isolated compounds, 3,8-dihydroxyserrulatic acid (20) and serrulatic acid (21) showed potency with the latter being most potent against *Staphylococcus aureus* [33].

A study carried out by Ndi et al., in 2007 evaluating the antibacterial activity of 72 different extracts of *Eremophila* towards 68 clinical isolates of multi-resistant methicillin-resistant *Staphylococcus aureus* (MRSA). Results shaded the light on the antimicrobial property of several tested *Eremophila* species particularly *E. virens* that prohibited the growth of all the examined isolates at a concentration of 31µg/ml [34]. *E. longifolia* ethanol extracts of the stem lethally affect *Streptococcus mutans* and *Streptococcus sobrinus* growth owing to the prevalence of phenolics [35]. *E. maculata* leaves ethanol extract was effectively potent against three Gram-positive bacteria [36]. Additionally, *E. longifolia* offered antimicrobial agents which are neryl ferulate (22) and neryl *p*-coumarate (23). The former showed moderate effect against several various Gram-positive bacterial strains whereas the latter was effective only towards *Enterococcus faecium* only [37].

The antimicrobial efficacy of *E. microtheca* and its isolated compound jaceosidin (24) recorded relevant antibacterial activity against *Staphylococcus aureus* strains [38]. In a mechanistic study to trace the antimicrobial potency of 8-hydroxyserrulat-14-en-19-oic acid (17) isolated from *E. neglecta*, results revealed that its bactericidal effect can be interpreted by its effect on the logarithmic-phase, stationary-phase, and adherent *Staphylococcus epidermidis*, as well as against methicillin-susceptible and methicillin-resistant S. aureus making it unable to produce polysaccharide intercellular adhesion-mediated biofilm. Thus, this clarified its multi-target effect through its hydrolytic properties on the cell membrane together with the general inhibition of macromolecular biosynthesis [39]. The smoke extract obtained from *E. longifolia* exhibited potent antimicrobial properties against the Grampositive species *Staphylococcus aureus*, *Bacillus subtilis* and the yeast *Candida albicans* owing to the presence of genifuranal that is formed as a result of rearrangement of geniposidic acid upon heating that does not exist naturally in the leaves [40].

Besides, the antimicrobial properties for a number of *Eremophila* species have been reported comprising *E. bignoniiflora* and *E. maculata*. The former with its major compounds fenchyl-acetate and bornyl-acetate successfully inhibited pathogenic *Trichophyton* species associated with dermatophytosis, however substantial activity was proven against *Candida albicans* and *Staphylococcus epidermidis* [41]. Meanwhile the volatile oil of the of the leaves

and flowers latter species showed a relevant antimicrobial activity against a panel of Gram

positive, Gram negative, MRSA and fungi [42].

3.3. Leucophyllum

Leucophyllum is a small genus comprising about 15 species of evergreen shrubs,

native to Mexico as well as the southwestern United States [43]. The methanol extracts

obtained from the roots and leaves of L. frutescens revealed a high antimicrobial potency

against the drug-resistant strain of Mycobacterium tuberculosis; with MICs of 62.5 and 125

µg/ml, respectively [44]. Besides, the diterpene leubethanol (25) isolated from L. frutescens

showed notable antibacterial potency towards multidrug-resistant strains as Staphylococcus

aureus, and is of great interest for being applied in the strict control of bacterial biofilms

formation [45]. Additionally, the furolignan 2',5"-dimethoxysesamin (26), isolated from the

its root bark, exhibited promising antituberculous activity with a MIC equals to 63 µg/ml with

no observed cytotoxicity [46].

3.4. Myoporum

Myoporum is a genus including about 30 species of flowering plants and recently

included in the family Scrophulariaceae. It is native to the Australian areas as well as Pacific

islands and Indian oceans. Myoporum members are mainly shrubs or small trees mostly with

white flowers. Many Myoporum species have shown a potent antimicrobial activity including

M. acuminatum in which the essential oil of its fruits revealed a notable antimicrobial potency

against a panel of bacteria as Bacillis subtilis, Streptococcus pneumonia, Escherichia coli as

well as fungi comprising Aspergillus fumigates, Geotricum candidum and Syncephalastrum

racemosum. This may be attributed to the synergistic action of all its components that are

represented mainly by D-limonene (27) and (-) carvone (28) as monoterpenoids in addition to

negaione (29) and myomontanone (30) as furanoid sesquiterpenes [47].

Moreover, the leaves of M. montanum was found to contain three toxic

furanosesquiterpenes namely (±)-myoporone (31), (-)-10,11-dehydromyoporone and

11-hydroxymyoporone that showed a prominent antibacterial activity towards *Enterococcus*

faecalis, Moraxella catarrhalis and Staphylococcus epidermidis with immeasurable

cytotoxicity against a number of cancer cell lines as well as the normal breast cells [48].

Also, the essential oil of its leaves and stems revealed a potent antimicrobial activity [49].

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Meanwhile, the ethereal oil of M. crassifolium showed a considerable antimycobacterial

activity against Mycobacterium bovis [50]. Regarding M. bontioides, it showed a relevant

antibacterial activity particularly towards Staphyloccocus aureus and Escherichia coli with its

constituent, 5,7-dihydroxyflavone (32) showing substantial of antibacterial property with

MIC of 62.50 µg/ml [51]. Besides, their isolated compounds namely tangeretin (33),

sinensetin (34), dihydrokaempferol (35), luteolin (36) exhibited significant antifungal activity

towards Colletotrichum musae [52].

Additionally, in a recent study that was done using Chromobacterium violaceum assay

to evaluate the ability of the substances to hinder and disrupt the bacterial communication

system expressed in terms of quorum sensing (QS). Leaves of M. laetum revealed an

interesting anti-quorum sensing that undoubtedly reflected in its potent antimicrobial activity

via prohibiting the secretion of virulence factors facilitating the eradication of the organisms

[53]. Additionally, a significant antiviral, antibacterial as well as antifungal activity was

shown by its essential oil that showed its richness by as ngaione (29), myoporone (31), and

myomontanon (**30**) [47].

3.5. Scrophularia

Scrophularia with its 200 species represents a large genus of herbaceous plants that

are widely distributed in Asia. Its name was derived from a form of tuberculosis termed

scrofula as the majority of its species are highly popular as antituberculous agents.

Traditionally, it was employed for the relief of many ailments owing to its richness with

various secondary metabolites particularly iridoids [54, 55].

Additionally many species of Scrophularia have been used for the alleviation of

infectious diseases as S. buergeriana that showed potent antimicrobial and anti-viral

properties [56]. Moreover, the leaves extract of S. ningpoensis together with its isolated

saponin glycoside, scrokoelziside A (37), showed a great activity against beta-haemolytic

streptococci using disc-diffusion as well as the micro-well dilution methods [57]. Recent

antimicrobial investigations done on the highly reputable traditional antimicrobial herbaceous

herb S. striata applying micro broth dilution assay towards a wide array of bacteria and fungi

resulted in a potent antimicrobial properties. Actually, this activity was found to be directly

related to the total phenolic content as reveled from the total phenolics and flavonoids

assessments [58]. Moreover, S. deserti showed a relevant antimicrobial activity that is

probably relied on the presence of 3(beta)-hydroxy-octadeca-4(E),6(Z)-dienoic acid,

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ajugoside (38) and scropolioside B that exhibited substantial antimicrobial activity towards

multidrug strains as well as methicillin-resistant Staphylococcus aureus (MRSA) and a panel

of rapidly growing mycobacteria showing MICs values between 32 and 128 µg/ml [59].

However, the essential oil isolated from the aerial parts of S. subaphylla that is composed

mainly of terpenoids and fatty acids namely, linalool, phytol, geraniol and palmitinic acid that

showed mild antibacterial activity comparable to other *Scrophularia* members [60].

3.6. Verbascum

The genus Verbascum is commonly named by velvet plant belongs to the figwort

family and comprises about 250 species. They mostly spread in Asia, Africa, and North

America and Europe. Many beneficial internal and external effects have been attributed to

leaves and flowers of many of its species. Traditionally, they were highly adopted to induce

diuresis, expectoration as well as sedation [61].

Many Verbascum species as V. bombyciferum, V. olympicum, V. thapsus and

V. xanthophoeniceum showed potent antimicrobial activity in many recent researches [62,

63]. Additionally, the ethanol/water (70:30 v/v) extract of *V. macrurum* leaves revealed potent

antibacterial activity [64]. It is worthy to mention that the antimicrobial activity of

V. densiflorum and V. phlomoides is greatly attributed to their richness in flavonoids and

phenylethanoids. Diosmin (39) and tamarixetin 7-rutinoside (40) are highly prevalent in the

flower V. phlomoides meanwhile verbascoside (1) and luteolin 7-glucoside (41) (were greatly

abundant in V. densiflorum flower [65].

V. bottae was proved to be a highly efficacious antibacterial agent exhibiting its effect

against a wide panel of Gram-positive bacteria including multi-resistant ones owing to the

existence of various classes of compounds like flavonoids and terpenoids [66]. Moreover,

the antimicrobial evaluation done on various extracts of V. pinetorum and V. antiochium

revealed that its methanol and methanol/chloroform extracts are highly potent on a broad

range of microorganisms particularly *Haemophilus influenzae*, whereas the acetone extract of

the former was highly active against and Candida albicans. Undoubtedly the observed

activity relied upon its phytoconstituents mainly iridoid glycosides, flavonoids, saponins and

phenolic compounds [67-69].

Besides, V. leptostychum flower showed a notable antimicrobial activity against

Proteus sp., Pseudomonas aeruginosa, Shigella dysenteria, Salmonella enteritidis,

Salmonella typhi, Staphylococcus aureus, Streptococcus faecalis, and Candida albicans [70].

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Hydroalcoholic extracts of *V. sinaiticum* showed sustainable antimicrobial activity towards *Staphylococcus aureus* and *Trichophyton mentagrophytes* that could provide evidence about its popularity in the folk medicine as a cure for various skin disorders [71].

4. IN SILICO MOLECULAR MODELING STUDY OF THE MAJOR CONSTITUENTS ISOLATED FROM THESE GENERA

In this section, molecular modelling studies of the isolated phytoconstituents from the 6 major Scrophulariaceae genera that showed relevant antimicrobial activity were done on important target enzymes implicated in the occurrence and dissemination of infection. This was done in an effort to explore the exact mechanism of action of these naturally occurring entities in fighting bacterial infections. Noteworthy to mention that plethora of enzymes could be targeted to either treat bacterial infections or prohibit the development of bacterial drug resistance. Herein, six enzymes which are crucial in the survival and division of bacteria as well as the development of resistance were chosen in the molecular modelling studies using C-docker protocol [72, 73]. The enzymes were downloaded from the protein data bank and are as follows: DNA-gyrase (PDB ID 4Z2D; 3.38 A°); topoisomerase IV (PDB ID 4Z3O; 3.44 A°); dihydrofolate reductase (PDB ID 4KM2; 1.4 A°); transcriptional regulator TcaR (protein) (PDB ID 4EJV; 2.9 A°), β -lactamase (PDB ID 3NBL; 2.0 A°) and aminoglycoside nucleotidyl transferase (PDB ID 4WQL; 1.73 A°).

DNA-gyrase is a vital enzyme that regulates the supercoiling of DNA in addition to relieving the topological stress resulting from the translocation of transcription and replication complexes within DNA. However, topoisomerase IV is an enzyme responsible for the decatenation and separation of interlinked daughter chromosomes consequently after DNA replication [74]. Folic acid is of great necessity to the bacterial growth as well as multiplication [75]. Dihydrofolate reductase is responsible for the catalysis of the NADPH-dependent reduction of dihydrofolate to tetrahydrofolate; consequent metabolites of tetrahydrofolate are required for incorporation of single carbon units into purines, pyrimidines and amino acids. Thus prohibition of dihydrofolate reductase resulted in a deficiency of the components of nucleic acids and proteins with concomitant inhibition of DNA synthesis and eventually cell death and can undoubtedly employed as antibacterial agents [76, 77].

Regarding bacterial resistance, β -lactamases constitute enzymes formed by the bacteria that are responsible for the development of multi-resistance to antibiotics containing

 β -lactam ring as penicillins *via* cleavage of the β -lactam ring and thus destroying the activity

of the antibiotics. Thus development of β -lactamases inhibitors is important for the prevention

of bacterial resistance [78]. Additionally, aminoglycoside nucleotidylyltransferase is an

enzyme involved in bacterial resistance to aminoglycoside antibiotics that changes the

structure of the antibiotics through adenylylation and thus deactivating the drug causing it to

be inactive towards bacteria [79].

Results of the molecular modelling of forty one (Fig. 2) previously reported

antimicrobial secondary metabolites from selected species belonging to Family

Scrophulariaceae in the active sites of six important enzymes implicated in the incidence of

bacterial infections as well as development of bacterial resistance using molecular modeling

experiments calculated in Kcal/mol were illustrated in Table 1 and Fig. 3 and 4. Results

revealed that tamarixetin 7-rutinoside (40) showed the highest binding in the active sites of

dihydrofolate reductase, β-lactamase and aminoglycoside nucleotidyl transferase as evidenced

from its biding free energies meanwhile diosmin (39) showed the highest inhibition to

topoisomerase IV. However, scrokoelziside A (37) exhibited the highest inhibition towards

DNA-gyrase and transcriptional regulator protein TcaR.

5. CONCLUSIONS

Family Scrophulariaceae with most of its genera offers a hidden mine for diverse

promising secondary metabolites that could be of high relevant antimicrobial activity.

Nevertheless, more thorough phytochemical and biological studies should be done on many

of its species to discover many antimicrobial leads that can overcome the resistance exerted

by many microbial strains.

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Figure 2. The structure of the secondary metabolites that showed antimicrobial activity from selected genera belonging to Schrophulariaceae

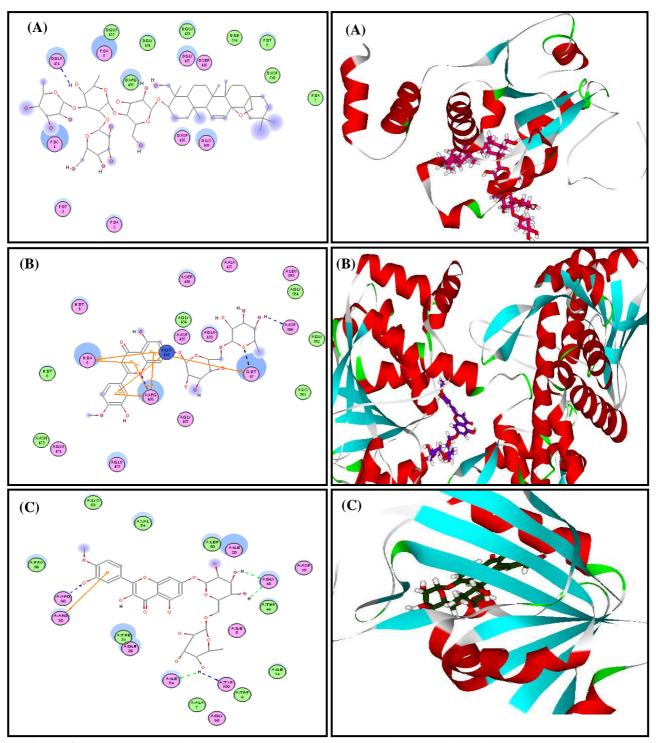


Figure 3. 2D and 3D binding modes of scrokoelziside A within the active sites of DNA-gyrase (A), 2D and 3D binding modes of diosmin within the active sites of topoisomerase IV (B) and 2D and 3D binding mode of tamarixetin 7-rutinoside within the active sites of dihydrofolate reductase (C)

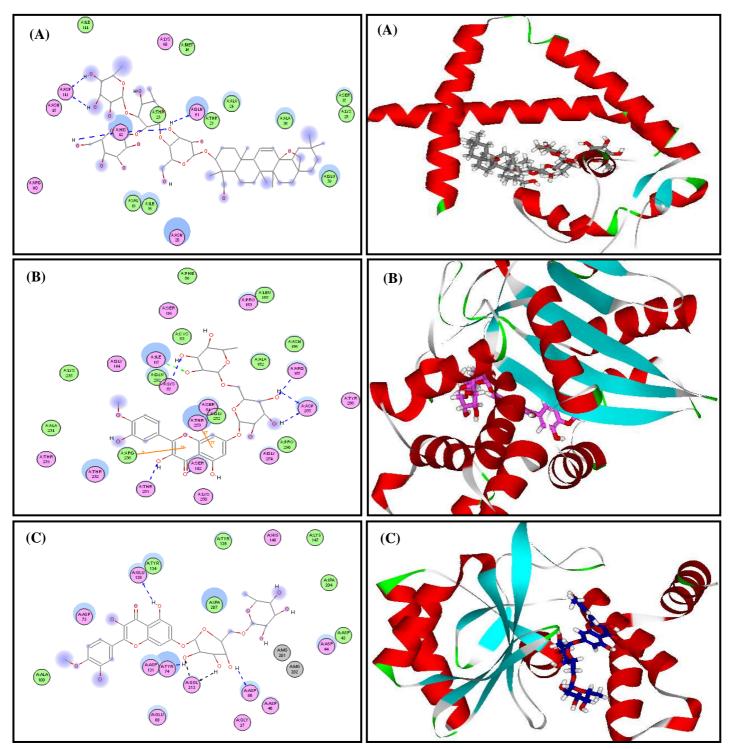


Figure 4. 2D and 3D binding mode of scrokoelziside A within the active sites of transcriptional regulator TcaRprotein (A), 2D and 3D binding mode of tamarixetin 7-rutinoside within the active site of β -lactamase (B) and 2D and 3D binding mode of tamarixetin 7-rutinoside within the active site of aminoglycoside nucleotidyl transferase (C).

Table 1. Free binding energies (ΔG) of some previously reported antimicrobial secondary metabolites from selected species belonging to family Scrophulariaceae in the active sites of six important enzymes implicated in the incidence of bacterial infections as well as development of bacterial resistance using molecular modeling experiments calculated in Kcal/mol

Compounds	Binding energy (Kcal/mol)							
	DNA- gyrase	Topoisomerase IV	Dihydrofolate reductase	Transcriptional regulator TcaR	β- lactamase	Aminoglycoside nucleotidyl transferase		
Verbascoside (1)	FD	FD	FD	-49.58	FD	-76.83		
Buddlejone (2)	-25.73	31.20	- 30.83	-22.74	FD	-51.02		
Maytenone (3)	-37.14	FD	FD	-33.88	FD	-33.75		
Buddledin A (4)	-23.99	-26.78	-29.53	-26.22	-33.26	-44.33		
Buddledin B (5)	-21.86	-24.57	-29.63	-21.25	-29.07	-32.70		
Deoxybuddlejone (6)	-28.84	-27.70	-30.16	-25.77	-32.76	-39.66		
Oleanolic acid (7)	-33.01	-33.28	-42.22	-31.24	-42.80	-43.21		
2[4'-Hydroxyphenyl]- ethyl lignocerate (8)	-49.64	FD	-48.91	-42.39	-57.09	-65.69		
4'-Hydroxyphenyl ethyl vanillate (9)	-32.84	-43.32	-39.88	-27.22	-45.32	-47.07		
Quercetin (10)	-31.17	-41.63	-40.28	-23.31	-48.82	-45.00		
Caryophyllene oxide (11)	-23.06	-23.21	-28.62	-22.22	-28.83	-27.58		
Citronellol (12)	-23.61	-24.34	-25.84	-24.64	-29.62	-32.05		
β-caryophyllene (13)	-21.32	-21.41	-22.80	-20.32	-28.84	-25.47		
Serrulat-14-en-7,8,20- triol (14)	-32.59	-39.46	-41.26	-28.76	-44.20	-46.57		
Serrulat-14-en- 3,7,8,20-tetraol (15)	-37.05	-40.00	-44.69	-28.01	-40.94	-58.80		
8,19- Dihydroxyserrulat- 14-ene (16)	-32.81	-36.61	-37.34	-28.66	-43.51	-40.32		
8-Hydroxyserrulat- 14-en-19-oic acid (17)	-33.63	-40.72	-40.25	-29.98	-48.19	-44.63		
O-naphthoquinone (18)	-29.32	-33.18	-34.71	-27.14	-36.66	-47.44		
20-Acetoxy-8- hydroxyserrulat-14- en-19-oic acid (19)	-40.95	-43.44	-45.86	-34.20	-49.20	-60.25		
3,8- Dihydroxyserrulatic acid (20)	-37.40	-35.95	-42.52	-33.54	-45.17	-51.87		
Serrulatic acid (21)	-36.26	-37.57	-41.33	-37.53	-44.66	-52.56		
Neryl ferulate (22)	-31.31	-42.40	-40.91	-29.80	-41.80	-45.98		
Neryl p-coumarate (23)	-35.63	-47.57	-37.78	-29.28	-48.74	-46.09		
Jaceosidin (24)	-33.18	-41.79	-40.13	-27.25	-41.39	-48.96		
Leubethanol (25)	-31.30	-36.26	-33.68	-28.17	-38.94	-41.86		
2',5"- Dimethoxysesamin	-32.42	FD	FD	FD	-46.28	FD		

Compounds	Binding energy (Kcal/mol)							
	DNA- gyrase	Topoisomerase IV	Dihydrofolate reductase	Transcriptional regulator TcaR	β- lactamase	Aminoglycoside nucleotidyl transferase		
(26)								
D-Limonene (27)	-18.69	-21.12	-16.59	-14.13	-19.74	FD		
(-) Carvone (28)	-20.43	-23.00	-21.81	-18.53	-26.21	FD		
Negaione (29)	-27.11	-32.36	-31.77	-27.55	-34.14	-42.06		
Myomontanone (30)	-23.90	-28.70	-27.92	-23.83	-31.57	-37.35		
(±)-Myoporone (31)	-27.99	-34.66	-33.07	-24.85	-42.89	-43.16		
5,7-Dihydroxyflavone (32)	-25.89	-33.10	-29.32	-22.81	-38.57	-41.31		
Tangeretin (33)	-34.19	-46.10	-42.48	-27.13	-45.40	-49.42		
Sinensetin (34)	-36.09	-48.18	-42.20	-32.07	-45.57	-49.46		
Dihydrokaempferol (35)	-28.83	-36.70	-34.59	-32.79	-41.00	-49.33		
Luteolin (36)	-32.86	-40.12	-39.00	-26.62	-38.15	-45.04		
Scrokoelziside A (37)	-62.28	FD	FD	<u>-56.01</u>	FD	FD		
Ajugoside (38)	-36.64	-34.36	-47.84	-35.96	-58.17	-56.70		
Diosmin (39)	-48.97	<u>-69.75</u>	-60.76	-46.97	-66.76	-66.76		
Tamarixetin 7- rutinoside (40)	-46.12	-64.60	<u>-61.32</u>	-44.79	<u>-67.11</u>	<u>-84.85</u>		
Luteolin 7-glucoside (41)	-43.77	-54.73	-52.81	-40.91	-53.29	-56.29		
Trimethoprim	ND	ND	-35.69	ND	ND	ND		
Chloramphenicol	ND	ND	ND	-29.02	ND	ND		
Cefuroxime	ND	ND	ND	ND	-70.87	ND		
Levofloxacin	-36.26	ND	ND	ND	ND	ND		
Moxifloxacin	ND	-53.10	ND	ND	ND	ND		
Kanamycin	ND	ND	ND	ND	ND	-73.94		

ND: not done FD: fail to dock

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