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REVIEW ARTICLE

Antibiotic Resistance and Emerging Alternatives for Controlling Foodborne Pathogens

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Antibiotic resistance in foodborne pathogens poses a significant challenge to public health and food safety. Major foodborne pathogens, such as Salmonella, Campylobacter, Escherichia coli, and Listeria monocytogenes, are commonly linked to contaminated meat, poultry, dairy products, and fresh produce, and their resistance to antibiotics is a growing concern. This resistance arises through genetic mechanisms like mutations and horizontal gene transfer, facilitated by resistance factors such as efflux pumps and biofilm formation. Geographical and temporal trends reveal disparities in resistance levels, with higher prevalence in low- and middle-income countries due to unregulated antibiotic use. Both agricultural and clinical antibiotic applications contribute to resistance, necessitating urgent intervention. Alternatives to antibiotics offer promising strategies to control foodborne pathogens. Bacteriophages, which target specific bacteria, are already applied in food safety but face limitations like phage resistance. Probiotics and prebiotics enhance gut health and inhibit pathogens, with effective combinations demonstrating synergistic effects. Antimicrobial peptides (AMPs) from natural and synthetic sources offer a broad spectrum of activity, while essential oils and plant extracts provide natural antimicrobial solutions in food preservation. Nano-based interventions, such as silver and chitosan nanoparticles, show potential but raise concerns about toxicity and environmental impacts. Vaccination strategies targeting livestock and humans offer a proactive approach but face challenges in development and implementation. CRISPR/CAS systems enable precise gene editing to eliminate resistance genes, while emerging solutions like lysozymes and competitive exclusion products further expand the toolkit. Applications of these alternatives span food processing, preservation, and animal agriculture, significantly reducing pathogen prevalence. Regulatory and policy frameworks must support their integration into food systems. Global collaborations and innovations are crucial for combating antibiotic resistance effectively, thereby ensuring safer food production and enhanced public health protection.

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INTRODUCTION

Foodborne pathogens include foodborne bacteria, viruses, and parasites that are capable of causing disease in humans. Such organisms are a global public health concern since they cause a significant level of morbidity and mortality as well as economic loss (Akinsemolu & Onyeaka, 2024). According to the World Health Organization (WHO), every year around 600 million people get sick due to foodborne diseases, resulting in around 420,000 deaths annually (Oduoye *et al.*, 2024). These foodborne pathogens can be transmitted to people

through the ingestion of infected meat, milk, or eggs, and contamination from the environment, such as water runoff from farms (Gobena *et al.*, 2024; Yitbarek, 2024). These infections cut across all social groups but are most rife among children, pregnant women, the elderly, and immune-compromised individuals (Mfeka, 2021). Antibiotics have played a significant role in the control of foodborne illnesses caused by bacterial pathogens since the mid-20th century (Lawani-Luwaji, 2024). These drugs work by inhibiting or killing bacteria within the body, making the drugs effective against infections and alleviating the symptoms (Karunarathna *et al.*, 2024). Antibiotics are widely used in both clinical and agricultural settings. Clinically, they are used to manage intensive bacterial infections, including those caused by foodborne pathogens, which include Salmonella, E. coli Campylobacter, etc. In agriculture, antibiotics are also used to treat diseases in livestock, as a preventive measure as well as for growth stimulation. The use of antibiotics for agricultural purposes is one of the major factors contributing to antibiotic resistance and dissemination patterns (Yaqub et al., 2024). Antibiotics are used not only to treat illnesses but also in healthy animals where they are used as prophylactic drugs or growth enhancers, and their usage is justified as they improve production. The nontherapeutic use of antibiotics fosters and selects for the survival and expansion of resistant bacteria.

Antibiotic resistance (AR) occurs when bacteria develop mechanisms that allow them to survive and proliferate despite the presence that would normally inhibit or kill them (Bai et al., 2024). While there are serious illnesses for which antibiotics are irreplaceable, their extended and misused application in practice and agriculture has triggered a surge of AR. To address this challenge, a balanced and integrated strategy that optimizes the available treatment options and at the same time seeks to protect the antibiotic efficacy for the future is mandatory (Ahmed et al., 2024; Legrand et al., 2024). Antibiotic resistance in foodborne bacterial pathogens is not only widespread but also one of the most significant global health challenges, joined with climate change further complicates the control of infectious diseases, including those caused by foodborne pathogens. Shifts in temperature, extreme weather, and humidity can alter the survival, distribution, and transmission pattern of pathogenic microbes, potentially increase the risk of outbreaks and reducing the predictability of disease management strategies.

People can suffer severe and sometimes lifethreatening infections due to pathogens that most of antibiotics cannot help in treating the diseases (Lewies et al., 2015). Foodborne pathogens such as Salmonella spp., E. coli, Campylobacter spp., and Listeria monocytogenes have constantly developed resistance to numerous antibiotics, including those that are considered to be of utmost importance in modern medicine (Grudlewska-Buda et al., 2023). Strains of these pathogens that are multidrugresistant (MDR) have been isolated from foodstuffs, from agricultural and clinical settings, making treatment guidelines overly complex and leading to catastrophic health consequences (Alara & Alara, 2024). One of the chief determinants of AR in foodborne bacteria is the misuse of antibiotics in food-producing animals. The globalized nature of the supply chain of food has made it easy for antibiotic-resistant bacteria to be transmitted across nations, making AR a global phenomenon (Tang et al., 2017; Sagar et al., 2023).

When faced with the problem of antibiotic resistance, it is necessary to take multidisciplinary and cooperative action. Together, public health agencies, scientists, and politicians can strengthen surveillance systems, improve antibiotic practices, and come up with alternative means for pathogen eradication (Muteeb *et al.*, 2023). At the same time, awareness campaigns are useful to promote the most appropriate behavior of consumers and providers of healthcare services (Eruaga, 2024). The increased use of antibiotics has generated concern regarding the emergence of antibiotic-resistant pathogens, which necessitate the use of bacteriophages. These are viruses that invade bacterial cells, and bacteriophages have shown promise in the targeted approach to foodborne pathogens. Advisory bodies like the Food and Drug Administration (FDA) have approved the use of bacteriophage-based interventions (Qiao *et al.*, 2024).

Fermented products: containing essential oils or phenolic compounds have also been shown to possess some antimicrobial activity against foodborne pathogens. There has been an increased interest in these natural antimicrobials as additives in food preservatives and coatings for fresh-cut produce (Di Matteo et al., 2024). Today, high-pressure or UV light treatment has been developed as a safe and effective method to reduce bacterial contamination. Other crucial aspects in the control of foodborne pathogens include proper biosecurity measures and hygiene practices in both agriculture and the food industry (Ndraha et al., 2024). This review aims to highlight the major foodborne pathogens, their mechanisms of action, antimicrobial resistance, and alternatives to antibiotics for controlling foodborne pathogens.

Mechanism of antimicrobial resistance: This mechanism includes the production of enzymes like β -lactamases that degrade antibiotics, modification of antibiotic target sites (e.g., DNA Gyrase or mutation in ribosomal proteins), and activation of efflux pumps that expel antibiotics from the cell. Additionally, changes in membrane permeability reduce antibiotic uptake, and biofilm formation provides a physical barrier protecting bacteria from antimicrobial agents. Many of these resistance traits are carried on mobile genetic elements such as plasmids, transposons, and integrons, enabling rapid horizontal gene transfer within and between bacterial species (Fig. 1) (Table 1). The widespread use of antibiotics in food animals for growth promotion and disease prevention has accelerated the emergence and dissemination of these resistance mechanisms, posing a significant threat to food safety and public health.

Major foodborne pathogens: Foodborne pathogens pose major public health concerns worldwide and lead to millions of illnesses each year. These include bacteria,



Fig. 1: Antibiotic efflux pump and enzymatic inactivation (Retrieved from biorender) (Kim & Ahn, 2022).

Table I: Examples of	significant mechanisms of antimicrobial resistance to	ogether with the correspondi	ng bacteria and antimicro	bials Defense
Antibiotic resistance mechanism	Classical examples	Antibiotics affected	Examples of Bacteria	References
Antibiotic	ß lactamases	Penicillin	S aureus	(Vilvanathan, 2021)
destruction	Penicillinases	narrow spectrum	N. Gonorrhoeae	(**************************************
		cephalosporins	Haemophilus	
			Influenzae	
			Enterobacteriaceae	
	Extended- β	Penicillin, first-, second,	Enterobacteriaceae	(Mizrahi et <i>al.</i> , 2020)
	Spectrum	third-generation	Pseudomonas	
	lactamases	cephalosporins and	aeruginosa	
	A	aztreonam Deniaillin finat accord	Fatanah antariana ar	(D
	AmpC enzymes	third		(Doern, 2021)
		generation cephalosporins	Pseudomonas spp.	
		aztreonam, and cephanosponing,	i seudomonus spp.	
	Carbapenemases	Carbapenems and almost	Enterobacteriaceae	(Cooper et al., 2021)
	•	Hydrolysable β-lactams		· · · /
Antibiotic modification	Aminoglycoside	Aminoglycosides	numerous bacteria	(Thacharodi &
	Modifying Enzymes (AMEs)			Lamont, 2022)
	Chloramphenicol	Chloramphenicol	numerous bacteria	(Biswas et al., 2012)
	acetyltransferases (CATs)			
Modifications of	Mutations in the nitroreductase genes nfsA and	Nitrofurantoin	Enterobacteriaceae	(Christaki et al.,
antibiotic-activating	nfsB			2020)
enzymes Target	Panicillin Pinding		Strabtococcus	(Eani at al. 2012)
rangel	Protoin (PBP) replacement	p lactams	Streptococcus	(Fani el ul., 2012)
bypass	Acquisition of a novel Penicillin-Binding Protein	ßlactams	S aureus	(Wacnik et al. 2022)
bypass	(PRP)	plactariis	J. durcus	
	Replacement of the terminal D-Alanine D-Alanine	Glycopeptides	Enterococci	(Wacnik et al., 2022)
	Moiety of the peptidoglycan precursors	- / - F - F	S. aureus (rarely)	(,, . ,
	Acquisition of a novel dihydrofolate reductase	Trimethoprim	Gram-negative	(Faltyn et <i>al.,</i> 2019)
			bacteria	
			Staphylococci	
			L. monocytogenes	
	Acquisition of a novel dihydropteroate synthase	Sulfonamides	Gram-negative bacteria	(Gauba & Rahman,
		-	F	2023)
	Utilization of folic	I rimethoprim	Enterococci	(Fernandez-Villa et al.,
Target site	exogenous acid	Linezolid	Entorococci S aurous	$(X_{\text{ang of } d}, 202E)$
alteration	Flutations of the 2551 KINA	Macrolides	numerous bacteria	(Tallg et ul., 2023) (Petinaki &
(mutation or		Lincosamides	numerous bacteria	Papagiannitsis, 2019)
enzymatic		Streptogramin B		·
alteration)	Mutations in the bacterial Gyrase or	Quinolones	numerous bacteria	(Badshah & Ullah,
	Topoisomerase IV			2018)
	Mutations in the RNA Polymerase β subunit	Rifampicin	numerous bacteria	(Ning et al., 2021)
	Mutational or Recombinational changes in the	Trimethoprim	numerous bacteria	(Sánchez-Osuna et
	Dihydrofolate Reductase gene	• •• • •		al., 2020)
	Mutational or recombinational changes in the	Sulfonamides	numerous bacteria	(Capasso & Supuran,
	dinydropteroate synthase gene	Macrolidas Lincocomidas	numerous basteria	2019) (Cattoir & Loclored
	Theory action of the 2551 KINA by erringenes	Streptogramin B	numerous Dacteria	(Catton & Leciercq, 2017)
	Methylation of the 23SrRNA by the cfr gene	Linezolid	S aureus	(Yang et al. 2025)
		Chloramphenicol	Stabhylococci	(Vester, 2018)
		Clindamycin		(*******
Target site protection	Ribosomal Protection Proteins (RPPs)	Tetracyclines	numerous bacteria	(Yadav et <i>al.,</i> 2021)
	Qnr proteins	Quinolones	numerous bacteria	(Ruiz, 2019)
Target	Overproduction of Dihydrofolate Reductase	Trimethoprim	E. coli	(Yu, 2024)
overproduction				
Decreased	Mutations affecting the expression or function of	Hydrophilic antibiotics (e.g.,	Gram-negative bacteria	(Gauba & Rahman,
permeability of the	porins	β lactams, quinolones,		2023)
bacterial outer		tetracyclines,		
Antibiotic offlux	Multidrug offlux pumps	A wide range of antibiotics	numorous bactoria	(Blanco et al. 2014)
Anubiouc eniux	Substrate-specific	A wide failge of antibiotics	numerous bacteria	(Henderson et al
	efflux numps	l'ett acyclines	numerous bacteria	2021)
		Macrolides	Streptococci and other	(Lekshmi et al., 2018)
			gram-positive organisms	()
		Chloramphenicol	numerous bacteria	(Seukep et al., 2022)
Global cell adaptations	Mutations in genes implicated in phospholipid	Daptomycin	Enterococci	(Johnston, 2021)
	metabolism and those encoding regulatory			•
	mechanisms governing cell envelope homeostasis			
	Gene mutations affecting important cell membrane	Daptomycin	S. aureus	(Poshvina et al., 2023)
	processes and cell wall alterations	Income disconsistent de la	C	(Destation of 20022)
	Cell wall thickening is caused by mutations in genes	intermediate susceptibility to	s. aureus	(rosnvina et al., 2023)
	regulate cell envelope homeostasis	8ixcopeptides		
	· · · · · · · · · · · · · · · · · · ·			

Та	able	: :	Examp	les o	f signi	ficant	mecha	anisms c	of antir	nicrol	bial	resistance	togethe	er with	the	e correspon	ding	bacteria and	l antimicro	bials	
_																					_

viruses, parasites, and prions that, at one point or another during the production, processing, and preparation chain, contaminate foods (Pal & Ayele, 2020). Major foodborne pathogens such as Salmonella, E. coli, L. monocytogenes, Campylobacter, Clostridium botulinum, and Norovirus have been responsible for severe diseases, with some of them being fatal without timely intervention (Afreen & Bağdatlı, 2021). Salmonella is the major cause of bacterial foodborne illnesses worldwide. This pathogen is highly common in raw or undercooked poultry, eggs, and unpasteurized milk. Salmonellosis is caused by the ingestion of contaminated food and is manifested by diarrhea, abdominal cramps, fever, and vomiting. While the illness is usually self-limiting, severe infections may be present in immunocompromised individuals, children, and the elderly. Good hygiene practices and proper cooking of food are some of the necessary preventive measures (Ehuwa et al., 2021).

Another major foodborne pathogen is *E. coli*, particularly the Shiga toxin-producing strain, O157:H7; this is normally associated with undercooked ground beef and raw vegetables, and also with contaminated water (Singha *et al.*, 2023). Symptoms also include severe diarrhea, with severe abdominal pain and vomiting, leading to serious complications such as hemolytic uremic syndrome. Kidney failures caused by Hemolytic Uremic Syndrome (HUS) may be manifested, which can affect children and older adults. Preventing *E. coli* contamination requires proper cooking to avoid cross-contamination by thoroughly washing produce (Possas & Pérez-Rodríguez, 2023).

L. monocytogenes is a kind of bacteria that can be particularly dangerous to pregnant women, newborn babies, and older adults. It can be found in refrigerated, ready-to-eat foods such as deli meats, soft cheeses, and smoked seafood. Infection with Listeria causes listeriosis, mostly leading to serious complications like meningitis, pregnancy complications, sepsis, and including miscarriage or stillbirth. Refrigeration does not kill this bacterium, so strict adherence to safe food storage and handling is very important (Wang et al., 2021a). Campylobacter is a major cause of bacterial diarrhea worldwide and is commonly associated with raw or undercooked poultry, unpasteurized milk, and contaminated water (Rehman, 2022). The infection generally presents with diarrhea, abdominal pain, fever, and nausea. Infection is usually self-limiting, but complications include Guillain-Barré syndrome, a rare neurological condition. Proper cooking of poultry and prevention of cross-contamination during food preparation are important modes of prevention (Akinsemolu & Onveaka, 2024).

C. botulinum produces a neurotoxin that is responsible for a rare but potentially lethal disease called botulism. It mostly occurs due to improperly canned or preserved foods. Symptoms include blurred vision, difficulty swallowing, muscle weakness, and, in severe cases, respiratory failure. Good food preservation practices and immediate disposal of damaged cans are very important safety considerations (Poulain & Popoff, 2019). Other leading contributors to foodborne illness are viruses. Norovirus is very contagious, acquired via food, water, or by touching any surface contaminated by an infected person. vomiting, diarrhea, nausea, and stomach cramps are the symptoms of Norovirus. It leads to outbreaks in group environments such as schools and cruise ships. Prevention includes good hand hygiene and cleaning of surfaces (Harris & Dabritz, 2024).

Other notable pathogens include Staphylococcus aureus, which elaborates heat-stable enterotoxins that can cause the rapid onset of food poisoning symptoms (Bao & Wu, 2024), and C. perfringens, whose outbreaks are commonly associated with improperly stored food (García et al., 2019). Bacillus cereus is another cause of two notable types of illness: one causing diarrhea and the other causing emesis: these have been associated with rice dishes and improperly cooked foods (Rodrigo et al., 2021). Among others, Toxoplasma gondii is a protozoan parasite found in undercooked meat that creates significant risks to pregnant women and people with immunocompromised conditions (Ducrocq et al., 2021). Mild flu-like signs to severe neurological complications are included in the symptoms. Cryptosporidium and Giardia lamblia are transmitted through contaminated water and thus are also among the significant contributors to foodborne parasitic infections, causing protracted diarrhea (Rossi et al., 2024).

Alternatives to antibiotics

Small Molecules (SMs): With drug-like qualities, SMs are non-peptide chemical compounds that may interact with biological molecules, such as proteins and nucleic acids, and change how they normally operate. Their strong hydrophilicity and low molecular weight (~200-500 Da) enable efficient absorption through both host and pathogen barriers. SMs can be altered to improve their mass applicability and antibacterial effectiveness. The creation of antibacterial medications and the discovery of SM candidates that prevent bacterial growth in whole-cell tests or the action of a primary bacterial enzyme or protein are two typical uses for high-throughput screening (HTS) (Alcalde-Rico et al., 2016). Targeting bacterial membranes, SMs have a minimal possibility for pathogens to acquire resistance, enhance the effectiveness of several antibiotics, and work well against biofilms and slowgrowing bacteria. Plant pathogens such as Xanthomonas species, Erwinia tracheiphila, Acidovorax citrulli, and Salmonella infections have recently been treated using SMs (Deblais et al., 2019; Kathayat et al., 2020). The activity of SMs within the recipient's body, the possibility of binding to non-target molecules within the human body, and structural design limits are some of their functional limitations, despite their positive impacts. These drawbacks include the inability to precisely identify the modifying proteins that SMs should target, the difficulty of engineering SMs to target unstructured disordered polypeptides, and their poor affinity to bind or alter protein surface mediators (Gurevich & Gurevich, 2014).

Quorum-sensing/Antivirulence inhibitors: Bacterial cells use a technique known as quorum sensing (QS) to communicate with one another and adjust their density and activity in response to their surroundings (Gao *et al.*, 2023). Bacteria secrete extracellular signaling molecules known as autoinducers (AIs), which play a crucial role in regulating the production of virulence factors, enzymes, biofilm formation, stress adaptation, secondary metabolite

synthesis, and swarming motility. AIs are released into the environment by active transport or diffusion to facilitate effective bacterial cell-to-cell communication. AIs accumulate in the environment as the density of bacteria increases, and once this exceeds a specific threshold, bacteria employ them as extracellular signaling molecules to coordinate their gene expression and modify their density. Three basic ideas underlie QS systems: (1) the density of bacterial cells; (2) receptors produced in the cytoplasm or on the membrane; and (3) the recognition of Als not only stimulates the expression of genes necessary for cooperative behaviors but also increases the bacterial synthesis of AIs. Bacterial QS systems are generally divided into three categories: (3) oligopeptide-twocomponent-type QS, (2) LuxI/LuxR-type QS, and (3) luxSencoded autoinducer-2 (AI-2) QS (Dutta et al., 2021). The AIs are divided into three categories: furanosyl borate diester, which is used by both Gram-positive and Gramnegative bacteria, oligopeptides, and acylated homoserine lactones (AHLs), which are used by Gram-negative bacteria (Koley et al., 2023).

Bacterial biofilm development and pathogenicity are decreased when the connecting mechanism between bacterial cells is disrupted. As a result, several tactics have been created to break this link and manage bacterial infections that are dependent on QS. Low-mass substances with excellent selectivity for the QS regulator and no harmful side effects on the bacteria or a possible eukaryotic host are known as ideal QS inhibitors (QSIs). They also need to be very effective and chemically stable. OSIs work through a variety of mechanisms, including (1) inhibiting AI synthesis; (2) antagonistic interactions with AI receptors; (3) blocking targets downstream of receptor binding; (4) using antibodies to sequester AIs; (5) breaking down AI-catalytic antibodies (abzymes) or enzymes (such as lactonases, acylase, and oxidoreductase); (6) attenuating AI secretion/transport; and (7) competing with autoinducing signal molecules to bind to the transcriptional protein regulators of bacterial QS systems.

One of the key limitations of QSIs is the potential development of resistance, as bacteria possess multiple OS systems and may adapt mechanisms to bypass or resist QSI activity. . By acquiring mutations in the amino acid residue of the LuxR protein regulator, which codes for virulence factors, motility, biofilm formation, and the manufacture of antibiotics, gram-negative bacteria can avoid the effects of QSIs. (1) Indole signaling: In a nutrient-poor environment, E. coli and Salmonella's QS systems react to indole by producing virulence factors, plasmid stability, adaptability, and antibiotic resistance (Escobar-Muciño et al., 2022). Under these circumstances, indole and AIs compete for binding to the SdiA transcriptional regulator's AHL domain. (2) Disturbance of microbiota homeostasis: OSIs disrupt human microbiota homeostasis by influencing human microflora activities such as adhesion, biofilm formation, and the synthesis of antimicrobial metabolites, as well as by causing the dispersion of AI-2 signaling (Juszczuk-Kubiak, 2024).

Probiotics: When given in sufficient quantities, probiotic microorganisms that coexist in a symbiotic relationship with the host perform a number of biological tasks and offer health advantages (Ahire *et al.*, 2024). They fall into

four groups: next-generation probiotics, viable and active viable/non-active probiotics. probiotics, and dead/nonviable probiotics (Silva et al., 2020). Few studies have examined probiotics' antibacterial qualities as innovative antibiotic substitutes, despite a large body of research on their biological qualities. Viable and active probiotics benefit the host's health by increasing the concentration of hydrogen ions in the gut (Liu et al., 2025), improving the synthesis of vital vitamins and enzymes, generating antimicrobial compounds, restoring intestinal microbiota following diarrhea, lowering serum cholesterol, strengthening the immune system, generating antioxidants, lowering food allergy sensitivities, and improving lactose and calcium absorption. Saliva enzymes, gastric fluid (acid and enzymes), bile salts, competitive gut bacteria, and inhibiting Gastrointestinal tract (GIT) conditions should all be avoided by them as they are moving from the mouth to the gut (Han et al., 2021).

Postbiotics and parabiotics, often known as dead or nonviable probiotics, are viable probiotics that lack metabolic activity (Nataraj et al., 2020). They are subjected to harmful stressors such as excessive O₂ supplementation, high osmotic pressure, temperature, and severe pH levels (Bustos et al., 2025). As inactive viable probiotics, Bacillus species, including B. coagulans, B. subtilis, B. clausii, and B. licheniformis are utilized in human diets and to treat intestinal and urinary issues (Sadrimovahed & Ulusoy, 2024). Because Bacillus species generate bacteriocins that are efficient against both Gram-positive and Gramnegative bacteria and fungi present in food, they are frequently utilized as natural preservatives in the food Dead/nonviable probiotics business. can use dead/nonviable probiotic cells to defend the body against pathogenic microorganisms. To generate nonviable, inviable, inactivated, or dead probiotic cells, many methods are used, such as ionizing radiation (10 kGy), heat treatment at 121 °C for 560 minutes, and ultraviolet (UV) radiation exposure for 530 minutes (Helmy et al., 2023). Among the structural and functional alterations linked to the inactivation process are protein denaturation, enzyme inactivation, nucleotide destruction, DNA breakage, and cell structural deformation. This is known as parabiotic/postbiotics, and it is seen as a new frontier in both the food business and microbial medicine (Piqué et al., 2019).

decreasing pathogenic cell adhesion and By translocation into intestinal cells, heat-inactivated L. plantarum has reduced S. enterica infection in mice's liver, spleen, and blood, among other organs (Tao et al., 2021). Heat-inactivated Leuconostoc mesenteroides cells inhibited the invasion of Caco-2 cells by L. monocytogenes by boosting the host immunological responses (Yap et al., 2024). Yogurt that has been inactivated by heat kill prevents cytokines from rupturing the human intestinal epithelial cells' barrier. It has been discovered that inactivated L. paracasei and L. rhamnosus cells can prevent stomach and colon cancer by enhancing cancer cell apoptosis and decreasing proliferative activity. Grampositive bacteria produce levochoic acid (LTA), a microbeassociated molecular pattern (MAMP) that is recognized by the Toll-like receptor 2 (TLR-2) on the surface of intestinal enterocytes (Frosali et al., 2015). An inflammatory cytokine response is triggered when LTA

binds to TLR-2, starting cellular signaling. By preventing the production of new biofilms and eliminating existing ones, LTA produced from probiotic *Lactobacillus* strains has anti-biofilm characteristics against oral and enteric pathogens, including *S. mutans*, *S. aureus*, and *E. faecalis* (Giordani *et al.*, 2021). Compared to live probiotics, prebiotics, and postbiotics have a number of benefits, such as being simpler to produce and store, having unique mechanisms of action shown in Fig. 2, having better accessibility of MAMP when it comes into contact with pattern recognition receptors (PRR), and having a higher chance of triggering specific reactions through particular ligand–receptor interactions (Li and Wu, 2021).

Antimicrobial Peptides (AMPs): The innate immune systems of many different animals, plants, and microbes depend heavily on AMPs, which are spontaneously generated by immune cells (Li et al., 2021). In addition to biological processes, including immunological regulation, angiogenesis, anticancer activity, and wound healing, they exhibit a broad range of antimicrobial action against bacteria, fungi, viruses, and parasites (Verhoef et al., 2019). A component of the innate immune system, AMPs have several advantages over antibiotics, including the ability to act on multiple target sites on the intracellular targets and plasma membranes of pathogenic bacteria, have strong killing activity against drug-resistant bacteria, save time and energy compared to antibody synthesis via acquired immunity, and reach the target sites more quickly than immunoglobulin (Seixas et al., 2022). Based on their protein structure, origins, net charge, and amino acid sequences. AMPs are divided into several classes. These subgroups include cationic α -helical AMPs, which are ≤ 40 amino acids long (50 percent hydrophobic) and have a charge of +2 to +9 with the C-terminus amidated, and anionic AMPs, which have 5-70 amino acid residues with a net charge range of -1 to -8. The stability of β -sheet AMP and biological processes depend heavily on cationic AMPs, which are peptides with 2-8 cysteine residues that form 1-

4 pairs of intramolecular disulfide bonds (Koehbach & Craik, 2019).

Tryptophan, arginine, proline, histidine, and glycine are among the amino acids found in extended cationic AMPs, which are devoid of the typical secondary structures. The development of harmful bacteria and fungi is significantly impacted by fragments of antimicrobial proteins with a broad-spectrum bactericidal activity, including lysozyme (Starling, 2017). Numerous antimicrobial peptides that have been extracted from various sources have demonstrated efficacy against a broad range of harmful microorganisms described in Table 2. Both the kind and nature of AMPs and the bacterial infections they target are major determinants of their antibacterial action. AMPs have the ability to kill bacteria directly by disrupting their membranes, which ultimately results in the death of the bacterium, or indirectly by modifying the immune system (Erdem Büyükkiraz & Kesmen, 2022).

Nanoparticles (NPs): Because of their distinct chemical and physical properties, nanoparticles (NPs) as an antibiotic substitute for managing multidrug-resistant microbes. NPs are simple to interact with target organisms



Fig. 2: Probiotics as an antibiotic.

AMP Name	Structure Type	Source	Target Microorganisms	Notable Features	References
Magainin-2	α-helix	Frog	P. gingivalis, F. nucleatum, P. intermedia, E. coli, S. aureus	Broad-spectrum activity	(Genco et al., 2003)
Cecropin & PI	α-helix	Silk moth (Cecropin), Pig (P1)	E. coli ML35p, S. aureus, B. subtilis, M. luteus, P. aeruginosa, S. Typhimurium, S. marcescens	High efficacy against Gram- positive & Gram-negative bacteria	(Franco et <i>al.,</i> 2013)
Apo-lactoferrin	α-helix	Bovine & Human	E. coli 0157:H7	Found in milk; strong anti- <i>E. coli</i> action	(Franco <i>et al.,</i> 2013)
Melittin	α-helix	Bee venom	S. salivarius, S. mitis, S. mutans, S. sanguinis, S. sobrinus, L. casei, E. faecalis	Potent oral pathogen inhibitor	(Leandro et al., 2015)
Temporins A & L	Not specified	Frog	MRSA, B. megaterium Bm I I, S. aureus Cowan I, E. coli D2 I	Effective against resistant strains	(Wang et <i>al.,</i> 2021b)
Buforin II	α-helix	Toad	B. subtilis, S. aureus, E. coli, S. Typhimurium, E. coli ML35p, L. monocytogenes	Penetrates bacterial cells	(Steinberg et al., 1997).
Clavanin A	α-helix	Styela clava (tunicate)	Same as above (Buforin II)	Marine AMP with broad activity	(Steinberg et al., 1997).
Protegrin-I	β-sheet	Pig & Human	P. aeruginosa, MRSA	β-sheet; strong pore-forming activity	(Han et al., 2024)
Tachyplesin-I	β-sheet	Horseshoe crab	S. Typhimurium	Stable β-structure AMP	(Han et al., 2024)
Hepcidin	β-structure	Human	S. aureus, S. epidermidis, E. coli	Iron-regulating hormone with AMP activity	(Michels et al., 2015).
Daptomycin	Cyclic lipopeptide	Streptomyces roseosporus	MRSA	Clinical antibiotic	(Blaskovich et al., 2018)
Nisin	Lantibiotic	Lactococcus lactis	S. pneumoniae, Enterococci, C. difficile, MRSA	Widely used food preservative and clinical interest	(Shin et al., 2016)
Peptides P1–P3	Short peptides	L. rhamnosus GG (in Chicken)	Avian pathogenic E. coli (APEC)	Novel peptides with activity	(Kathayat et al., 20.21)

because of their tiny size (1-100nm) and vast surface area. Their diverse chemical structures and ability to be produced from a variety of sources enable a range of functions. By focusing on important active areas in pathogens, NPs can partially or completely limit their antibacterial activity (Zaidi et al., 2017). Metal and metal oxide are examples of inorganic NPs that have bactericidal action against bacteria through a variety of methods, earning them the moniker "nanobactericides." The structural elements and growth rate of bacterial cells determine their sensitivity to NPs. While non-porous cell walls function as barriers to penetration, gram-positive bacteria's permeable and negatively charged cell walls make them more vulnerable to NPs. Because fast-growing bacteria express distinct stress-response genes, slowgrowing bacteria are more vulnerable to NPs (Khan & Al-Khedhairy, 2017).

Damage to the cell membrane, inhibition of electron transport and oxidative phosphorylation, modification of bacterial metabolism through disruption of enzymes, DNA, and ribosomes, inhibition of biofilm formation, oxidative stress, and elicitation of host immune responses are all part of NPs' deadly effects on microbial cells (Jagadeeshan & Parsanathan, 2019). The accumulation of silver nanoparticles (Ag) in human organs, oxidative damage caused by CuO, ZnO, or TiO2 NPs, accumulation of metallic NPs in tissues that may cause renal damage and liver or lung toxicity, the absence of a well-described standard technique that is not influenced by NP properties, and the possibility of bacterial resistance to NPs after exposure to metal NPs or metal-oxide NPs are some of the major limitations of NPs (Dianová *et al.*, 2022).

Essential Oils (EOs): EOs are intricate natural mixtures with anywhere from 20 and 60 different components in different ratios. Terpenes, terpenoids, and aromatic and aliphatic groups from various natural sources make up the majority of their constituents (85%) (Chouhan et al., 2017: Sen & Gençer, 2023)). Many pathogens, including P. aeruginosa, S. pyogenes, S. mutans, S. sanguis, S. salivarius, and E. feacalis, have been demonstrated to be antagonistically affected by EOs. Extracted from different sources, thymol and carvacrol have demonstrated the ability to inhibit the growth of S. aureus, C. hystoliticum, C. perfringens, E. coli O157:H7, S. Typhimurium, S. Enteritidis, and L. monocytogenes. P. aeruginosa, K. pneumoniae, E. coli (DH5a), E. coli (MTCC 723), S. Typhimurium, S. aureus, S. epidermidis, and S. mutans have all been demonstrated to be inhibited by linalyl acetate, α -terpineol, nerol, and G. citrata Ehrh (Verma et al., 2016; Bajer et al., 2017).

Because of their hydrophobic properties, EOs and their constituents can interact with the lipids found in microbial cell membranes. Bacterial cell death, cytoplasmic membrane breakdown, and leakage of vital intracellular components can result from their ability to sensitize cells and seriously damage membranes. By binding to them, lowering virulence factors linked to cell walls, and translating certain target microorganism regulatory gene products, EOs can also directly target the production of biofilms. Because food ingredients contain a lot of fat, proteinaceous, and sugary substances that might interfere with their action, using EOs in food preparation decreases their availability as antimicrobial agents. Furthermore, EOs given to food have doses and concentrations that are 10-100 times lower than those seen in vitro, which reduces their effectiveness. Additionally, adding EOs to food products may alter their physical characteristics, including flavor and odor (Omonijo *et al.*, 2018).

Animal studies have shown that essential oils (EOs) are natural substitutes for antibiotics as part of livestock diets. Pan et al. (2023) discussed the biological effects of EOs, like antimicrobial, antioxidant, and anti-inflammatory activity, and pointed out their capability to enhance feed proficiency, boost immune functions, and decrease pathogenic burden in livestock, particularly in swine and poultry. Mo et al. (2022) showed that microencapsulated essential oils were superior to unencapsulated oils and antibiotics in weaning piglets. These benefits included better gut morphology, greater integrity of intestinal barrier function, and an improved gut microbiota status with more beneficial bacteria. In the same context, Giannenas et al. (2013) summarized a wide number of trials and found that EOs such as thymol, carvacrol, and cinnamaldehyde are responsible for improved growth performance, feed conversion, and microbial equilibrium in the poultry, pig, and ruminant gastrointestinal tract. Simitzis (2017) elaborated on this by highlighting the function of EOs in not only enhancing growth and immunity but also the quality of meat and products resulting from it, such as via increased oxidative stability and flavor. Lastly, Coles et al. (2023) conducted a review on several studies of broiler chickens and concluded that supplementation with essential oils in the diet markedly decreased the prevalence and severity of necrotic enteritis, a serious gut disease, by inhibiting Clostridium perfringens while enhancing gut health without impairing performance. Together, these studies prove that EOs can be a powerful and natural approach to replace antibiotics in animal feed to improve animal health, growth, and product quality and reduce the risk of antimicrobial resistance.

Bacteriophages: Bacteriophages, viruses that infect bacteria, have also been contemplated as specific biocontrol agents. They can substantially reduce bacterial populations without eliminating other useful microorganisms (Ke et al., 2024). Since phages are very active against antibiotic-resistant bacteria, they have been exploited to control L. monocytogenes and C. jejuni in food products (Yang et al., 2024). Phages can also be used for surface decontamination if applied at the time of food processing (Hill, 2019). Various Animal studies have shown the promise of bacteriophages as alternatives to antibiotics, especially against multidrug-resistant bacteria.

The research favors phage therapy as a specific and flexible method with limited effect on positive microbiota. Likewise, Joerger (2003) indicated that bacteriophages hold promise in poultry models by targeting Salmonella and Campylobacter effectively, which reduces colonization and enhances food safety, but without the downfalls of developing antibiotic resistance. In a systematic review by Taati Moghadam et al. (2020) reported evidence from various models of animals (mice, rabbits, and rats), demonstrating that phage therapy can effectively decrease the severity of wound infection, bacterial load, and inflammation triggered by multidrugresistant bacteria like A. baumannii and K. pneumoniae. Kutateladze and Adamia (2010) reporting on effective animal trials in which phage treatment precluded or cured infection by antibiotic-resistant strains with low toxicity and good tolerability. Plumet et al. (2022) surveyed several animal model studies of S. aureus infections, namely skin, bone, and systemic infections. The phage therapy in these models produced significant decreases in bacterial burden, enhanced healing rates, and, in a few instances, superior efficacy than conventional antibiotic regimens. Taken overall, these trials very clearly suggest that bacteriophages are a viable alternative or adjunct to antibiotics in veterinary medicine and could be an important weapon in the war against antibiotic resistance.

Organic Acids: Organic acids have also been widely investigated as possible natural alternatives to antibiotic growth promoters in agriculture. Saki et al. (2012) found that supplementing broiler diets with organic acids highly improved growth performance, intestinal integrity through the suppression of pathogenic bacteria like E. coli, and beneficial microorganisms like *Lactobacillus*, and enhanced immune function, thus suitable for organic poultry farming (Saki et al., 2012). Khan et al. (2022) further proved that organic acids such as formic, propionic, lactic, and butyric acids lower gut pH, enhance nutrient digestibility, increase enzyme activity and gut morphology, and exhibit strong antimicrobial, antifungal, and immunomodulatory effects and thus can help in sustainable poultry production considering growing pressure on antibiotic usage (Khan et al., 2022). Mroz (2005) showed that in pigs, organic acids enhance feed intake, protein digestion, daily gain, and feed efficiency, especially under unsatisfactory hygienic conditions, by lowering gastric pH а (

and	inhibitir	ng	pathog	gens	such	as	Salmonel	la a	and
Clost	ridium,	as	well	as	enhanc	ing	intestinal	bar	rier
Table 3: Organic acids as an antibiotic alternative									

functions and immune responses (Mroz, 2005).

Common organic acids (OAs) have a complex mode of action, which may be partly due to factors such as inhibiting development of pathogenic microbes in the the gastrointestinal tract by reducing gut pH, reducing gastric emptying rates, and maintaining endogenous enzyme secretion, mineral chelation and stimulation on intermediary metabolism, and facilitation of proper digestion due to lower gastric pH and enhanced pepsin secretion. Organic acids are weak acids that can easily diffuse across cell membranes, influencing cell metabolism and disrupting normal microbial cell functioning. They have a stronger effect on inhibiting gram-positive bacteria than gram-negative bacteria due to structural differences in the cell membrane described in Table 3 (Rathnayake et al., 2021).

The application of organic acids as antibiotic alternatives has registered encouraging outcomes in various animal studies. By Long et al.'s (2018) study conducted a mixture of organic acids such as formic, lactic, and propionic acids considerably enhanced growth performance, strengthened serum immunity (especially IgG and IgA), and improved intestinal morphology and microbiota of weaned piglets. Acids increased villus height and enhanced the villus-to-crypt ratio, and also encouraged good bacteria such as Lactobacillus and inhibited bad E. coli. Likewise, Dibner and Buttin (2002) presented a critical review that showed that organic acids decrease the pH in the gastrointestinal tract to make it less favorable for pathogens such as Salmonella and Clostridium. They also reported that organic acids enhance nutrient digestion, activate the enzymes, and improve mineral absorption, all of which validate their effectiveness as good alternatives to antibiotics. Hassan et al. (2010) demonstrated in poultry that replacing antibiotic growth promoters with citric and propionic acids in broiler rations improved growth performance and modified gut microflora by suppressing

Organic Acid	Dietary Dose	Observations	Reference
Formic Acid, Propionic Acid,	Not specifically quantified	Improved growth performance, gut health, and reduced	(Saki et al., 2012)
Citric Acid	(inclusion in organic broiler diets)	pathogenic bacteria (<i>E. coli</i>).	
Formic, Acetic, Propionic,	0.1-0.5% inclusion in broiler diets	Lowered gut pH, enhanced nutrient digestibility, and	(Khan et al.,
Butyric Acids	(variable)	antimicrobial and immunomodulatory effects.	2022)
Lactic Acid, Formic Acid,	I–2% in pig diets	Improved feed intake, feed efficiency, protein digestion, and	(Mroz, 2005)
Fumaric Acid		reduced gastric pH.	
Oregano-derived organic acids	0.05–0.1% in pig diets	Enhanced intestinal health, increased villus height, and improved	(Talavera et al.,
(e.g., Carvacrol)		growth performance.	2020)
Water-soluble Organic Acids	0.2% in the weanling pig water supply	Improved growth rate and feed efficiency post-weaning.	(Nhara et <i>al.,</i> 2024)
Benzoic Acid	0.5–1.0% inclusion	Improved nutrient utilization, gut health, and performance similar to antibiotic growth promoters.	(Kiarie <i>et al.,</i> 2018)
Formic Acid, Lactic Acid, Essential Oils Combination	0.2–0.4% in piglet diets	Improved digestive enzyme activities, gut morphology, and intestinal health.	(Xu et al., 2018)
Organic Acid Blend (Formic + Lactic + Propionic)	0.3–0.5% in nursery pig diets	Improved growth performance and gut microbial balance.	(Ferronato & Prandini, 2020)
Organic Acid Mixture (Orgacids™)	0.3% in growing pig diets	Decreased gut pH, improved gut microflora, reduced meat cholesterol.	(Tugnoli et al., 2020)
Benzoic Acid + Essential Oils	0.5% total blend in piglet diets	Effective against <i>E. coli</i> F4 infection, improved intestinal health, and growth.	(Rodrigues et al., 2020)
Citric Acid, Malic Acid,	Analytical guantification, not	Detected and quantified via capillary electrophoresis; relevance	(Liu et al., 2020)
Tartaric Acid, Lactic Acid,	feed-related	for guality control.	(,
Succinic Acid (in			
beverages/fruits study)			
Organic Acids during Soy	Not dietary; fermentation study	Organic acid profiles associated with bacterial community shifts	(Liu et al., 2020)
Sauce Fermentation		during fermentation.	
Microencapsulated Organic	0.3% in piglet diets	Improved nutrient absorption, gut barrier function, and	(Choi et al.,
Acids + Essential Oils		immunity under E. coli F4 challenge.	2020)
Organic Acids as Feed Additives (Various types)	0.2–0.5% recommended	Potential replacements for antibiotic growth promoters; improve broiler health and performance.	(Dai et <i>al.,</i> 2021)

pathogenic bacteria and favoring the proliferation of beneficial bacteria. Moreover, Cai *et al.* (2024) researched that an organic acid mixture stimulated not only weaned piglet growth performance, but also improved their intestinal barrier function, antioxidant activity, and fecal microbial richness, indicating the multi-functionality of the compounds. Lastly, Dittoe *et al.* (2018) also pointed to the capacity of organic acids in positively modifying the avian gut with reduced pathogenic load and immune function support, hence offering a natural means of disease prevention and cure without antibiotics. In all, these studies suggest organic acids' capability to fully substitute for antibiotics in the production of livestock and poultry through enhancing gut health, immune function, and general performance.

The Minimum Inhibitory Concentration (MIC) is the lowest concentration of the antimicrobial agent, such as an antibiotic, antifungal, or antiseptic, which inhibits the visible growth of a microorganism after overnight incubation (Chen *et al.*, 2024). It's an important metric in microbiology that allows for the comparison of antimicrobial potency against specific pathogens shown in Table 4.

Phytochemicals: Phytochemicals, including essential oils, phenolics, and alkaloids, are strong antimicrobials. Eugenol, carvacrol, and thymol in plant essential oils such as oregano and clove act on a wide range of bacteria, viruses, and fungi described in Table 5. These natural antimicrobials disrupt microbial cell membranes and inhibit metabolic pathways; hence, they are promising food preservatives as shown in Fig 3 (Aljaafari *et al.*, 2021). AMPs are usually small peptides with broad-spectrum antimicrobial activity. They are found in various types of

organisms, including humans and animals, and plants. For example, defensins and cathelicidins can disrupt microbial membranes, leading to their death (Li *et al.*, 2024). They work effectively against bacteria like *E. coli* and *Salmonella*, food-borne fungi, and viruses. The synthetic ones are under development with better stability and effectiveness (Liu *et al.*, 2020).

Physical alternatives: High-pressure processing (HPP) is a non-thermal technology that inactivates pathogens by applying high pressure (up to 600MPa) to food products (Akanni *et al.*, 2024). The process is very effective against bacteria, viruses, and fungi without compromising food quality. HPP is particularly effective against *Listeria* and *Norovirus* in ready-to-eat foods (Govaris & Pexara, 2021).

UV radiation, especially UV-C (wavelength 200-280 nm), has strong germicidal effects. It penetrates and damages microbial DNA, thus inhibiting replication and causing cell death (Alessio *et al.*, 2024). UV treatment is an accepted procedure in water purification; it has also been applied for the decontamination of food contact surfaces and packaging materials. It is efficient against bacteria such as Salmonella, viruses like Hepatitis A, and spoiling fungi (Jing *et al.*, 2022).

Pulsed Electric Fields (PEF) is a therapy that employs extremely short pulses of high voltage applied to food goods, producing electroporation of microbial cell membranes (Lytras *et al.*, 2024). The method has demonstrated efficacy against germs, viruses, and fungi while maintaining the food's nutritional value and flavor. PEF has the ability to significantly lower the levels of *Listeria* and *E. coli* in liquid foods like milk and juices (Martínez *et al.*, 2020).



Fig. 3: Bacteriophage and Horizontal gene transfer.



Fig. 4: Alternatives of Antibiotic resistance.

 Table 4: The MIC value of different compounds obtained from plants against various foodborne microorganisms

Plant Derivatives	Plant Name	MIC Value	Bacterial sp.	References
Alkaloids, terpenoids, flavonoids, saponins, and	Anogeissusa cuminata	0.29 mg/mL	S. aureus	(Mishra et al., 2017)
tannins		I.51 mg/mL	A. baumannii	. ,
		3.41 mg/mL	C. freundii	
		3.41 mg/mL	E. coli	
		1.51 mg/mL	K. oxytoca	
		0.67 mg/mL	K. pneumoniae	
		0.67 mg/mL	P. aeruginosa	
Flavonoids and β-sitosterol	Azadirachta indica	3.41 mg/mL	S. aureus	(Mishra et al., 2017, Cesa et al.,
		4.27 mg/mL	A. baumannii	2019)
		3.41 mg/mL	C. freundii	
		4.27 mg/mL	E. coli	
		9.63 mg/mL	K. oxytoca	
		4.27 mg/mL	K. pneumoniae	
		9.63 mg/mL	P. aeruginosa	
		I28 μg/mL	H. pylori	
Glucoside, tannins, saponins, terpenoids, flavonoids,	Bauhinia variegata	3.41 mg/mL	S. aureus	(Mishra et <i>al.,</i> 2017)
		9.63 mg/mL	A. baumannii	
		9.63 mg/mL	C. freundii	
		4.27 mg/mL	E. coli	
		9.63 mg/mL	K. oxytoca	
		4.27 mg/mL	K. pneumoniae	
		3.41 mg/mL	P. aeruginosa	
Flavonoids and β-sitosterol	Boerhaavia diffusa	4.27 mg/mL	S. aureus	(Mishra et <i>al.,</i> 2017)
		9.63 mg/mL	A. baumannii	
		9.63 mg/mL	C. freundii	
		NA	E. coli	
		9.63 mg/mL	K. oxytoca	
		9.63 mg/mL	K. pneumoniae	
		9.63 mg/mL	P. aeruginosa	
Punicalagin, ellagic acid, ellagitannin, and flavonoids	Punica granatum	0.29 mg/mL	S. aureus	(Mishra et <i>al.,</i> 2017)
	-	3.41 mg/mL	A. baumannii	
		0.67 mg/mL	C. freundii	
		0.67 mg/mL	E. coli	
		3.41 mg/mL	K. oxytoca	
		3.41 mg/mL	K. pneumoniae	
		0.67 mg/mL	P. aeruginosa	
Myricetin, luteolin 7-O-glucoside, methyl	Soymida febrifuga	067 mg/mL	S. aureus	(Mishra et al., 2017)
angolensate, flavonoids, and sitosterol		1.51 mg/mL	A. baumannii	. ,
-		3.41 mg/mL	C. freundii	

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		4.27 mg/mL	E. coli	
		3.41 mg/mL	K. oxytoca	
		3.41 mg/mL	K. pneumoniae	
		4.27 mg/mL	P. aeruginosa	
Alkaloids, terpenoids, flavonoids and flavins, tannins	Terminalia chebula	1.51 mg/mL	S. aureus	(Mishra et al., 2017)
and their derivatives, glycosides, and steroids		9.63 mg/mL	A. baumannii	
•		NA	C. freundii	
		9.63 mg/mL	E. coli	
		9.63 mg/mL	K. oxytoca	
		9.63 mg/mL	K. pneumoniae	
		9.63 mg/mL	P. aeruginosa	
Lactones, glycosides, steroids, phenolics, alkaloids,	Tinospora cordifolia	4.27 mg/mL	S. aureus	(Mishra et al., 2017)
and terpenoids		NA	A. baumannii	
		9.63 mg/mL	C. freundii	
		4.27 mg/mL	E. coli	
		9.63 mg/mL	K. oxytoca	
		9.63 mg/mL	K. pneumoniae	
		4.27 mg/mL	P. aeruginosa	
Alkaloids, steroidal saponins, flavonoids, and	Tribulus terrestris	3.41 mg/mL	S. aureus	(Mishra et <i>al.,</i> 2017)
flavonol glycosides		9.63 mg/mL	A. baumannii	
		9.63 mg/mL	C. freundii	
		4.27 mg/mL	E. coli	
		4.27 mg/mL	K. oxytoca	
		9.63 mg/mL	K. pneumoniae	
		3.41 mg/mL	P. aeruginosa	
Coumarins, flavonoids, and phenolic	Petroselinum crispum	EO 22.68 µL/mL	B. cereus	(Semeniuc et al., 2017)
	(Essential oil)	10.80 µL/mL	S. aureus	
		47.62 µL/mL	P. aeruginosa	
		10.80 µL/mL	E. coli	
		47.62 µL/mL	S. typhimurium	
Semantic anhydride, d-terpineol, terpenoids, n-	Levisticum officinale EO	47.62 µL/mL	B. cereus	(Semeniuc et al., 2017)
butylidene phthalide, n-butyl phthalide, l, l phenolic,		2.45 µL/mL	S. aureus	

22.68 µL/mL

10.80 µL/mL

47.62 uL/mL

10.80 µL/mL

2.45 µL/mL

22.68 µL/mL

10.80 µL/mL

22.68 µL/mL

0.56 µL/mL

0.06 µL/mL

0.56 µL/mL

0.27 µL/mL

0.56 µL/mL

8 mg/mL

90 µg/mL

10 µg/mL

50 µM

100 µM

0.01 mg/mL

0.045 mg/mL

0.045 mg/mL

0.045 mg/mL

0.09 mg/mL

25 and 50 µg/mL

P. aeruginosa

S. typhimurium

P. aeruginosa

S. typhimurium

P. aeruginosa

S. typhimurium

E. coli

B. cereus

S. aureus

B. cereus

S. aureus

S aureus

S aureus

E. coli

S. aureus

S. aureus

E. cloacae

S. typhimurium

P. aeruginosa

B. cereus

E. coli

E. coli

E. coli

E. coli

Terpenoid, phenol, and rosmarinic acid

Thymol, y-terpinene, and p-cymene

palbinium, palbine, palbidine, and ipomine

There are three types of eudesmol: β -, α -, and γ -.

and β-caryophyllene

Terpenoids and phenols

saponins

Cannabis sativa L., EO Flavonoids and phenol Flavonoids, phenol, and tannins of gallic and ellagic Acrosta phylosuvaursi acids Alkaloids and isoquinoline Coptis chinensis

essential oils, germacrene-D, α -pinene, β -ocimene, Hypericum olympicum

tannins, cardiac glycosides, steroids, flavonoids, and Eucalyptus tereticornis

121 µg/mL $\alpha\text{-pinene, p-cymene, and 1,8-cineole}$ 118 μg/mL Eucalyptus globulus Terpenoids, alkaloids, lignans, and flavonoids Larreatri dentata 60 µg/mL Myrcene, limonene, and α -pinene Alpinia galanga 3.12-25 mg/L Mycobacterium smegmatis mc2 155 Quercetin, kaempferol, and dioxyflavanol Ammannia spp. 125 µg/mL E. coli 250-1000 µg/mL Alkaloids, isoquinoline Berberis vulgaris P. aeruginosa 5 and 10 mg/mL E. coli Tannins and flavonoids Acer saccharum P. aeruginosa P. mirabilis Catharanthus roseus Terpenoid and limonene 25 mg/L P. aeruginosa Phytosterol, sitosterol, and triterpenoids Holarrhena antidysenterica 20 mg/L P. aeruginosa Steroids, alkaloids, phenols, flavonoids, glycosides, 5 mg/mL S. aureus Cuminum cyminum saponins, and tannins Rosemarinic acid, phenolics, and flavonoids 5 µl/mL Salvia fruticosa S. epidermidis Ascaridole, α-terpinene Chenopodium Ambrosioides 170.6 µl/mL S. aureus 12.1–97.5 µg/mL Gallic and ellagic acids Terminalia chebola E. coli 1.56 mg/L Persea lingue Flavonoids. S. aureus

Ipomoea muricata

Alkanna orientalis

Salvia officinalis

Occimomum basilicum EO

Thymus vulgare EO

(Roy et al., 2012) (Dwivedi et al., 2014) (Aghayan et al., 2017) (Maisuria et al., 2015) (Dwivedi et al., 2018) (Siriyong et al., 2017)

(Semeniuc et al., 2017)

(Semeniuc et al., 2017)

(Zengin et al., 2018)

(Ruiz, 2017)

(Chovanová et al., 2015) (Limaverde et al., 2017) (Bag & Chattopadhyay, 2014) (Holler et al., 2012) (Maurya et al., 2013) (Shiu et al., 2013)

(Kakarla et al., 2017)

(Bame et al., 2013) (Dwivedi et al., 2014)

(Mocan et al., 2020)

 Table 5: Plant-derived substances' (PDSs') mode of action against a variety of microorganisms

Phytochemicals	Extract	Mode of Action	Antimicrobial-Resistant Microbes	References
Flavonoids	Cranberry, or Vaccinium macrocarpon alt	alters the production of biofilms	E. faecalis, E. coli, P. aeruginosa	(Sun et al., 2015; Wojnicz et al., 2016)
	Epigallocatechin, myricetin, and robinetin	prevents the production of DNA in bacteria	E. coli	(Cushnie & Lamb, 2005)
	Quercetin	GrYB protein increases β-galactosidase and extracellular phosphatase while inhibiting ATPase activity.	E. coli, S. aureus	(Simões <i>et al.,</i> 2011)
Essential oils (EOs)	Thymus vulgaris, Levisticum officinale, Ocimum basilicum, and Petroselinum crispum essential oils, as well as Cannabis sativa L.	Cell permeability increases, cell components leak out, bacterial cell wall and membrane changes, ATP loss, protein synthesis inhibition, pH disruption, intracytoplasmic damage, DNA damage, and quorum sensing inhibition prevents the production of biofilms	B. cereus, S. aureus, P. aeruginosa, E. coli, S. Typhimurium, S. aureus	(Willers <i>et al.,</i> 2017; Mocan <i>et al.,</i> 2020)
Tea tree oil (TTO)	Terpenes, sesquiterpenes, and monoterpenes	causes cell death, destroys the cell membrane, interferes with cell development, and alters membrane permeability.	E. coli, S. aureus, C. albicans	(Carson et <i>al.,</i> 2006; Sharifi-Rad et <i>al.,</i> 2017)
Plant-derived peptides	Moringa oleifera	Membrane disruption	E. coli, S. aureus, P. aeruginosa, S. Typhimurium	(Suarez <i>et al.,</i> 2005)
Natural Efflux	Carnosic acid, chalcones,	inhibit several bacterial efflux pumps (EtBr	MDR Uropathogenic E. coli,	(Sabatini et al., 2017; Agbayan et al., 2017;
inhibitors (EPIs)	piperine, reserpine, and gallotannin		isolates)	Willers et al., 2017)

Challenges and future perspectives: For instance, bacteriophage resistance arises by mutations in the bacteria binding that prevent phage or through CRISPR/Cas-mediated immunity (Kadkhoda et al., 2024). Similarly, biofilms may protect pathogens against AMPs and essential oils. Again, this illustrates the importance of multi-target strategies to minimize risks of resistance. The cost of developing, producing, and deploying such alternatives like CRISPR/Cas systems, nano-based antimicrobials, and bacteriophages is inhibitive (Ahmed et al., 2024). Most technologies are yet to overcome scaling challenges for application in agricultural and food processing systems.

Antibiotic resistance in foodborne pathogens is a very complex global issue. This requires creative solution strategies, harmonization of responses, and the future perspective must point toward multi-strategy approaches, advanced research and technology, and well-developed cohesive global policies (Jia et al., 2024). Precision microbiology, synthetic biology, and AI are the grounds for the new generation of technologies facing the resistance management challenge. Under precision microbiology, the analysis of pathogen genomes will make way for effective targeted interventions for highly targeted antimicrobials (Branda & Scarpa, 2024). Utilizing the CRISPR/Cas system, new approaches have been developed that are capable of knocking out resistant genes selectively without affecting useful bacteria. In this respect, governmentindustry-academia collaborations may be useful in promoting research funding, sharing of data, and harmonized regulatory environments (Purnell, 2024). Such actions would specifically serve to support LMICs, which are relatively speaking more challenged in taking full approaches to resistance. Any future effort at ensuring innovation, collaboration, and sustainable practices can potentially serve in prevention to protect the world's food systems from antimicrobial resistance towards foodborne pathogens.

Conclusions: The emergence and increase of antibiotic resistance among foodborne pathogens including Salmonella, Campylobacter, E. coli, and L. monocytogenes are an immense risk to public health and food safety on a worldwide basis, even further increased due to the use of antibiotics under non-regulated manners in farming and clinical fields, notably within low- and middle-income economies. This review highlights, the latest research and developments have brought forward a large range of physical, chemical, and natural substitutes to reduce antibiotic use without impairing effective control of pathogens. Natural substitutes such as bacteriophages, probiotics, antimicrobial peptides (AMPs), and essential oils have demonstrated promising results in the inhibition of pathogenic bacteria and food preservation and safety. Physical means, such as nano-based systems and CRISPR/Cas systems, present very effective and targeted means of managing bacterial populations with minimal reliance on conventional antibiotics. Chemical means, such as the creation of novel food-grade preservatives and the application of nanoparticles, provide other means of managing microbial development and increasing the shelf life of foods. Together, these options hold great promise to upgrade food processing, preservation, and animal husbandry operations into sustainable and health-friendly systems. To move from promise to practice, global efforts must prioritize; Research investment, policy and public-private capacity building, and regulation, collaboration. For their effective inclusion in conventional food production, there needs to be intense global coordination, the setup of effective regulatory systems, and ongoing oversight to ensure the safe, efficient, and sustainable utilization of these advanced solutions.

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