# **Chapter 9 Natural Products from Plants with Antimicrobial Action**



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**Abstract** In recent years, there has been a continuous increase in resistance to traditional antibiotics developed by pathogenic bacteria. However, in many parts of the world, several medicinal plants are traditionally used to control infectious microorganisms. Because of this, antimicrobial agents derived from natural products have received a lot of attention, both for their effectiveness and also for being more economically accessible. Therefore, the research discussed in this chapter aims to conduct a systematic review on the use of medicinal plants and isolated compounds as a potential antimicrobial agent. The study focuses on an investigation of several electronic databases: Scopus, Web of Science, Academic Google, SciELO, PubMed, SciFinder, and ScienceDirect. The results may contribute to the increase of strategies for the treatment of infections caused by microorganisms. Since medicinal plants play an essential role in health, they may represent a signifcant source of new antimicrobial drugs to combat microorganisms resistant to multidrug.

**Keywords** Natural products · Antimicrobial action · Plants · Antimicrobial resistance

# **Abbreviations**



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## **1 Introduction**

Starting from the twentieth century, there were signifcant advances in the feld of science with the use of antimicrobial drugs, such as penicillin which was discovered in 1928 by Sir Alexander Fleming, intending to improve human life expectancy and control microbial infections (Aminov [2010](#page-11-0)). Antimicrobials can be chemotherapy (synthetic) or antibiotics (natural substances) that inhibit the development of microorganisms or even destroy them. Antimicrobial drugs are considered the secondmost used drug class globally (Saez-Llorens et al. [2000](#page-14-0); Abushaheen et al. [2020\)](#page-11-1).

As these drugs are more widely used, microorganisms develop resistance, quickly causing a global problem (Anand et al. [2020](#page-11-2)). The use of antimicrobial drugs without a medical prescription is a problem reported at least half a century ago (Scheckler and Bennett [1970;](#page-14-1) Kunin et al. [1973](#page-13-0)). This is due to the misdiagnosis between a bacterial infection and a viral infection, the absence of educational programs for the rational use of antimicrobials or the wide distribution of antimicrobials. In this way, the use of these drugs and improper self-medication exacerbate antimicrobial resistance (Mota et al. [2010](#page-13-1)).

This resistance to antibiotics has become one of the main problems of humanity since the end of the twentieth century. Therefore, the search for new antimicrobials effective in eliminating microbes becomes necessary, given that the traditional approaches to fnd new antimicrobial drugs are no longer effective due to the rapid resistance developed against them (Abreu et al. [2012](#page-11-3)).

One of the strategies applied for the treatment of infectious diseases would be natural products. For several years, medicinal plants derived from these natural products play an essential role in health. They could represent a signifcant source of new antimicrobial drugs to combat multidrug-resistant microorganisms (Aleksic Sabo and Knezevic [2019\)](#page-14-2). Besides, antimicrobial resistance to plant extracts would be less likely, since a wide variety of their active compounds are found in these extracts that could reverse antibiotic resistance and minimize man's exposure to resistant bacteria (Gupta and Birdi [2017](#page-12-0)).

Bioactive compounds called natural products come from the biosynthesis of plants, lichens, fungi, bacteria, and microorganisms. Antibiotics, terpenoids, alkaloids, and polyketides are the main microbial natural products (Gunatilaka and Wijeratne [2012](#page-12-1); Huang and Lin [2017](#page-13-2); Newman and Cragg [2016\)](#page-14-3).

It is noteworthy that approximately 50% of approved drugs are related to plants or natural products, and 10% of all antimicrobial agents are from natural plant products (Al-Marzoqi et al. [2016;](#page-11-4) Cowan [1999](#page-12-2); Veeresham [2012\)](#page-15-0). In this way, natural products are still used as a powerful therapy against pathogenic bacteria and are considered the pillars of discovering new antibiotics (Wright [2017\)](#page-15-1).

According to Moore et al. ([2017\)](#page-13-3), there is an evident lack of new antimicrobial drugs. They are emphasizing the need to search for new bioactive therapeutic agents as treatment strategies for infections caused by microorganisms. In addition to medicinal plants, critical antimicrobial properties have been evidenced, demonstrating the enormous potential that does not control infectious diseases or that suggests as a valuable source of research for the removal of antimicrobial compost.

#### **2 Antimicrobial Resistance**

Antimicrobial resistance is a global public health problem, which has worsened over time, as infections caused by resistant microorganisms, in many cases, do not respond to treatment, which can increase hospital costs and increase the risk of death (Fair and Tor [2014](#page-12-3)). Some studies report that if there are no measures to neutralize this problem, there may be a beginning of a post-antibiotic era, with a shortage of effective treatments based on antimicrobials (Vasoo et al. [2015;](#page-15-2) Vanegas-Múnera and Jiménez-Quiceno [2020](#page-15-3)).

As these antimicrobials are indiscriminately used, microorganisms were developed by creating different forms of resistance, which led to a global problem (Davies and Davies [2010](#page-12-4)). Alexander Fleming, who received the Nobel Prize in Medicine in 1945, warned about the severe problem of antimicrobial resistance. Still, human being did nothing, and thus resistant bacteria emerged as a result of natural selection (Ventola [2015\)](#page-15-4). This problem worsened over time due to the inappropriate use of antibiotics, was also favored by the lack of standards and control of their service, such as the ease of acquiring these drugs without a prescription, in addition to false marketing of antimicrobials (Organización de las Naciones Unidas para la Alimentación y la Agricultura [2016](#page-14-4)).

The frst reports of antimicrobial resistance were published in 1944, three years after starting the use of penicillin. The isolates of *Staphylococcus aureus* were reported, these being the frst isolates resistant to this antibiotic. Since then, the situation has only worsened; at times in the 1980s, there were few options of antibiotics available (Tang et al. [2014\)](#page-14-5). Currently, the resistance rate is so high that some doctors use antibiotics that were previously discarded, because they have specifc toxicity, such as colistin (Loho and Dharmayanti [2015\)](#page-13-4).

Antimicrobial resistance can be of two types: natural or acquired. Natural or intrinsic, occur when there are bacteria of the same species naturally resistant to some families of antibiotics, which leads these bacteria to have a competitive advantage over other strains; therefore, the antibiotic has no effect. The acquired resistance can be attributed to mutations in chromosomes (stable, spontaneous, and transmitted vertically from generation to generation) or to changes in resistance genes (conjugation, translation or transformation and transmitted horizontally) (Guimarães et al. [2010;](#page-12-5) Pérez [2017\)](#page-14-6).

The increase in antimicrobial resistance has severe consequences for public health, limiting the therapeutic variability of antimicrobial in combating pathologies in humans (Semret and Haraoui [2019](#page-14-7); Costa et al. [2020\)](#page-11-5). This increase in antimicrobial resistance is due to the misuse of these agents to resistant strains, such as methicillin to *Staphylococcus aureus*, vancomycin-resistant enterococci, resistance to drugs *Streptococcus pneumoniae* and *Mycobacterium tuberculosis*, enterobacteria, which produce beta-lactamase, extended-spectrum, and resistant to carbapenems, *Pseudomonas aeruginosa*, resistant to multiple drugs and *Acinetobacter baumannii* (Sabo and Knezevic [2019\)](#page-14-2).

Thus, fungi became resistant to polyols, azoles, and echinocandins, which contributed to the emergence of drug-resistant strains and were found in all species of fungi (Robbins et al. [2017\)](#page-14-8). An aggravating factor given the remarkable emergence of resistant protozoa and viruses is a limited number of antiprotozoal and antiviral agents (El-Taweel [2015](#page-12-6); Irwin et al. [2016\)](#page-13-5).

Plants are of great importance in modern medicine. Thus, the plant kingdom can be an alternative to combat antimicrobial resistance, so there is a search for new antimicrobial drugs that can combat resistant microorganisms. This can play an essential role in combating microbial resistance (Sabo and Knezevic [2019\)](#page-14-2). However, there is little incentive from the pharmaceutical industries to produce new antimicrobial, mainly due to the low fnancial return, thus limiting research for this type of medicine (Aslam et al. [2018](#page-11-6)).

With the current antimicrobial resistance, a post-antibiotic era is getting closer and more concrete every day. This problem is due to not only the evolution and adaptation of these microorganisms but also by those who prescribe these drugs, by those who use them, by those responsible for the control and distribution of these drugs (Vanegas-Múnera and Jiménez-Quiceno [2020](#page-15-3)). Thus, public policies are needed to raise awareness and educate the population about the use of antimicrobial, as well as health professionals can limit the unnecessary use of antimicrobial (Hu et al. [2020\)](#page-13-6).

# **3 Classifcation and Mechanism of Action**

Antimicrobials are classifed according to their chemical structure (derived from amino acids, sugars, acetates, propionates, among others), types of microorganisms on which they act, or even the effect of the organism (Manrique and Galvão [1997\)](#page-13-7). Besides, they are classifed through factors, such as the mechanism of action, bacterial activity, susceptible microorganisms, the spectrum of action, and mainly through the assessment of their natural or synthetic origin (Anvisa [2020](#page-11-7)).

The mechanism of action of an antimicrobial starts when it reaches an ideal concentration at the infection site, then actively or passively passes through the cell wall, showing an affnity for the binding site inside the bacteria and remaining long enough to exercise its inhibitory activity. Antibiotics can have bactericidal action, killing microorganisms directly, acting in strong reactions to the infecting cell, or just inhibiting bacterial growth through a bacteriostatic action, keeping the bacteria in a stationary phase. In bacteriophage, the host triggers its defenses, such as phagocytosis and the production of antibodies to control the invading microorganism. In this case, the inhibition can be reversible, as the bacteria can continue to produce toxins or become resistant to the drug if the host's defenses are not efficient Fig. [9.1](#page-4-0). (Katzung [2007](#page-13-8); Lago [2011](#page-13-9); Brunton et al. [2012](#page-11-8); Tortora et al. [2012;](#page-15-5) Pankey and Sabath [2004;](#page-14-9) Engleberg et al. [2013](#page-12-7)).

<span id="page-4-0"></span>

Source: Persona data.

**Fig. 9.1** Mechanism of action of antibiotics. (*Source*: Personal data)

Regarding natural antibiotics, together with their semi-synthetic derivatives, they comprise most of the antibiotics used clinically and can be classifed into β-lactams (penicillins, cephalosporins, carbapenins, oxapenins, and monobactams), tetracyclines, aminoglycosides, glycopeptides, lipodepsipeptides, streptogramins, licosamines, chloramphenicol, among others. Antibiotics of synthetic origin are classifed as sulfonamides, fuoroquinolones, and oxazolidinones (Abraham [2003](#page-11-9); Patrick [2005;](#page-14-10) Pupo et al. [2006](#page-14-11)).

Other important examples of antimicrobial classes are Macrolides (Bologa et al. [2013\)](#page-11-10), Glycopeptides (Guimarães et al. [2010\)](#page-12-5), Amphenicols (Von [2004](#page-15-6)), and Quinolones (Emmerson and Jones [2003](#page-12-8)).

### **4 Herbal Medicines with Antimicrobial Action**

The frst antimicrobial drugs used successfully against potentially fatal infectious diseases were Salvarsan (arsphenamine) and its derivative neoarsphenamine, used against syphilis (infection with *Treponema pallidum*) in 1911 (Mann [1984](#page-13-10); Gaynes [2017\)](#page-12-9). During the late 1930s and early 1940s, sulfamide antibiotic derivatives (e.g., prontoil, sulfamethazine) effective against lobar pneumonia (caused mainly by *Streptococcus pneumoniae*), meningococcal meningitis (caused by *Neisseria meningitidis*), and gonorrhea (caused by *Neisseria gonorrhoeae*) were introduced to the market (Kumar and Clark [1998](#page-13-11); Gaynes [2017\)](#page-12-9). However, until the discovery of penicillin (USA: 1945; United Kingdom 1946), these synthetic agents remained the only chemotherapeutic agents used for bacterial infections (Mann [1984\)](#page-13-10).

However, the widespread use of antibiotics created an evolutionary adaptation of bacteria, leading to the so-called resistance to multiple drugs (Neu [1992\)](#page-14-12). The consequence of the increased resistance of bacterial strains to biocidal agents is an increased risk of chronic infections and diffculties in their treatment (wound infections, osteomyelitis, septic arthritis, endocarditis, etc.) (Karam et al. [2016\)](#page-13-12). However, there is a strong need to develop new compounds that are not only highly effcient but also that should not cause resistance to development in bacteria (Kyzioł et al. [2020\)](#page-13-13).

In search of these new antibiotics, research on herbal medicines may prove promising since these medicines have been used for centuries to treat infectious diseases (Landers et al. [2012\)](#page-13-14). They are readily available for patients to purchase themselves and appear to be increasingly popular, especially in the United Kingdom. This growing public interest in herbs can reduce dependence on antibiotics, especially for self-limited infections (Hu et al. [2020](#page-13-6)). A recent WHO report on traditional medicines noted that most of the world's population depends on traditional medicines for health care, including treatment of infections (WHO [2002\)](#page-15-7).

Thus, compounds isolated from medicinal plants are preferable to synthetic compounds due to their use in conventional medicine (Rakholiya et al. [2013\)](#page-14-13). Therefore, these compounds are considered a substituted source of antimicrobial drugs (Savoia [2012](#page-14-14)).

## **5 Plant Antimicrobial Activity**

#### *5.1 Extracts with Antimicrobial Action*

Natural products are used since ancient times for the treatment and, or control of infectious diseases among them from fungi, bacteria, or pathogenic microorganisms (Dkhil et al. [2020;](#page-12-10) Frassinetti et al. [2020;](#page-12-11) Silva [2006](#page-14-15)). These products include extracts, fractions, essential oils, isolated compounds, among other medicinal plant derivatives, which may be promising for the advancement of new studies, therapies, and treatments for pathologies (Frassinetti et al. [2020;](#page-12-11) Mehlhorn [2014;](#page-13-15) Hayek et al. [2013\)](#page-12-12). Thus, some tasks related to the development and progress of studies on prominent plant extracts, in potential, with antimicrobial action will be addressed.

Antimicrobial activities of the ethanolic extracts of the fowers and leaves of ten species of the genera *Senna* and *Cassia* were evaluated against seven bacteria and three strains of fungi, oral aerobic and anaerobic bacteria, and Candida spp. by the microdilution broth method. This investigation was due to the use of these species in treatments of infections in the Brazilian traditional medicine, used as laxative agents, analgesics, and antifungals for mycosis and other fungal infections of the skin. Nascimento et al. ([2020\)](#page-14-16) developed this work, showing that among the tested species, *Cassia fstula* L.*, Senna macranthera* (Collad)*, Cassia bakeriana* Craib, and *Senna spectabilis* DC*.*, showed moderate activity against two bacterial strains with MIC values (minimum inhibitory concentration) varying between 200.0 and 400.0 μg.mL−<sup>1</sup> , while the ethanolic extract of *S. macranthera* fowers showed very low values of MIC (23.4, 11.7, and 5.9  $\mu$ g mL<sup>-1</sup>) in the antifungal test.

Thus, the ethanol extract of *S. macranthera* flowers was subjected to liquidliquid extraction with solvents of different polarities, which include n-hexane, dichloromethane and ethyl acetate. The ethyl acetate fraction of *S. macranthera* fowers showed better antifungal activity (MIC values of 5.9 μg mL−<sup>1</sup> for *Candida glabrata,* 23.4 μg mL−<sup>1</sup> for *Candida albicans,* and (0.1 − 0.2 μg mL−<sup>1</sup> ) for *Candida tropicalis* using amphotericin B as a positive control, so this fraction indicated potential antifungal action and selectivity against strains evaluated. It is noteworthy that the MIC represents the lowest concentration of the extract capable of preventing microbial growth. From this fraction, analysis by UHPLC-ESI-MS/MS (High-Effciency Liquid Chromatography Coupled by Tandem Mass Spectrometry) were performed, which identifed different types of phenolic compounds, in particular, proanthocyanidins in several isomeric forms.

This class of secondary metabolites, proanthocyanidins (condensed tannins), are phenolic compounds (Kyraleou et al. [2020;](#page-13-16) Monteiro et al. [2005\)](#page-13-17), active against species of *Candida* and thus may be associated with the antifungal action of the fraction cited (Piccinelli et al. [2016](#page-14-17); Freitas et al. [2018\)](#page-12-13). This study has collaborated to strengthen the traditional use of the species from these genera *Senna* and *Cassia*, particularly the fowers of *S. macranthera,* as a good and promising source of discoveries of compounds as antifungal agents.

Weremczuk-Jeŝyna et al. ([2019\)](#page-15-8) evaluated the anti-pathogenic potential of the hydromethanolic extract of *Dracocephalum forrestii* sprouts cultivated in nutrientsprinkled bioreactors. These antimicrobial evaluations were performed against selective strains of bacteria, six Gram-pathogenic, and four Gram-negative, and three pathogenic strains of fungi, which showed moderate activity limiting the growth of pathogens. The minimum bactericidal concentration (MBC) and MIC activity of the extract were analyzed, where most of the strains tested were within the range of 2.5–5 mg.mL<sup>-1</sup>.

A greater effect was observed on *Bacillus cereus*, *Escherichia coli*, and *Pseudomonas aeruginosa* with MIC 2.5 mg mL<sup>-1</sup>. The hydromethanolic extract at a concentration of 2.5 mg mL−<sup>1</sup> . showed antifungal activity against *Candida albicans* and *Candida glabrata. They were determined by the microdilution broth method according to the European Committee for Antimicrobial Susceptibility Testing (EUCAST). Rosmarinic acid, gentamicin, and fuconazole (all from Sigma-Aldrich) were used as reference antimicrobials with values lower than 1–5 mg.mL*−<sup>1</sup> .

In this work, the phenolic compounds present in the cultivated extract were evaluated qualitatively by UPLC-PDA-ESI-MS (ultra-performance liquid chromatography (UPLC) coupled to photodiode array detection (PDA) and electrospray ionization (ESI) mass spectrometry) and quantitatively by UPLC-PDA. Rosmarinic acid (17.90 mg g−<sup>1</sup> DW) and salvianolic acid B (6.50 mg g−<sup>1</sup> DW) were verifed in larger quantities. Deba et al. ([2008\)](#page-12-14) and Estevinho et al. [\(2008](#page-12-15)) reported that plant material rich in polyphenols, bioactive phenolic compounds, may be associated with anti-pathogenic effects due to the interaction between these constituents. The author suggests further studies to improve the research and thus to estimate a possible synergistic effect of *D. forrestii* extract on a likely antibiotic therapy against multi-resistant bacteria.

Fei et al. [\(2018](#page-12-16)), in their studies, reveal that the potent antioxidant activity of two polyphenols can be considered as one of the ways this class of natural products inhibits the development of pathogenic microorganisms.

#### *5.2 Phytoconstituents with Antimicrobial Action*

The use of antibiotics and/or synthetic or natural compounds to treat infectious diseases is growing widely (Hemaiswarya et al. [2008;](#page-13-18) Bimani and Hossain [2020\)](#page-11-11). Thus, scientists are increasingly improving their search for fnding natural compounds bioactive that may be able to exercise biological activities to treat, improve, or resolve the resistance of microorganisms to synthetic drugs (Bimani and Hossain [2020;](#page-11-11) Vidhya et al. [2020\)](#page-15-9).

Andrade et al. ([2020\)](#page-11-12) evaluated the antimicrobial activity of two isolated natural molecules, Braquidines BR-A and BR-B, against *S. aureus*, *E. coli*, and *C. albicans*. The isolation and analysis were performed from the purifcation of the dichloromethane fraction of *Arrabidaea brachypoda* fowers, by HPLC-PCDA

(High-Effciency Liquid Chromatography with photodiode arrangement detection) and UV-VIS (Ultraviolet Spectroscopy-visible) revealing and identifying phenolic compounds, dimeric favonoids differentiating only in the aromatic C ring by the presence of a hydroxyl group in BR-A, already in BR-B there is a methoxyl group, giving purity of 95 and 97%, respectively. BR-A showed no antimicrobial activity against the microbial strains tested, since MIC values above 1000 μg.mL−<sup>1</sup> , presenting a MIC of 1024 μg.mL−<sup>1</sup> , were determined by microdilution assay. Thus, according to the literature, these are considered clinically irrelevant results (Houghton et al. [2007\)](#page-13-19). However, the BR-B showed antifungal activity against *C. albicans* of 161 μg.mL−<sup>1</sup> , showing promising future studies and improvements.

A phytochemical study on *Trianthema decandra* leaves was performed by Geethalakshmi and Sarada ([2018\)](#page-12-17), where two compounds of chloroform extract from the leaf were isolated and characterized using HPLC, UV, FT-IR, RMN, LC-MS, and CHNS techniques. Identifying a new sterol named as 17-(5-ethyl-6 methylheptan-2-yl) – 4, 4, 10, 13-tetramethyl-hexadecahydro-1H-cyclopenta  $(\alpha)$ phenanthren-3-ol and the favonoid as 2-(3´,4'dihydroxyphenyl) 3,5,7-trihydroxychromen-4-one. The antimicrobial activity of the sterol and favonoid isolates were analyzed by disc diffusion and broth dilution assays, showing good results against the tested microorganisms, in particular sterol presented an MIC of 39.05 μg mL−<sup>1</sup> against the strain *Salmonella typhi* and the favonoid presented MICs of 78.10 and 39.05 μg mL−<sup>1</sup> against *Vibrio cholerae* and *P. aeruginosa,* respectively.

## *5.3 Essential Oils with Antimicrobial Action*

A variety of oils essential (OEs), derived from aromatic plants, are used as insecticide, antiparasitic, fungicide, virucidal, bactericide, cosmetics, food, and agricultural industries (Dhif et al. [2016;](#page-12-18) Atif et al. [2020\)](#page-11-13). These OEs also act as preservatives, sedatives, antimicrobial, and spasmolytic analgesics. The chemical composition of OEs includes sesquiterpenes, monoterpenes, and their oxygenated compounds, such as oxides, phenols, ketones, ethers, esters, aldehydes, molecules containing nitrogen or sulfur (Ahmed et al. [2019](#page-11-14)).

With the presence of secondary metabolisms, OEs play a protective role against various microbes in plants. The actions generated by these metabolites can inhibit or slow the growth of bacteria, yeasts, and molds, whose components have a variety of targets, particularly on the microbial membrane and cytoplasm, and in some situations, they drastically alter cell morphology (Chorianopoulos et al. [2008](#page-11-15); De Martino et al. [2009\)](#page-12-19).

Because they have antimicrobial activity against several microorganisms, OEs are considered an alternative to conventional antibiotics (Silva-Santos et al. [2017\)](#page-14-18).

Herbs such as cloves, cinnamon, oregano, peppers, rosemary, and thyme have OEs that generate intense antibacterial activity against *S. aureus, S. typhi*, and *P. aeruginosa* (Conner and Branen [1993\)](#page-11-16). Among all the essential oils tested, clove

oil was the most effective. In addition, it was found that carvacrol has antimicrobial activity against a broad spectrum of bacterial strains (Fernández-Pan et al. [2015;](#page-12-20) Kristo et al. [2008](#page-13-20)).

Mohamed et al. ([2013\)](#page-13-21) reported effective inhibition in pathogenic bacterial strains such as *N. gonorrhoeae*, *E. coli*, *Bacillus subtilis*, *S. aureus,* and *P. aeruginosa*, developed by *Syzygium cumini* OE, containing 1,3,6 limonenoene, transcarcinogenic-octatriene, β-pinene, δ-3-carene, α-pinene and α-caryophyllene; these compounds are responsible for the bactericidal action.

Because they come from plant sources, OEs can also develop synergistic activity, which effectively combat the growth of various microorganisms. Some constituents such as carvacrol, γ-terpinene, and p-cymene are more effective when combined (Elshafe and Camele [2017\)](#page-12-21).

OEs are mentioned as potent antimicrobial agents, as they have substantial antibacterial activities against Gram-positive and Gram-negative pathogens (Yap et al. [2014\)](#page-15-10). For example, OEs derived from medicinal aromatic plants, such as peppermint (*Mentha piperita*), thyme (*Thymus vulgaris*), and fennel (*Foeniculum vulgare*), are cataloged as effective against viruses, Gram-negative and Gram-positive bacteria, fungi, and yeast (Reichling [2018](#page-14-19)).

Various volatile, terpenic, and phenolic substances present in OEs show remarkable antimicrobial activity (Marchese et al. [2016](#page-13-22)). And according to the literature, OEs that contain these terpenes, phenolic, and aldehydes have an excellent application in biomedicine for properly eliminating many viral, fungal, and bacterial pathogens (Swamy et al. [2016](#page-14-20)).

#### **6 Perspectives with the Use of Secondary Plant Metabolites**

Antimicrobials have played an essential role in human health since their discovery. This relevance is refected in the results obtained using antimicrobial pre-agents, which are also among the drugs most used by medicine. The discriminated and illicit use of antibiotics has resulted in the emergence of drug resistance among pathogens, which in many parts of the world, especially in developing countries, have reached critical levels (Ayukekbong et al. [2017\)](#page-11-17).

This practical and negligent use of antibiotics is a growing concern worldwide, fueling antimicrobial resistance. In the clinical context, particularly in the health care sector, it is vital to validate a prescription for antibiotics before administering the medication. Besides, the dosage of antibiotics must be such that pathogenic bacteria are eliminated in a complete cycle of antibiotics. However, there are few effective techniques for a quick discovery of susceptibility to drugs. Continuous unintended exposure of bacteria to antibiotics can promote antimicrobial resistance that can increase in the body for longer in most living organisms. The development of technologies to detect antibiotics (mainly used in the therapy of human infection) is essential to containing the threat of antimicrobial resistance to guarantee its ideal use in the treatment of human bacterial infection (Nag et al. [2020\)](#page-13-23).

Antimicrobial products characterized as drugs have a shelf life, changes in the metabolic and genetic level, a faster rate of evolution such as variable global temperature, a catalog of widely documented side effects. However, the main concerns of doctors and scientists are the prolonged use of antimicrobials, the high cost of clinical research and drug development. The main obstacle to antibiotic resistance is developing effective classes of antibiotics with a new mode of action (Anand et al. [2020\)](#page-11-2).

It should be noted that more than 80% of the world population depends on conventional pharmaceutical products made up of medicinal plants to meet different conditions of human medicine (Ekor [2014;](#page-12-22) Oyebode et al. [2016](#page-14-21); Pan et al. [2014;](#page-14-22) Mahomoodally [2013\)](#page-13-24). Worldwide interest in research aimed at discovering new drugs from plant sources is growing (Anand et al. [2020\)](#page-11-2).

The natural products are characterized as a diverse group with different bioactivities for therapeutic purposes. Natural sources are privileged source in searching for new antimicrobial molecules (Agrawal et al. [2017\)](#page-11-18). More than half of medicines in clinical use approved by the US Food and Drug Administration (FDA) are derived from natural products (Newman and Cragg [2016](#page-14-3)).

Therefore, the work appears with the perspective of contributing to the development of new antimicrobial drugs to combat microorganisms resistant to multidrug drugs, in addition to enabling positive actions about natural compounds derived from plants, which may later be the basis of pre-clinical studies in humans and in a future perspective, which will contribute to implementation of these compounds of natural origin in the treatment of pathologies.

#### **7 Conclusions**

This chapter pointed out that phytochemicals, extracts, and derivatives of natural compounds could be effective in the treatment of resistant microorganisms, since they have antimicrobial activity against microorganisms, especially in the therapeutic approach or acting as drugs. In this way, natural products may serve as a promising therapeutic source for various pathogenic microorganism, and therefore are pillars for discovering new antibiotics. However, the pharmacological utility of these products will require further studies. Thus, this review becomes extremely important, as it shows medicinal plants as a source of new therapies that can contribute to the discovery of potential candidates for the treatment of resistant microorganisms.

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