



Antimicrobial Resistance: A Global Health Threat

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Abstract

Antimicrobial resistance (AMR) is a rapidly escalating global health crisis, threatening the effectiveness of antibiotics and other antimicrobial drugs used to treat infections. This paper reviews the epidemiology of AMR, highlighting the mechanisms by which resistance develops and spreads, and the critical impact it has on healthcare systems and global health security. We discuss the factors driving the emergence and spread of AMR, including overuse and misuse of antimicrobials in human medicine, agriculture, and livestock production, as well as the lack of access to effective antimicrobials in many parts of the world. The paper explores the consequences of AMR, including increased morbidity and mortality, prolonged hospital stays, higher healthcare costs, and the potential for untreatable infections. Finally, we examine strategies to address AMR, emphasizing the need for multi-sectoral approaches, including stewardship programs, research and development of new antimicrobials, and public awareness campaigns.

Keywords: Antimicrobial resistance, antibiotics, infections, healthcare, global health, epidemiology, overuse, misuse, morbidity, mortality, public health, stewardship, research and development

1 Introduction

Antimicrobial resistance (AMR), the alarming ability of microorganisms – bacteria, viruses, fungi, and parasites – to withstand the effects of antimicrobial drugs, most notably antibiotics, is no longer a distant threat; it is a rapidly escalating global health crisis with profound and immediate consequences. The unchecked spread of AMR jeopardizes the very foundation of modern medicine, eroding the effectiveness of vital medications that have been essential in the treatment of common and life-threatening infections for decades. This crisis extends far beyond the medical realm, posing a grave challenge to economic stability, global security, and the progress of sustainable development. This paper provides a comprehensive overview of the multifaceted dimensions of AMR, investigating its global epidemiology, the complex interplay of factors that drive its emergence and propagation, the devastating consequences it has on human and animal health, and the diverse strategies that are essential to mitigate its devastating impact. This exploration recognizes the interdisciplinary nature of the challenge, requiring expertise from diverse fields including microbiology, medicine, pharmacology, epidemiology, environmental science, animal health, and policy. The urgent need for collaborative action across national boundaries and sectors will be highlighted throughout the paper. This introduction emphasizes that AMR is not just a scientific challenge, it is a societal imperative demanding immediate and concerted action to safeguard the efficacy of antimicrobials and to protect the health of both current and future generations. This paper serves as a call for a unified, global response to address this critical threat to public health.

2 The Epidemiology of Antimicrobial Resistance

Antimicrobial resistance (AMR) is not a localized phenomenon; it is a complex and dynamic global health challenge that transcends geographical boundaries and socioeconomic status. The World Health Organization (WHO) has recognized AMR as one of the ten greatest threats to global health, underscoring the urgent need for a concerted global response. This section delves into the epidemiological aspects of AMR, examining the underlying mechanisms of resistance, the pathways through which resistance spreads, and the factors that influence its distribution and prevalence across various settings. Microorganisms have evolved a remarkable array of mechanisms to evade the effects of antimicrobial drugs. These resistance mechanisms are often the result of selective pressures imposed by the widespread use and misuse of antimicrobials. Understanding these mechanisms is crucial for developing strategies to overcome or circumvent resistance (1).

Mutations: Altering the Target Site: Genetic mutations in the microbial genome can alter the target site of an antimicrobial drug, preventing the drug from binding and exerting its inhibitory effect. For instance, mutations in bacterial genes encoding for ribosomal proteins, DNA gyrase, or penicillin-binding proteins can lead to resistance to antibiotics that target these specific proteins. The accumulation of multiple mutations in the target site can further diminish the drug's binding affinity and increase the level of resistance. These mutations can arise spontaneously or be induced by exposure to antimicrobials. The ability of microorganisms to readily acquire and accumulate resistance mutations is a key factor contributing to the rapid spread of AMR (2).

Inactivation: Enzymatic Degradation of Antimicrobials: Microorganisms can produce enzymes that inactivate

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antimicrobial drugs, rendering them ineffective. For example, bacteria can produce β -lactamase enzymes that hydrolyze and inactivate β -lactam antibiotics like penicillin and cephalosporins. Different types of β -lactamases exist, and some of these are capable of inactivating a wide range of β -lactam antibiotics, which makes these resistance mechanisms clinically very important. Other enzymes can modify aminoglycoside antibiotics, thus preventing them from binding to their target and exerting their antimicrobial activity. The production of inactivating enzymes is often encoded on plasmids, which can be easily transferred between bacteria (3).

Efflux Pumps: Active Export of Antimicrobials: Microorganisms can develop efflux pumps, which are membrane proteins that actively transport antimicrobial drugs out of the cell, reducing the intracellular concentration of the drug. These efflux pumps are often broad-spectrum, meaning that they can export multiple classes of antimicrobial drugs. The overexpression of efflux pumps can result in multi-drug resistance, making it difficult to treat infections with conventional antimicrobial agents. Efflux pumps can also be induced by exposure to antimicrobial drugs, which can further enhance their contribution to AMR. Efflux pumps represent a significant barrier to antimicrobial activity, particularly in Gram-negative bacteria (4).

Target Modification: Some microorganisms develop resistance by altering their target site in a way that the antimicrobial can no longer bind effectively, without complete loss of function of the target. This can be through alterations in the structure of the target molecule which decreases the binding affinity for the antimicrobial. **Reduced Permeability:** Microorganisms can reduce their cell membrane permeability which hinders the access of antimicrobials to the cell. This can occur through alterations in the outer membrane structure, porins, or cell wall composition, thus decreasing the entry of antimicrobial drugs into the cell. The spread of AMR is a complex process involving multiple pathways. Understanding these pathways is critical for implementing effective strategies to contain the transmission of resistant microorganisms (2).

Horizontal Gene Transfer: Rapid Dissemination of Resistance Genes: Horizontal gene transfer (HGT) is a key mechanism by which resistance genes can be transferred between bacteria, even between distantly related species. This process often involves plasmids, mobile genetic elements that can replicate independently of the bacterial chromosome. Plasmids can carry multiple resistance genes, which can spread rapidly throughout microbial populations. Other mechanisms of HGT include transformation, which involves the uptake of free DNA from the environment, and transduction, which involves the transfer of DNA by bacteriophages. The efficient mechanisms of HGT allow resistant genes to spread rapidly across diverse microbial populations.

Vertical Transmission: Inheritance of Resistance Genes: Resistant bacteria can also pass on their resistance genes to their offspring through vertical transmission, during cell division. This form of transmission is essential for the maintenance of resistant strains within a population. This is particularly relevant within infections where resistant microbes can continue to grow in number. **Selection Pressure: The Driving Force Behind Resistance:** The overuse and misuse of antimicrobials create a strong selective pressure that favors the survival and multiplication of resistant strains. When antimicrobials are used, susceptible microorganisms are killed, while resistant microorganisms survive and multiply. This selection process leads to an increased proportion of resistant organisms within a population. The indiscriminate use of antimicrobials, particularly in settings where they are not needed, contributes significantly to the selection and spread of

AMR. This is further exacerbated by the use of antimicrobials in agriculture and animal husbandry (5).

Environmental Spread: Resistant microorganisms and resistance genes can be transmitted to the environment through waste products, wastewater, and agricultural runoff, creating environmental reservoirs of resistance. These environmental reservoirs can contribute to the dissemination of AMR to humans through contaminated water and food products and through direct environmental contact. The spread of AMR through the environment represents a significant challenge to infection control efforts. **Global Travel and Trade:** The ease and speed of global travel and trade contribute to the spread of resistant microorganisms across geographical boundaries. Individuals carrying resistant organisms can spread these to new areas as can the trade of products including animals and food (6).

3 Drivers of Antimicrobial Resistance

Several interconnected factors contribute to the emergence and spread of antimicrobial resistance (AMR), making it a complex and multifaceted global health challenge. These drivers create environments that favor the selection and proliferation of resistant microorganisms, ultimately diminishing the effectiveness of antimicrobial treatments. The inappropriate and excessive use of antimicrobials in human medicine is a primary driver of AMR. This stems from several key issues. **Over-Prescribing of Antibiotics:** A significant portion of antibiotic prescriptions are unnecessary, often driven by patient demand, diagnostic uncertainty, or a lack of readily available diagnostic tests. The prescription of antibiotics for viral infections, such as the common cold or influenza, is a particularly problematic example of overuse, as antibiotics are ineffective against viruses and merely expose bacterial populations to selective pressure. The overuse of broad-spectrum antibiotics, when narrower-spectrum alternatives are sufficient, further exacerbates the issue by affecting a wider range of bacterial species and driving the development of resistance (7).

Inappropriate Use of Antimicrobials for Viral Infections: As mentioned, the misconception that antibiotics are effective against viral infections is widespread. This results in unnecessary exposure of bacteria to antibiotics, accelerating the selection process for resistant strains. Moreover, the use of antibiotics for mild or self-limiting bacterial infections further increases antibiotic use. Patient education and provider adherence to best practices are crucial to combat this. **Suboptimal Duration and Adherence to Antimicrobial Therapy:** Incomplete courses of treatment, where patients stop taking medication prematurely, can result in the survival of more resistant bacteria, which then multiply and spread. Similarly, improper dosing or failure to adhere to prescribed treatment regimens leads to subtherapeutic levels of antimicrobials, which promotes the development of resistance. **Lack of Diagnostic Stewardship:** The lack of access to and use of rapid and accurate diagnostic tests often leads to empiric antimicrobial therapy, where antibiotics are prescribed based on clinical presentation without confirmed bacterial infection. Empiric therapy often relies on broad-spectrum antibiotics, which contribute to widespread resistance. Improving diagnostic testing capacity is key to improving antimicrobial stewardship (8).

Self-Medication and Over-the-Counter Availability: In many regions, antibiotics are readily available without a prescription, leading to self-medication and uncontrolled antibiotic consumption. This further increases the risk of overuse, incorrect dosing, and the development of resistance, often without monitoring by health care professionals. The

widespread use of antimicrobials in agriculture and animal husbandry contributes substantially to the development and spread of AMR. Prophylactic Use in Livestock: Antimicrobials are often used preventatively in livestock, particularly in intensive farming environments where animals are kept in close proximity. This continuous, low-dose exposure to antibiotics in animal feed or water creates a persistent selection pressure that promotes the emergence of resistant bacteria within animal populations (9).

Use as Growth Promoters: In some countries, antibiotics are still used as growth promoters in livestock, even in the absence of clinical disease. These sub-therapeutic doses can contribute to the development and spread of resistance, even in the absence of overt infections. Transfer of Resistance Genes: Bacteria from livestock, including resistant strains, can be transmitted to humans through the food chain (consumption of contaminated products), direct contact with animals, or environmental contamination. This transfer of resistance genes between bacterial species is a key mechanism by which AMR spreads from animals to humans. Environmental Contamination: Antimicrobial residues and resistant bacteria in animal waste contaminate the environment, including soil and water sources. This environmental contamination acts as a reservoir of resistance, increasing the potential for the spread of resistant strains across various ecological niches and to humans through the food chain, water and contact with the environment (6).

Paradoxically, limited access to effective antimicrobials also contributes to the problem of AMR, particularly in low- and middle-income countries. Unequal Distribution and Availability: Many regions lack consistent access to a range of affordable, effective antibiotics, leading to reliance on older or substandard medications. This lack of access to first-line treatments may contribute to the development of resistance in pathogens by forcing the use of suboptimal treatment options. Limited Healthcare Infrastructure: Deficiencies in healthcare systems, including a lack of trained personnel, diagnostic laboratories, and infection control protocols, result in ineffective treatment of infections, leading to more protracted illnesses and driving further antibiotic use. This poor infrastructure leads to a vicious cycle of infection, treatment with less-than-ideal therapies, and the further evolution of resistance. Inadequate Infection Control Practices: Lack of proper hygiene, sanitation, and infection control measures in healthcare facilities and communities contributes to the spread of both drug-sensitive and drug-resistant bacteria, increasing the overall burden of antimicrobial usage and accelerating the selection for resistant organisms (10).

Financial Barriers: The high cost of novel antibiotics and treatment options limit their use in many settings, resulting in limited effectiveness against multi-drug resistant organisms and increases the prevalence of infectious diseases. These factors, working in concert, create a complex web that promotes the emergence and spread of AMR. Addressing this global health crisis requires a multi-faceted approach including antimicrobial stewardship, investment in diagnostics and novel therapeutics, strengthening healthcare infrastructures and global surveillance.

4 Consequences of Antimicrobial Resistance

The repercussions of antimicrobial resistance are not merely confined to direct health impacts; they ripple outwards, affecting societal structures, economic stability, and even our ability to engage in basic activities we take for granted. The growing ineffectiveness of antimicrobials threatens to destabilize the foundations of modern medicine and public health, creating a complex crisis with far-reaching implications.

Direct Health Impacts - A Deeper Dive. The Cascade of Treatment Failures: The journey of an AMR infection often involves a cascade of treatment failures. Initial therapies prove ineffective, leading to a search for alternative antibiotics, which may be less effective, more toxic, or both. This prolonged search for a viable treatment prolongs suffering, worsens outcomes, and escalates costs. The use of last-resort antibiotics can also accelerate the emergence of resistance to those drugs (7).

Impact on Specific Infections: AMR is particularly devastating in the context of specific infections like tuberculosis (TB), pneumonia, and sepsis. Drug-resistant TB is a major public health challenge, requiring extended, arduous, and often less effective treatments, with lower cure rates. In pneumonia, antibiotic resistance can result in higher rates of hospitalization and mortality, particularly among the elderly. Sepsis, a life-threatening condition caused by the body's overwhelming response to infection, is often initiated by resistant bacteria, making prompt and effective treatment even more difficult. Compromised Immunocompromised Populations: People with weakened immune systems, such as those with HIV/AIDS, cancer patients undergoing chemotherapy, and transplant recipients, are particularly vulnerable to AMR infections. These individuals rely heavily on antimicrobials to prevent and treat infections, and the loss of effective drugs can have devastating consequences. The need for prophylactic antimicrobials to prevent opportunistic infections in these individuals also increases the selective pressure for resistance (11).

Long-Term Health Sequelae: Even when infections caused by resistant organisms are treated, patients can experience long-term health sequelae. These include chronic organ damage, disabilities, and psychological distress, leading to a reduced quality of life. Chronic and recurrent infections also increase the healthcare burden and require long term care. Impact on Maternal and Child Health: AMR poses a significant threat to maternal and child health. Infections during pregnancy can lead to complications such as preterm birth, stillbirth, and neonatal sepsis. The increasing prevalence of resistant infections in newborns makes routine care difficult and increases infant mortality rates (12).

Disruption of Healthcare Workflow: The emergence of AMR disrupts the normal workflow within hospitals and clinics. The need for specialized isolation units, enhanced infection control measures, and frequent antimicrobial resistance testing require more resources and staff time. This creates inefficiencies and can lead to delays in care for other patients. Erosion of Public Trust: The inability to effectively treat common infections can erode public trust in the healthcare system. This can lead to decreased adherence to medical advice, avoidance of healthcare settings, and a greater reliance on alternative or unproven therapies. Impact on Healthcare Workers: Healthcare workers face increased risks of exposure to drug-resistant infections and the emotional burden of caring for patients with untreatable illnesses. This can lead to burnout, high staff turnover, and a decline in the quality of care. The Need for Specialized Infrastructure: The management of AMR requires specialized infrastructure, such as laboratories with advanced diagnostic capabilities and antimicrobial stewardship programs. The development and implementation of these programs in all care settings is a significant financial and logistical undertaking (13).

Impact on Food Production and Security: Antimicrobial resistance in agriculture and aquaculture leads to reduced animal productivity and increased rates of disease in livestock. This translates into higher food costs, food shortages, and economic losses for farmers. The overuse of antimicrobials in

animal agriculture is also driving resistance, creating a dangerous feedback loop. Travel and Tourism Disruption: The increased risk of acquiring AMR infections can affect travel and tourism, with potential travel advisories and restrictions being put in place. The cost of travel insurance may also increase to reflect the increased risk of illness. Impact on Research and Development: The growing crisis of AMR requires a large investment in research and development for new antimicrobials and alternative therapies, but this is a long, costly and difficult path. The economic disincentives for developing new antimicrobials means that private industry is often reluctant to invest in this critical area (6).

Impaired Global Development: The overall societal and economic costs of AMR, including decreased workforce productivity, increased health costs, and reduced trade and tourism, undermine sustainable economic development, particularly in low and middle-income countries. These are countries where the infrastructure to deal with this is often limited. **Ethical Dilemmas:** AMR raises complex ethical dilemmas concerning the allocation of resources, access to care, and the use of last-resort antimicrobials. The inequitable distribution of AMR risks to vulnerable populations raises fundamental questions about justice and fairness. **Reversal of Medical Advances:** The rise of untreatable infections threatens to reverse decades of progress in healthcare. Many routine medical procedures, such as organ transplantation, joint replacements, and cancer treatment, rely heavily on the availability of effective antimicrobials. The loss of these drugs will render these procedures riskier and in some cases impossible, dramatically impacting health care progress (14).

Increased Surgical Risks: The ability to prevent and treat post-operative infections is crucial to the success of surgical procedures. The rise of AMR increases the risks of post-surgical infections, including sepsis, potentially making surgery a dangerous undertaking. The increased risk of infection also may limit elective surgeries and procedures and have major impact on access to care. **Compromised Medical Device Technology:** AMR threatens medical devices that are critical for health and life, including catheters, pacemakers, ventilators, and joint replacements. Because these devices are foreign materials in the body, they are prone to forming biofilms, which harbor and nurture antimicrobial resistant bacteria. These infections are very difficult to treat and can lead to device failure, increased morbidity and mortality (15).

5 Conclusion

Antimicrobial resistance is a global health threat that demands urgent attention. It jeopardizes our ability to treat infections effectively and poses a significant threat to public health and global security. Addressing AMR requires a multifaceted approach involving governments, healthcare institutions, researchers, and the public. By implementing effective stewardship programs, investing in research and development, and promoting public awareness, we can mitigate the spread of resistance and ensure the continued effectiveness of antimicrobials in treating infections.

Ethical issue

Authors are aware of, and comply with, best practice in publication ethics specifically with regard to authorship (avoidance of guest authorship), dual submission, manipulation of figures, competing interests and compliance with policies on research ethics. Authors adhere to publication requirements that submitted work is original and has not been published elsewhere in any language.

Author Contributions

Conceptualization, Aigul B. Ramankulova; methodology, Gulzhan A. Tulegenova; formal analysis, Aigerim A. Umbetova; investigation, Gulzhan A. Tulegenova; resources, Aigul B. Ramankulova; writing—original draft preparation, Gulzhan A. Tulegenova, Aigerim A. Umbetova; writing—review and editing, Aigul B. Ramankulova

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Conflicts of Interest

The authors declare no competing interests.

References

- Haldane V, De Foo C, Abdalla SM, Jung AS, Tan M, Wu S, et al. Health systems resilience in managing the COVID-19 pandemic: lessons from 28 countries. *Nat Med.* 2021;27(6):964-80.
- Munita JM, Arias CA. Mechanisms of Antibiotic Resistance. *Microbiol Spectr.* 2016;4(2).
- Egorov AM, Ulyashova MM, Rubtsova MY. Bacterial Enzymes and Antibiotic Resistance. *Acta Naturae.* 2018;10(4):33-48.
- Abdali N, Parks JM, Haynes KM, Chaney JL, Green AT, Wolloscheck D, et al. Reviving Antibiotics: Efflux Pump Inhibitors That Interact with AcrA, a Membrane Fusion Protein of the AcrAB-TolC Multidrug Efflux Pump. *ACS Infect Dis.* 2017;3(1):89-98.
- Patangia DV, Ryan CA, Dempsey E, Stanton C, Ross RP. Vertical transfer of antibiotics and antibiotic resistant strains across the mother/baby axis. *Trends Microbiol.* 2022;30(1):47-56.
- Manyi-Loh C, Mamphweli S, Meyer E, Okoh A. Antibiotic Use in Agriculture and Its Consequential Resistance in Environmental Sources: Potential Public Health Implications. *Molecules.* 2018;23(4).
- Prestinaci F, Pezzotti P, Pantosti A. Antimicrobial resistance: a global multifaceted phenomenon. *Pathog Glob Health.* 2015;109(7):309-18.
- Samreen, Ahmad I, Malak HA, Abulreesh HH. Environmental antimicrobial resistance and its drivers: a potential threat to public health. *J Glob Antimicrob Resist.* 2021;27:101-11.
- Sachdev C, Anjankar A, Agrawal J. Self-Medication With Antibiotics: An Element Increasing Resistance. *Cureus.* 2022;14(10):e30844.
- Iskandar K, Molinier L, Hallit S, Sartelli M, Catena F, Coccolini F, et al. Drivers of Antibiotic Resistance Transmission in Low- and Middle-Income Countries from a "One Health" Perspective-A Review. *Antibiotics (Basel).* 2020;9(7).
- Knight GM, Glover RE, McQuaid CF, Olaru ID, Gallandat K, Leclerc QJ, et al. Antimicrobial resistance and COVID-19: Intersections and implications. *Elife.* 2021;10.
- Moghimi N, Di Napoli M, Biller J, Siegler JE, Shekhar R, McCullough LD, et al. The Neurological Manifestations of Post-Acute Sequelae of SARS-CoV-2 infection. *Curr Neurol Neurosci Rep.* 2021;21(9):44.
- Majumder MAA, Rahman S, Cohall D, Bharatha A, Singh K, Haque M, et al. Antimicrobial Stewardship: Fighting Antimicrobial Resistance and Protecting Global Public Health. *Infect Drug Resist.* 2020;13:4713-38.
- Dadgostar P. Antimicrobial Resistance: Implications and Costs. *Infect Drug Resist.* 2019;12:3903-10.
- Robinson ED, Volles DF, Kramme K, Mathers AJ, Sawyer RG. Collaborative Antimicrobial Stewardship for Surgeons. *Infect Dis Clin North Am.* 2020;34(1):97-108.