



Short review on the potential alternatives to antibiotics in the era of antibiotic resistance

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ABSTRACT

The importance of antibiotics in the treatment of infectious diseases of bacterial origin is well established. Despite the positive contribution of antibiotics in veterinary and clinical medicine, some of these molecules were regarded as harmful to both human and animal health with some undesirable effects. Moreover, microorganisms have developed resistance mechanisms to almost all commonly used antibiotics. Hence, there is an urgent need to find alternative ways against bacterial resistance to antibiotics which constitute a public health concern worldwide. The search for solutions to overcome this problem prompted researchers to seek new methods and molecules to continue the treatment of bacterial infections while limiting the spread of resistance and thwarting already resistant germs. The present study is new and current because it presents in a single document the most known alternatives to antibiotics. Therefore, the aim of this brief review was to discuss methods that can constitute a valid way to reduce the use or even replace conventional antibiotics. We found that phage therapy, probiotics, antimicrobial peptides, vaccines, medicinal plants, nanoparticles, antibodies, and cytokines are among the most promising alternatives and more investigation should focus on them.

INTRODUCTION

An antibiotic is a natural or synthetic substance that kills bacteria or blocks their growth (Lewis, 2020). Antibiotics act on the bacterium by attacking either their cell wall (Kobras *et al.*, 2020), their cytoplasmic membrane (Heesterbeek *et al.*, 2021), their protein synthesis metabolism (Stokes *et al.*, 2019), their nucleic acid (RNA and DNA), or even more than one of these elements (Grandclaoudon *et al.*, 2020). Since their discovery, many antibiotic molecules have emerged and their role in the treatment

of various bacterial infections in both humans and animals is no longer to be demonstrated. In the United States, recent estimates show that 80% of antibiotics are used in livestock breeding as feed additives and growth promoters to improve the quality and yield of production (Ventola, 2015). Unfortunately, these antibiotics are excreted by animals in feces and then end up in the groundwater environment or are used as fertilizers (Ventola, 2015). As a result of this practice, beyond the direct environmental impacts, we are witnessing the emergence of increasingly resistant and multiresistant microorganisms (Ventola, 2015). This adaptation phenomenon is mainly due to the enzymatic degradation of antibiotics by bacteria (Reis *et al.*, 2020), the modification of the target of the antibiotic (Schaezner *et al.*, 2020), the change in membrane permeability (Xu *et al.*, 2020), and alternative metabolic pathways (Pollock *et al.*, 2020). Interbacterial transmission of antibiotic resistance through horizontal gene

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transfer (conjugation, transduction, and transformation) has made the situation critical worldwide (Lima *et al.*, 2020). Broadly speaking, antibiotic resistance is defined as the ability of a bacteria to resist the inhibitory or destructive activity of an antimicrobial to which it was not resistant (Palma *et al.*, 2020). This problem concerns both veterinary medicine and human medicine, and every year numerous studies demonstrating the increasing resistance of bacterial strains are carried out (Mbarga *et al.*, 2020; Rabello *et al.*, 2020). In hospitals, the most resistant germs encountered are *Staphylococcus aureus*, Coagulase-negative *Staphylococci*, *Pseudomonas aeruginosa*, *Klebsiella pneumoniae*, *Escherichia coli*, *Enterobacter* spp., *Enterococcus* spp., and *Acinetobacter* (Antoinette and Dieudonne, 2020; Cleven *et al.*, 2006). These germs are generally involved in various classic infections and some of them are sometimes implicated in nosocomial infections (Cleven *et al.*, 2006). According to several researchers, the fight against antibiotic resistance requires a more measured use of antibiotics, a prescription exclusively reserved for professionals, systematic antibiograms before administration, and the search for new effective molecules on already resistant germs (Cleven *et al.*, 2006; Lima *et al.*, 2020; Palma *et al.*, 2020). However, beyond the search for new antibiotics commonly called classic, several studies have been carried out to propose alternative techniques and methods having an effect on the bacteria. This review aims to discuss, without claiming to be exhaustive, most of the possible alternatives to conventional antibiotics.

REVIEW METHODOLOGY

This review article was carried out by exploiting numerous review articles, original articles, and related books from *reputable* databases, such as Web of Science, PubMed, and Scopus. No open access published papers have been made available using the facilities provided by the People's Friendship University of Russia, Moscow, Russia. The literature investigation process was conducted between June 2020 and June 2021 and the literature investigations were conducted in English and French. The keywords explored during literature searching included combinations of the words constituting the titles of each section, that is, "resistance to antibiotics" "phage therapy", "antimicrobial peptides" (AMP), "combination of antibiotic and antimicrobial peptides", "vaccines and antibiotics", "herbal medicine", "antimicrobial activity of plant materials", "modulation of antibiotics with plant materials", "antibacterial properties of nanoparticles", "probiotics and antibiotics"; "phagothérapie", "plantes médicinales", "activité antibactérienne des nanoparticules", "peptides antimicrobiens", and "synergie entre les extrait des plantes et les antibiotiques classiques".

Resistance to Antibiotics and Current Issues

Antibiotic resistance poses one of the most serious threats to global health, food security, and development today (World Health Organization (WHO), 2019). An increasing number of infections, such as *pneumonia*, tuberculosis or gonorrhea, and salmonellosis, are becoming more difficult to treat as the antibiotics used to treat them lose their effectiveness (WHO, 2019). This situation is likely to affect people of all ages and genders in all corners of the globe and both the losses and the dangers it represents are considerable. Indeed, antibiotic resistance leads to

prolonged hospitalizations, an increase in medical expenses, and, increasingly, an increase in mortality (Jit *et al.*, 2020; WHO, 2019). Likewise, recent estimates have shown that antibiotic resistance is responsible for 700,000 annual deaths worldwide, 230,000 of which have resulted from multidrug-resistant tuberculosis (WHO, 2019). Furthermore, the WHO, 2019, estimates that if nothing is done to address this problem, drug-resistant diseases may cause 10 million deaths each year by 2050 and damage to the economy as catastrophic as the 2008–2009 global financial crises. In addition, economically (linked directly or not to agriculture and animal breeding), antimicrobial resistance could force up to 24 million people into extreme poverty by 2030 (WHO, 2019). However, in order to reduce the spread of this phenomenon, the researchers recommend more careful use of these substances both in animal breeding and in human medicine (Doidge *et al.*, 2020; Morel *et al.*, 2020). This consists mainly of stopping the use of antibiotics in breeding as growth factors and their use only when necessary to treat identified infections, under prescription, and with prior antibiograms (Mbarga *et al.*, 2020; WHO, 2020). Moreover, it is also suggested that a major part of the solution to this problem lies in the establishment of new therapeutic protocols which include new biologically active compounds (Morel *et al.*, 2020) including AMP, nanoparticles (NPs), antibodies, cytokines, phytochemicals, viral particles (bacteriophages), and microorganisms (probiotics).

Phage Therapy

The first use of phage therapy for the treatment of patients dates to 1919 when Félix d'Hérelle used them at the Institut Pasteur in Paris (Dublanche and Schwartz, 2017). After this discovery, the practice quickly spread and the establishment of therapeutic protocols involving phages as a means of combating bacterial diseases was recognized by many scientists (Summers, 2005). However, it was only in 1925 that the first mass treatments took place in Brazil at the Oswaldo Cruz Institute and the results published by this institute were very promising. Among the first 10,000 cases treated, only 1 patient remained ill although his recovery was observed later (Biacchesi *et al.*, 2017). Despite the good results obtained by this method, it has been abandoned in favor of the use of antibiotics because of their simplicity of handling, production, and use (Mireille *et al.*, 2020). Today, due to the growing problem of multidrug resistance to antibiotics, phages are gradually regaining their place in the fight against bacterial infections although their use is still almost limited to highly equipped centers.

Mechanism of action of phages and application in medicine

Unlike conventional antibiotics whose mechanism of action is to attack one or more elements of the bacterial cell (wall, cytoplasmic membrane, metabolic pathways, or nucleic acids), phages have their own mechanisms and present a highly narrow specificity with the target bacteria (Koskella and Meaden, 2013). Without the host, it is impossible for them to multiply. As shown in Figure 1, the phage replication mechanism is like that of normal viruses. This figure shows that, after the attachment or adsorption (1) of the phage on the bacterial cell, the latter makes (2) its DNA penetrate the bacterium and rout all the enzymatic machinery of the latter to synthesize (3) the genome and viral proteins. After this step, the synthesized constituents will assemble (4) and cause

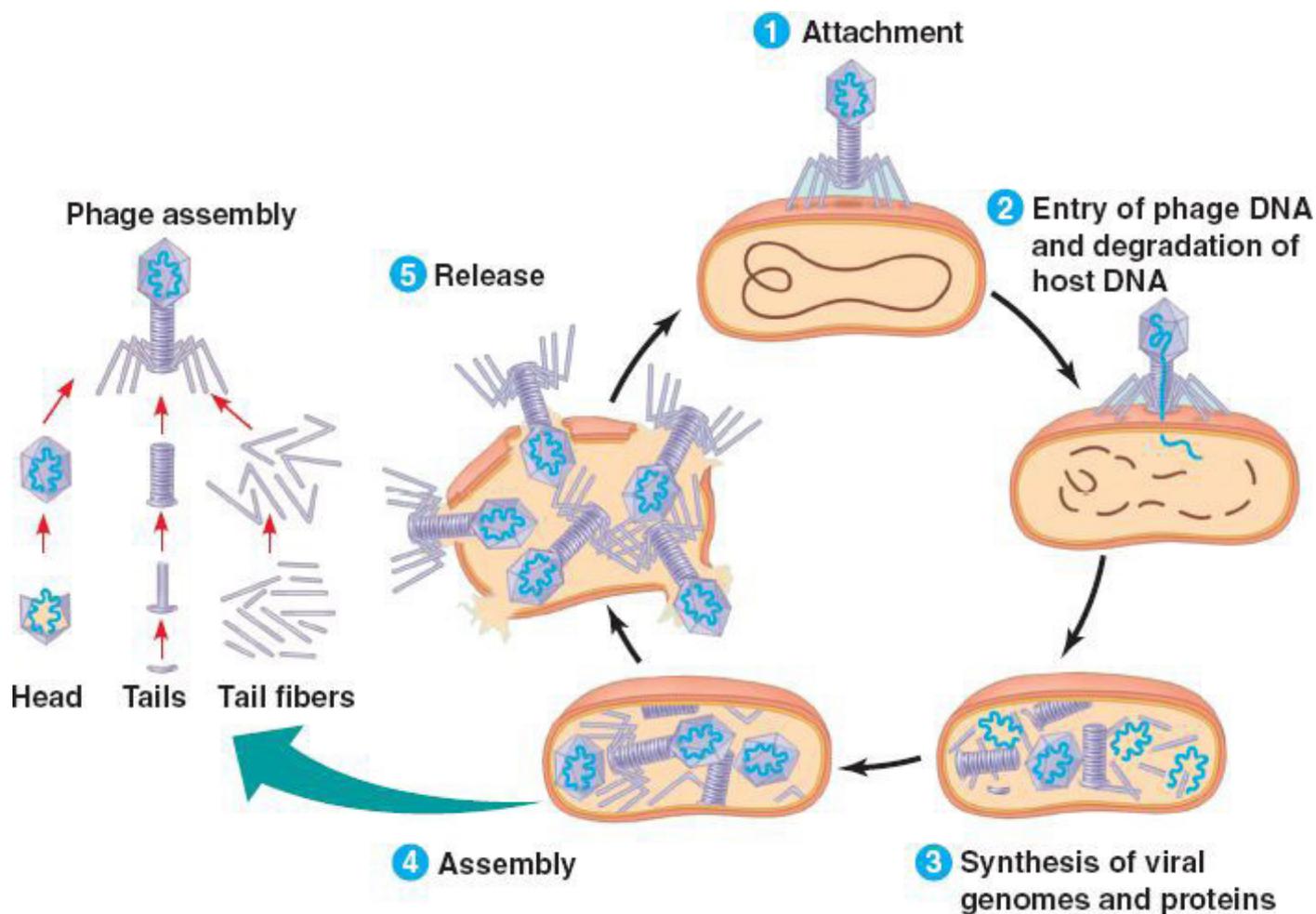


Figure 1. Phage mechanisms of action (Jun, 2017).

the lysis of the bacterial cell which will lead to the release (5) of the new phages formed and the phenomenon will continue until the lysis of all the target bacteria (Jun, 2017). The time interval between attachment and cell lysis generally depends on several parameters (Van Belleghem *et al.*, 2018). Indeed, modeling the pharmacokinetics of bacteriophages requires considering parameters related to their viral cycle, such as the adsorption constant of the viruses on the host bacteria, the number of virions released per cycle, the latency time, the initial number of bacteria values, and the bacteriophage/bacterium ratio (Van Belleghem *et al.*, 2018). Likewise, the minimum bacterial density to ensure initiation of viral multiplication/transmission must be known. The modeling becomes even more complex when one considers the dynamics of the bacterial population over time (the uninfected cells continuing to replicate) as well as the phenomena of resistance to bacteriophages and their amplification over time treatment (Skurnik and Strauch, 2006). This pharmacokinetics is barely addressed in experimental animal studies, typically because of its complexity (determination of viral constants *in vivo*, bacterial inoculum, spontaneous viral elimination constant, etc.). It is, however, a fundamentally important approach and failure to take this into account could explain several previous failures of phage therapy (Payne and Jansen, 2001). Theoretical modeling

of pharmacokinetics has thus made it possible to correlate the predicted results positively and significantly with what is observed *in vitro* (Cairns *et al.*, 2009) but with difficulty *in vivo* (Weld *et al.*, 2004), proving the need to continue studies in this area.

However, despite the complexity of mastering phage pharmacodynamics *in vivo*, many recent studies have demonstrated their efficiency in human medicine. This is the case of a study conducted in the UK in 2018 which demonstrated the therapeutic potential of three bacteriophages (*Escherichia* phage ECP311, *Klebsiella* phage KPP235, and *Enterobacter* phage ELP140) against *E. coli*, *K. pneumoniae*, and *Enterobacter cloacae* (Prasanth *et al.*, 2018). In addition, excellent results were obtained in terms of both safety and therapeutic efficiency in an Food and Drug Administration (FDA)-approved phase during the clinical trial (clinicaltrials.gov NCT00663091) on 42 patients with skin ulcers to evaluate the safety of 8 combinations of phages capable of lysing *S. aureus*, *P. aeruginosa*, and *E. coli* (Criscuolo *et al.*, 2017). Three years later, still in Spain, another study (clinicaltrials.gov NCT00937274) was conducted on the safety, tolerability, and efficiency of oral administration of phage T4 in the treatment of toxigenic *E. coli* diarrhea in children; the results have been shown to be an alternative to antibiotic therapy, especially in developing countries, without modification of the patients' intestinal flora

(Criscuolo *et al.*, 2017). Moreover, in a study carried out at the Ludwik Hirszfeld Institute in Poland on 157 patients (for 3 years) presenting various infections with multidrug-resistant bacteria, in particular urinary, bone, or respiratory, and treated with phage therapy for several weeks, beyond the observed efficiency, the biological parameters (blood count, liver function, renal function, and C reactive protein) demonstrated that no clinically relevant variation was observed, thus demonstrating the safety of bacteriophages under these conditions (Miedzybrodzki *et al.*, 2012). Similarly, in another study involving 15 people who took three oral doses per day (for 2 days) of the bacteriophage T4 targeting *E. coli*, in addition to the efficiency, no adverse effects or an increase in markers of hepatic cytolysis or the development of antibodies against the T4 bacteriophage has been observed (Bruttin and Brussow, 2005). Unfortunately, in recent years, some researchers have gradually evoked the resistance mechanisms of certain bacterial groups against the phages that once infected them (Jorge and Eduardo, 2019; Sadhana *et al.*, 2018).

Advantages and disadvantages of phage therapy

Table 1 presents the advantages and disadvantages of phages. One of the main drawbacks of phage therapy is the need to quickly establish the etiology of the bacteria causing the infection (Mireille *et al.*, 2020) due to the excessive specificity of many phages. However, this problem could be solved with the help of phage mixtures and selection of potent lytic phages from a collection prior to bacterial characterization which could form the basis of empirical combination therapy (Nobrega *et al.*, 2015). Otherwise, the possible involvement of humoral immunity with the production of neutralizing antibodies and the variable stability over time, influenced by storage conditions (pH, temperature, UV, etc.) as well as the potential capacity of phages to induce horizontal gene transfer by generalized transduction, is not negligible

(Jun, 2017). During cell lysis, bacterial endotoxins and other released substances are a specific disadvantage of phages and other antimicrobial agents (Jun, 2017). However, the zero impact on nontarget bacteria and eukaryotic cells, rapid bactericidal activity independent of antibiotic resistance (Bruttin and Brussow, 2005), and antibiofilm activity (Sadhana *et al.*, 2018) are certain assets that make this method a real alternative to conventional antibiotics.

Use of AMP as Alternatives to Antibiotics

AMP, also called host defense peptides, are essential components of the defense of multicellular beings and are part of the innate immune response (Andersson *et al.*, 2016). They are amphiphilic and cationic molecules, mostly 12–50 amino acids, which exhibit a wide variety of structures. They have a broad spectrum of antimicrobial action as well as antibiofilm, anti-inflammatory, and immunomodulatory properties and constitute a class of molecules of choice in the face of the problems of antibiotic resistance (Vasilchenko and Rogozhin, 2019).

Antibacterial properties of known AMPs

According to the antimicrobial peptide database (<http://aps.unmc.edu/AP/main.php>), 3,201 AMPs have been identified to date, of which 2,680 have antibacterial activity. Several AMPs have been shown to have a broad spectrum of activity against various microorganisms including Gram-positive and Gram-negative bacteria (Balter and Brown, 2011). In a recent study, the AMP PMAP-36PW and PMAP-36PK were found to have an expanded antibacterial spectrum, and in the evaluation of efficiency *in vivo* with mice infected with *Salmonella choleraesuis* C78-1 and *Listeria monocytogenes* CICC 21533, these peptide analogs have exhibited impressive therapeutic effect by reducing bacterial gene copies and decreasing inflammatory damage in mouse liver and lungs, resulting in reduction in mortality (Zhou *et al.*, 2019). Each year, many

Table 1. Summary of the advantages and disadvantages of the therapeutic use of bacteriophages from Mireille *et al.* (2020) with several modifications.

Characteristics	Advantage	Disadvantage
Narrow host spectrum	No impact on nontarget bacteria (microbiota).	For probabilistic approaches, this involves the use of bacteriophage cocktails to increase antibacterial coverage.
	Limitation of the number of strains likely to develop resistance to a given bacteriophage.	
Great diversity	High probability of isolating a bacteriophage infecting a pathogenic strain.	
Action mechanism	Rapid bactericidal activity.	Like antibiotics, release of endotoxins (and others) during lysis.
	Lytic activity independent of antibiotic resistance.	
	Antibiofilm activity of certain bacteriophages (capable of depolymerizing the polysaccharides composing biofilms).	
Viral nature (bacteria virus)	Proven diffusion in many tissues.	Possible involvement of humoral immunity with production of neutralizing antibodies.
	Inability to infect eukaryotic cells.	
	Replicative: increased concentration at the site of infection.	Variable stability over time, influenced by storage conditions (pH, temperature, UV, etc.).
	Found naturally in the environment, it is not necessary to manufacture them.	
	Rapid isolation of new bacteriophages possible (within a few hours or days) for many pathogens.	Potential ability to induce horizontal gene transfer by generalized transduction (virulence factor; antibiotic resistance).

similar studies are conducted and that provide baseline data for the design of clinically effective antibacterial peptides. In addition, unlike common antibiotics (Mahlapuu *et al.*, 2016), several AMPs such as melittin, LL37, and alamethicin have demonstrated their ability (*in vivo* and *in vitro*) to induce membrane permeability in spheroplasts in *E. coli* (Faust *et al.*, 2017). Furthermore, several antibacterial peptides act on other intracellular target processes, such as inhibition of DNA and RNA synthesis (buforin II, pleurocidin, and dermaseptin), protein synthesis (indolicidin and PR-39), and enzymatic activity (Jenssen *et al.*, 2006). As shown in Table 1, most antibacterial peptides are associated either with carbohydrates (glycoproteins) or with lipids + carbohydrates (lipoglycopeptides) and their action is mainly directed toward Gram-positive bacteria. Some antimicrobial peptide drugs approved by the Food and Drug Administration (FDA) are presented in Table 2 AMPs are generally active on resistant bacteria and can be used for the treatment of complex infections involving resistant bacteria (Lei *et al.*, 2019); such is the case with Telavancin and Teicoplanin which can be used either in the treatment of pseudomembranous colitis or treatment of diarrhea associated with *Clostridium difficile*; and the other in complicated skin infections, blood infections, endocarditis, bone and joint infections, and meningitis caused by methicillin-resistant *S. aureus* (Lei *et al.*, 2019). The same could be said for daptomycin which was marketed in 2003 as an anionic AMP for the treatment of skin infections caused by Gram-positive bacteria and has shown inhibitory effects on *Salmonellae* highly resistant to classical antibiotics and on *S. aureus* (Lei *et al.*, 2019).

Combinations of AMP–AMP and AMP–Antibiotics

AMPs have shown their effectiveness on biofilms. This is the case of gH625, its analog gH625-GCGKKKK, and the synergy of cyclic polymyxin B and gramicidin S AMPs which are active on *P. aeruginosa* biofilms (Rončević *et al.*, 2019). In the same sense, the combination of different substances, Geitani *et al.* (2019) demonstrated an interesting antibacterial activity of the AMPs–antibiotic association. Indeed, human cathelicidin (LL-37) and cecropin (1–7)–melittin A (2–9) amide in combination with the antibiotics colistin and imipenem has proven to be effective against antibiotic-sensitive or resistant *S. aureus* and *P. aeruginosa*. Moreover, their combination considerably reduces the minimum inhibitory concentration (MIC) of imipenem and colistin and reduces their toxicity (Geitani *et al.*, 2019).

AMP and brain infections

A highly desirable feature in the treatment of infectious agents is the ability of the antimicrobial to be able to cross the blood–brain barrier. To this end, several AMPs, such as oncocin, apidecin (137), and drosocin (Pro5Hyp), can cross the blood–brain barrier to selectively target brain cells. These AMPs could be potential therapies for brain infections (Li *et al.*, 2014). Concisely, considering all the advantages they present, AMP constitute a major alternative to fight against antibiotic resistance. However, although they are a serious option to replace antibiotics, they also have many drawbacks that limit their production and use. This is linked to their MIC which is often too high and can constitute a factor of toxicity (Raheem and Straus, 2019). Thus, it would be more judicious to use them: on resistant bacteria on which they are active; for infections where their MIC is not very high; in combination with antibiotics to overcome resistance and limit the use of conventional antibiotics.

Use of Vaccines to Reduce the Use of Antibiotics

Since the discovery of the first vaccine in the 1790s, many other vaccines have been developed and have since played a major role in the prevention of various diseases. Vaccines are defined as preparations based on microorganisms with attenuated virulence or on their constituents, which the administration aims to protect the host from the disease caused by the pathogen involved (Rappuoli *et al.*, 2014). The best-known antibacterial vaccines are those for diphtheria (caused by *Corynebacterium diphtheriae*), cholera (caused by *Vibrio cholerae*), tetanus (caused by *Clostridium tetani*), and pertussis (caused by *Bordetella pertussis*). In some countries, the oral Ty21a vaccine to prevent infections with *Salmonella typhi* (the causative agent of typhoid) is widely used. Certain vaccines can be combined, and this makes it possible to protect the host against several infectious agents at the same time; this is the case with the four-valent combination vaccines marketed under the names Infanrix Tetra®, Tetravac acellulaire®, Repevax®, and Boostrix Tetra®, which simultaneously prevent diphtheria, tetanus, pertussis, and poliomyelitis. The introduction of vaccines against resistant pathogenic bacteria could constitute a sufficient immunological barrier allowing the host to fight the incriminated bacteria without resorting to antibiotics (Terlizzi *et al.*, 2017). However, setting up a vaccine against a specific bacterial germ requires the presence of one or more factors likely to trigger an immune response (Terlizzi *et al.*, 2017). These factors typically include adhesins, iron receptors, secreted toxins, capsules (and their K antigens), capsule flagella (and their H antigens), outer membrane proteins, and lipopolysaccharides which may be transferred by plasmids, transposons, bacteriophages, and islets of pathogenicity (Terlizzi *et al.*, 2017). Numerous studies have shown that the use of vaccines in certain pathologies of bacterial origin can help reduce the use of antibiotics (Bloom *et al.*, 2018; Klugman and Black, 2018; Kobayashi *et al.*, 2016; Rappuoli *et al.*, 2014) and some have demonstrated the effectiveness of vaccines in the fight against antibiotic resistance (Klugman and Black, 2018; Ouldali *et al.*, 2019). The most recent example is the introduction of pneumococcal conjugate vaccines, which has led to a significant drop in the proportion of resistant pneumococcal strains. Unfortunately, due to the excessive consumption of antibiotics, some of the nonvaccine serotypes have also become resistant to antibiotics, raising (still moderately) the level of resistance (Ouldali *et al.*, 2019). In addition, a team of researchers from Johns Hopkins University have demonstrated the effectiveness *in vitro* (on macaques) of a vaccine against *S. aureus* (the major cause of nosocomial infections) whose treatment is generally complicated due to the resistance of this germ to antibiotics (Kuklin *et al.*, 2006). Ultimately, despite the difficulties presented by this alternative, it can constitute significant protective means against the bacterial strains for which vaccines exist and consequently limit the use of antibiotics in a possible infection that can be caused by these pathogens.

Use of Cytokines and Antibodies as an Alternative to Antibiotics

Cytokines are cell signaling glycoproteins, expressed by cells of the immune system, and antibodies are glycoproteins capable of specifically detecting and neutralizing pathogens (Larochette *et al.*, 2019). With the growth of antibiotic resistance, several researchers have proposed antimicrobial antibodies

and cytokines as possible alternatives to reduce the use of conventional antibiotics (Holland, 2000). The ability of antibodies and cytokines to fight against various bacteria *in vitro* (Bebbington and Yarranton, 2009; Conti *et al.*, 2000) and *in vivo* (Hancock *et al.*, 2012; Morita *et al.*, 2021; Rumfield *et al.*, 2020) has been demonstrated. Some studies have proposed antibody–antibiotic combinations to overcome antibiotic resistance and to cure serious bacterial infections (Song *et al.*, 2012; Sawa *et al.*, 2014). Furthermore, certain monoclonal antibodies such as pagibaximab (BSYX-A110) (chimeric monoclonal antibody directed against lipoteichoic acid, a key component of the wall of Gram+ bacteria) have been set up to overcome certain disadvantages of antibiotics in vulnerable people and fragile individuals such as premature babies (Hancock *et al.*, 2012). Finally, although antibodies cannot completely replace antibiotics, they can nevertheless help to reduce their use and fight against antibiotic resistance.

Phytochemicals and Plant Extracts as Alternatives to Antibiotics

Herbal medicine refers to the use of plants for therapeutic purposes. Depending on the case, the whole plant or parts of it are used, including the roots, stems, leaves, flowers, and fruits. To exploit their properties in various diseases, they generally proceed by extraction methods such as infusions, maceration, decoctions, or even extractions with solvents such as ethanol, methanol, and acetone. Plants with high antibacterial potential can contribute significantly to the diversification of antibacterials and fight against antibiotic resistance either by using them as they are or by using them in synergy with conventional antibiotics to thwart bacteria that are already resistant (Arsène *et al.*, 2021; Mbarga *et al.*, 2021; Nascimento *et al.*, 2000).

Antibacterial properties of plant extracts and phytochemicals

Between 2015 and 2019, around 11,000 studies were published on the antibacterial properties of plants across the world (Source: PubMed). These studies, whether *in vitro* or *in vivo*, have demonstrated the antibacterial properties of many plants on various bacteria (Nascimento *et al.*, 2000; Alrozeky and Nakahara, 2002; Arsène *et al.*, 2021; Mbarga *et al.*, 2021). Among these varied plants, certain spices and edible plants have also proved to be antibacterial (Alrozeky and Nakahara, 2002; Fankam *et al.*, 2011). In a study conducted in China in 2002, several edible Asian plants demonstrated antibacterial activity on *E. coli*, *Salmonella infantis*, *L. monocytogenes*, *S. aureus*, and *Bacillus cereus* (Alrozeky and Nakahara, 2002). In Cameroon, a study involving spices used in cooking showed that 15 of the 20 plants used had an antibacterial effect on *Mycobacterium tuberculosis* H37Rv and H37Ra and extract of *Echinops giganteus* exhibited the most significant activity with a MIC value of 32 and 16 µg/ml (Fankam *et al.*, 2011). The list of studies having obtained significant results with plant extracts cannot be exhaustive here but more detailed information on this subject can be found in our previous review (Mbarga *et al.*, 2021). However, it is necessary to emphasize that most of the studies carried out on the antibacterial properties of plant extracts were carried out *in vitro*. Such results cannot be transposed directly to a possible *in vivo* application without additional studies. Notwithstanding the above, given the use of certain plant extracts in traditional medicine as a treatment of infections of the urinary tract (Ibrahim *et al.*, 2015)

and diarrheal infections (Ullah *et al.*, 2016) among others, it clearly appears that plant extracts can play a major role in reducing the use of antibiotics.

Synergy between common antibiotics and phytochemical compounds to counteract antibiotic resistance

In most of the *in vitro* studies carried out on the antibacterial properties of plant extracts or phytochemical compounds, a part is reserved for the study of the effect of their combination with common antibiotics on multiresistant bacteria. This is, for example, the case with the study conducted by Nascimento *et al.* (2000) which demonstrated the synergistic effect of the combination of antibiotics with extracts of clove, jambolan, pomegranate, and thyme on *P. aeruginosa* (resistant to 19 antibiotics) and *K. pneumoniae*. Likewise, the combination of anacardic acid and totarol with methicillin has shown positive results in inhibiting methicillin-resistant strains of *S. aureus* (Nascimento *et al.*, 2000). The mechanism of action of this synergistic action would involve multidrug efflux pumps (EPs) located on the cytoplasmic membrane of bacteria. Indeed, plants are an important reservoir of bioactive compounds which can serve as a source of potent EP inhibitors (EPIs) (Shriram *et al.*, 2018). One study published in 2020 demonstrated that flavonoids could serve as EPIs and antimicrobials against both environmental and pathogenic intracellular mycobacterial species (Solnier *et al.*, 2020). In the same vein, the results obtained by Tran *et al.* (2020) demonstrated that carvotacetones from *Sphaeranthus africanus* could have an antibacterial activity and play an EPI role against Mycobacteria. Briefly, the antibacterial properties of several plant extracts and isolated phytochemical compounds are undeniable but more *in vivo* studies should be carried out to determine the optimal conditions under which these preparations could be safely used as such or in combination with conventional antibiotics.

Use of NPs in Synergy with Conventional Antibiotics

NPs are a wide class of materials that include particulate substances, which have one dimension less than 100 nm at least (Laurent *et al.*, 2010). NPs are considered a feasible alternative to antibiotics, especially with the emergence of bacterial multidrug resistance, since it has been discovered that they do not lead to bacterial resistance (Rai *et al.*, 2012). As shown in Table 3, several recent investigations have used different methods to synthesize various NPs and these NPs have shown very encouraging antimicrobial properties with very low minimum inhibitory and bactericidal concentrations. NPs have a high surface area to volume ratio and unique chemical and physical properties, which makes them promising antimicrobial agents (Crisan *et al.*, 2021; Yin *et al.*, 2020). It has been observed that NPs target multiple cellular pathways at once (Crisan *et al.*, 2021; Li *et al.*, 2012), which makes it extremely difficult for bacteria to develop resistance against them (Rai *et al.*, 2014). NPs with size less than 20 nm can kill bacteria by penetrating the cell wall and destruct cell organelles (Arakha *et al.*, 2015; Crisan *et al.*, 2021) and can hamper the synthesis of nucleic acids in several microorganisms (Fayaz *et al.*, 2010). In recent years, silver nanoparticles (AgNPs) were considered as a possible new class of antimicrobial (Arsène *et al.*, 2021; Crisan *et al.*, 2021; Rai *et al.*, 2014). Several studies have shown that AgNPs can enhance the effect of antibiotics

Table 2. Some antimicrobial peptide drugs approved by the FDA.

Drug	Type	Against	Administration	Clinical use	References
Bacitracin	Cyclic peptides	Gram +	Topical	Localized skin and eye infections and wound infections	Lei <i>et al.</i> (2019)
Dalbavancin	Lipoglycopeptide	Gram +	Intravenous	Complicated skin infections	Spann <i>et al.</i> (2004)
Daptomycin	Lipoglycopeptide	Gram +	Intravenous	Skin and skin structure infections caused by Gram-positive infections, <i>S. aureus</i> bacteremia, and right-sided <i>S. aureus</i> endocarditis	Lei <i>et al.</i> (2019)
Oritavancin	Glycopeptide	Gram +	Intravenous	Acute bacterial skin and skin structure infections caused by Gram-positive bacteria	Lei <i>et al.</i> (2019)
Teicoplanin	Glycopeptide	Gram +	Intravenous and intramuscular	Bacterial infections and in the treatment of pseudomembranous colitis and <i>C. difficile</i> -associated diarrhea, with comparable efficiency with vancomycin	Lei <i>et al.</i> (2019)
Telavancin	Lipoglycopeptide	Gram +	Intravenous	Complicated skin and skin structure infections	Higgins <i>et al.</i> (2005)
Vancomycin	Glycopeptide	Gram +	Oral and intravenous	Complicated skin infections, bloodstream infections, endocarditis, bone and joint infections, and meningitis caused by methicillin-resistant <i>S. aureus</i>	Hamilton, (2015)

against susceptible and resistant bacteria (Birla *et al.*, 2009). Combinations between antibiotics and NPs have also shown efficiency against multiresistant bacteria. Indeed, in a study conducted by Mala *et al.* (2016) where they impregnated urinary catheters with a synergistic combination of antibiotics and AgNPs to evaluate antibiofilm activity *in vitro* and *in vivo*, the authors reported that the synergistic combination showed a 90% inhibition of bacterial adhesion and concluded that a synergistic combination of antibiotics and AgNPs is an efficient method for preventing biofilm formation (Mala *et al.*, 2016). Besides AgNPs, copper

nanoparticles (CuNPs) also demonstrated high antibacterial and antifungal effects (Amer and Awwad, 2021; Ramzan *et al.*, 2021). Examples showing the antibacterial activity of other NPs are presented in Table 3. Although NPs have proven to be a great substitute for conventional antibiotics, they can be used not only to treat antibiotic-resistant bacterial infections but also to avoid the potential for resistance development. However, research should no longer stop at the *in vitro* aspect but should propose formulations likely to be applied to treat bacterial infections.

Table 3. Some recent research on the green synthesis of NPs and their antibacterial activity.

NPs	Abbreviation	Plant material used	Antibacterial activity against	Sources
Silver	AgNPs	Grapefruit peel	Gram-positive (<i>S. aureus</i> and <i>E. faecalis</i>) and Gram-negative bacteria (<i>E. coli</i>)	Arsène <i>et al.</i> (2021)
		<i>Nymphaea odorata</i>	<i>Staphylococcus aureus</i> and <i>E. coli</i> at very low concentration (25µg/ml)	Gudimalla <i>et al.</i> (2021)
		<i>Symplocos racemosa</i>	<i>Pseudomonas aeruginosa</i> .	Panda <i>et al.</i> (2021)
		<i>Citrus limon</i> fruits	<i>Staphylococcus aureus</i> and <i>E. coli</i> (with high inhibition diameter)	Amer and Awwad, (2021)
Copper and copper oxide	CuNPs and CuONPs	<i>Cedrus deodara</i> leaf	<i>Escherichia coli</i> and <i>S. aureus</i> with the highest antibacterial activity on <i>E. coli</i>	Ramzan <i>et al.</i> (2021)
		<i>Cassia fistula</i> and <i>Melia azedarach</i>	Multidrug-resistant <i>K. pneumonia</i> and <i>Helicobacter pylori</i> biofilms	Naseer <i>et al.</i> (2021)
Zinc and zinc oxide	ZnNPs and ZnO-NPs	<i>Justicia adhatoda</i>	<i>Staphylococcus aureus</i> , <i>E. coli</i> , and <i>P. aeruginosa</i>	Pachaiappan <i>et al.</i> (2021)
Gold	AuNPs	<i>Garcinia kola</i>	<i>Bacillus cereus</i> , <i>S. aureus</i> , <i>E. coli</i> , and <i>P. aeruginosa</i> with the highest activity on <i>P. aeruginosa</i>	Akintelu <i>et al.</i> (2021)
Nickel and nickel oxide	NiNPs and NiO-NPs	Onion extract		Nawaz <i>et al.</i> (2020)
Iron and iron oxide	FeNPs and FeO-NPs	<i>Carica papaya</i> leaf	Moderate antibacterial activity against <i>Klebsiella</i> spp., <i>E. coli</i> , <i>Pseudomonas</i> spp., and <i>S. aureus</i>	Bhuiyan <i>et al.</i> (2020)
Magnesium and magnesium oxide	MgNPs and MgO-NPs	<i>Dalbergia sissoo</i> leaf	High antibacterial activity against <i>E. coli</i> and moderate against <i>Ralstonia solanacearum</i>	Khan <i>et al.</i> (2020)
Manganese and manganese oxide	MnNPs and MnO ₂ -NPs	<i>Aloe vera</i>	<i>Streptococcus mutans</i> , <i>S. aureus</i> , and <i>E. coli</i> ,	Joshi <i>et al.</i> (2020)
Selenium	SnNPs	<i>Ceropegia bulbosa</i> Roxb	<i>Bacillus subtilis</i> and <i>E. coli</i>	Cittrarasu <i>et al.</i> (2021)

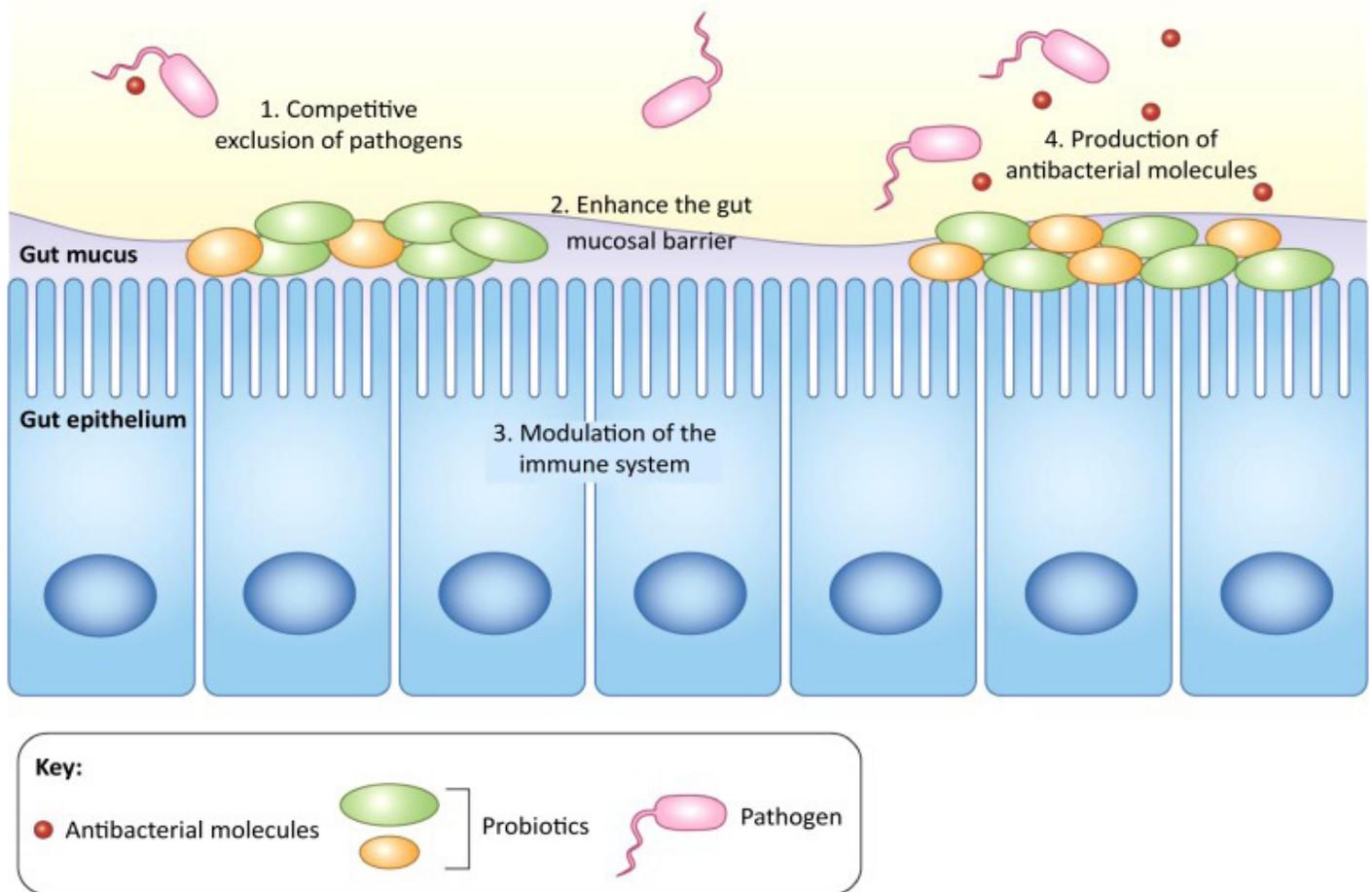


Figure 2. Mechanism of protection of probiotics against infections (Ghosh *et al.*, 2019).

Probiotics as a Potential New Antibacterial Strategy

Probiotics are living microorganisms (bacteria or yeasts), which when consumed in adequate quantities produce a beneficial effect on the health of the host beyond the traditional nutritional effects (WHO and FAO, 2002). The main strains recognized as probiotics in humans are bacteria belonging to the genera *Lactobacillus*, *Bifidobacterium*, *Enterococcus*, and *Streptococcus* and yeasts of the genus *Saccharomyces* (Ayivi *et al.*, 2020; Joseph *et al.*, 2021). As described in Figure 2 (Ghosh *et al.*, 2019; Markowiak and Śliżewska, 2017), it has been shown that certain probiotics are able to inhibit the growth of pathogenic bacteria by specific and nonspecific competition for adhesion to epithelial cells (Bajaj *et al.*, 2021), by the production of antimicrobial substances (Arsene *et al.*, 2021; Silva *et al.*, 2020), by competition in the use of nutrients (Abd El-Hack *et al.*, 2020), and by the stimulation of immune defense mechanisms (Abd El-Hack *et al.*, 2020). With the resurgence of the resistance to antibiotics issue, the antagonist and immunomodulatory properties of probiotics are today the most used (Arsene *et al.*, 2021; Manga *et al.*, 2019; Silva *et al.*, 2020). Thanks to these properties, probiotics could constitute a credible alternative to antibiotics in the fight against specific bacterial infections for which their effectiveness has already been

proven. Our previous review comprehensively presented the main probiotics that can effectively replace antibiotics in breeding (Arsène *et al.*, 2021). Otherwise, several *in vitro* and *in vivo* studies, both in clinical medicine and in veterinary medicine have demonstrated the efficiency of probiotics against specific germs (Abd El-Hack *et al.*, 2020; Bajaj *et al.*, 2021; Huang *et al.*, 2014; Homan and Orel 2015; Makras, 2006). For example, the study conducted by Makras (2006) demonstrated the antagonist activity of six strains of *Lactobacilli* (*L. acidophilus*, *L. amylovorus*, *L. casei*, *L. johnsonii*, *L. plantarum*, and *L. rhamnosus*) on diarrhea caused by *Salmonella enterica*. In addition, certain combinations of probiotics with each other or with antibiotics have also demonstrated their effectiveness (Homan and Orel, 2015). In the search for new antibacterial control strategies and the reduction of the use of antibiotics, probiotics have various advantages such as the possibility of being included in food such as yogurts, cheeses, dietary supplements, and fermented drinks (Mbarga *et al.*, 2019), their immunomodulatory capacity (Silva *et al.*, 2020), and their ability to cleanse and balance the intestinal flora (Wang *et al.*, 2020). However, although they have positive effects on certain enteric infectious pathologies, probiotics have various drawbacks such as their ineffectiveness on major bacterial infections affecting other organs such as the skin, heart, or brain.

CONCLUSION

All the methods discussed in this review can help to reduce the use of antibiotics, but none alone can replace them. Our analysis of the situation shows that the development of vaccines can effectively contribute to reducing the use of antibiotics by strengthening the immune system in order to protect it against specific pathogens as is already the case with bacteria such as *Neisseria meningitidis*, *Haemophilus influenzae*, *Streptococcus pneumoniae*, *M. tuberculosis*, *C. diphtheriae*, and *C. tetani*. In addition, probiotics, thanks to their multiple barrier effects on the intestinal flora, their antagonist effect, and their immunomodulation capacity of the immune system, can be a significant contribution in reducing the use of antibiotics. As for phages, cytokines, and antimicrobial antibodies, given their narrow specificity, they appear to be pragmatic solutions to drastically limit the use of antibiotics in the treatment of diseases caused by bacteria on which their effect is recognized. However, phages have some drawbacks such as the release of enterotoxin and other metabolites during lysis, the possible involvement of humoral immunity with the production of neutralizing antibodies, their variable stability over time influenced by storage conditions (pH, temperature, UV, etc.), and their potential capacity to induce horizontal transfer of genes by generalized transduction (virulence factor; resistance to antibiotics). Although the antibacterial efficiency of several plants has been demonstrated, there is an eternal dilemma as to the isolation of active molecules because the totum of preparations based on these plants would be more bioavailable and more effective than isolated phytochemicals and their potential ability to induce horizontal gene transfer by generalized transduction (virulence factor; antibiotic resistance). In addition, the green synthesis of NPs (silver, gold, zinc, copper, iron, etc.) is also an important option against the scourge of antibioresistance and the combination of common antibiotics with other antibacterial molecules (phytochemicals, NPs, and AMP) can give a second life or improve the effectiveness of certain antibiotics which are losing their effectiveness. Finally, research should continue to be carried out to effectively exploit all the alternatives presented in this study to face this major public health issue of antibiotic resistance.

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All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; agreed to submit to the current journal; gave final approval of the version to be published; and agree to be accountable for all aspects of the work. All the authors are eligible to be an author as per the international committee of medical journal editors (ICMJE) requirements/guidelines.

CONFLICTS OF INTEREST

The authors report no financial or any other conflicts of interest in this work.

ETHICAL APPROVALS

Not Applicable

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