

## Antimicrobial Resistance through the One Health Lens

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Antimicrobial resistance (AMR) has silently evolved into one of humanity's most pressing health threats. This issue transcends conventional boundaries and extends beyond hospitals and clinics to farms, rivers, markets, and even residential kitchens <sup>1,2</sup>. The One Health approach, which recognizes an inseparable triad of humans, animals, and the environment, provides a holistic framework for confronting AMR <sup>3</sup>. It demands that specific fields, such as medicine, agriculture, environmental science, and policy-making, interact and coordinate, even though historically each has functioned individually <sup>4</sup>.

At its essence, AMR — a biological inevitability — is bound to become a socio-behavioral and ecological crisis. Resistant genes do not respect borders; they traverse through trade, travel, and even the simple flow of water <sup>4</sup>. A study conducted by Velazquez-Meza and colleagues reminds us that the evolution of microbial communities is linked intimately to human action <sup>5</sup>. The misuse of antibiotics, inadequate infection control, and agricultural pollution together are promoting an invisible evolutionary race that the humans may not be able to win <sup>6</sup>.

The escalating crisis of AMR demands a multifaceted, globally coordinated response. Recognizing that AMR is not limited to clinical settings but rather represents a complex, cross-sectoral threat to human, animal, and environmental health is essential for addressing the burden more holistically <sup>5-7</sup>. The One Health strategy requires integrated action across human medicine, veterinary practice, agricultural management, and environmental stewardship to curb both the emergence and transmission of resistant pathogens <sup>8</sup>.

### The Perfect Storm: Overuse, Misuse, and Cross-Contamination

From penicillin prescriptions in urban clinics to prophylactic tetracycline in poultry farming, the overreliance on antimicrobials creates sustained selective pressure, producing resilient microbial strains

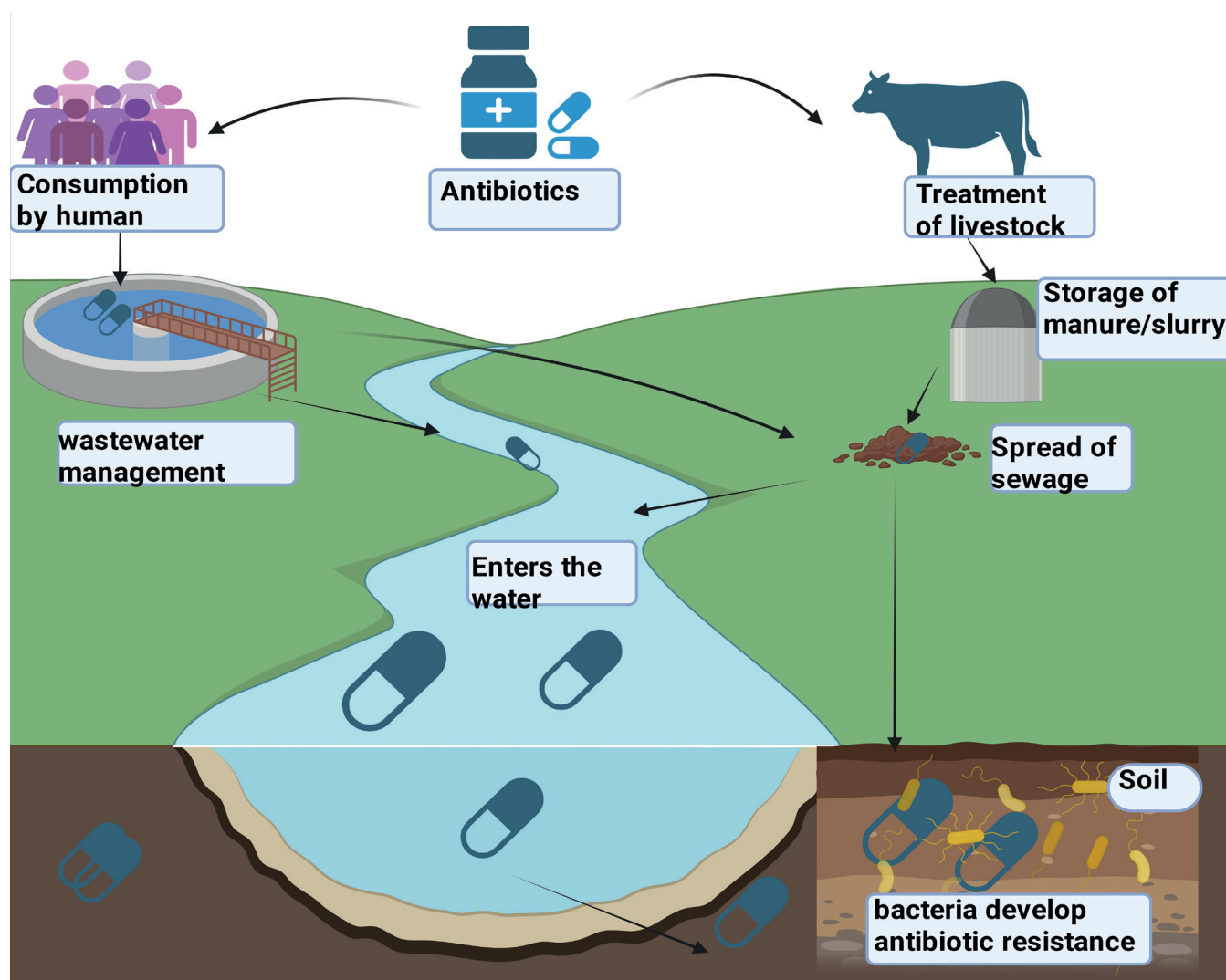
<sup>9</sup>. Antimicrobials that were once considered a triumphant achievement of the human beings have now become a means for the microbials to adapt and transform <sup>10</sup>. A study noted that 75%–90% of antimicrobials administered to animals are excreted unmetabolized into the environment, introducing resistance genes into the crops, water, and soil <sup>5,11</sup>.

This contamination chain converts fields into bacterial laboratories and rivers into corridors of genetic exchange <sup>12</sup> [Figure 1]. Studies have found resistance plasmids in bacterial isolates from vegetables and aquaculture systems—suggesting that environmental pathways spread AMR far more widely than clinical misuse alone <sup>13-15</sup>. The agricultural-industrial complex, remarkably, remains a stubborn frontier for the mitigation of AMR. Therefore, both economic and behavioral change are essential <sup>16</sup>.

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**Figure 1:** Display of the pathways of spread of antibiotic resistance in the environment by human consumption and animal treatment with antibiotics. This figure was drawn via the premium version of BioRender (<https://biorender.com/>), accessed on October 31<sup>st</sup>, 2025, with license number GM28XUBU04 <sup>17</sup>

**Illustration Credit:** Rahnuma Ahmad.

### A Historical Continuum: From Virchow to the Present Crisis

The roots of One Health trace back to Rudolf Virchow's 19th-century concept of zoonosis and Calvin Schwabe's later advocacy for "one medicine" <sup>18</sup>. Therefore, it can be noted that the One Health model is not new; however, it has been overlooked during the antibiotic century <sup>5,18,19</sup>. Modern globalization has only magnified this interconnection. Resistant pathogens can now cross continents in days, turning local epidemics into global challenges <sup>5,20</sup>. Even though the Manhattan Principles marked a turning point in recognizing disease

transmission within ecosystems, nearly two decades later, AMR rates continue to rise <sup>5,20</sup>. Thus, it can be noted that recognition alone is insufficient without radical, sustained intervention.

### Global Action and the Surveillance Imperative

The World Health Organization's Global Action Plan and the creation of the Global Antimicrobial Resistance Surveillance System (GLASS) are examples of large-scale institutional commitments for tracking microbial evolution <sup>21</sup>. These frameworks aim to standardize data across countries. This is an enormous challenge, particularly for regions where the

health infrastructure is weak. To introduce effective interventions, surveillance is essential. Data obtained reveal where and how resistance proliferates, alerting to the rational use of antimicrobials and clarifying cross-sectoral transmission routes<sup>21,22</sup>. The cross-institutional collaboration among the United Nations Food and Agriculture Organization (FAO), the World Health Organization (WHO), and the World Organization for Animal Health (WOAH—formerly OIE) forms a tripartite mechanism to coordinate AMR responses across sectors. These alliances symbolize the necessary interdependence between animal and human health agencies<sup>19,20,23</sup>.

### Antibiotic Use Across Sectors: The Shared Burden

Among the antibiotic classes used across species and ecosystems, the examples of colistin, cephalosporins, and fluoroquinolones are particularly illuminating<sup>24-27</sup>. Colistin, once a reserve antibiotic for life-threatening infections, has led to the development of the plasmid-mediated *mcr-1* gene. This gene has been detected across continents within months of its first identification in China<sup>28</sup>. A similar pattern has occurred in the case of third-generation cephalosporins—ceftiofur in poultry, ceftriaxone in hospitals—producing cross-resistance that bridges farms and hospitals<sup>29</sup>. Fluoroquinolones, too, follow this pattern: their use in poultry promotes resistance in *Campylobacter jejuni*, an agent of foodborne disease that directly impacts human health<sup>30</sup>. Resistance flows bi-directionally between animals and humans, mediated by market forces, inadequate regulation, and environmental leakage<sup>31</sup>.

### The Socioeconomic Dimension

AMR is not only a microbiological problem but also a socioeconomic crisis. The economic burden of resistance includes prolonged hospital stays, lost productivity, and increased mortality. These costs are disproportionately borne by low- and middle-income countries, where antimicrobials are often available over the counter and infection control systems remain fragile<sup>32</sup>. This imbalance is ethically troubling. Bacteria do not discriminate, but economic inequities shape who bears the brunt of their resistance. There is a need to strengthen governance, ensure legislative uniformity, and ensure equitable access to clean water and sanitation<sup>33</sup>.

### Human Costs and the Veterinary Paradox

From a clinical viewpoint, AMR reverses decades of medical progress. Common infections become

untreatable<sup>34</sup>. Complex surgeries face renewed risk. Neonatal sepsis, urinary tract infections, and wound infections that were once treatable now threaten mortality in resource-constrained facilities. At the same time, it is becoming increasingly challenging to create a balance between the needs of the food productivity system and the dangers to public health. The practice of using antimicrobials as growth promoters exemplifies this tension. Such practices need to be banned, as they are a scientific necessity<sup>5,35</sup>.

Within agriculture and veterinary medicine, there is the need for continued restraint and judiciousness in antimicrobial use. Curtailing the use of antibiotics as growth promoters in livestock and aquaculture is fundamental to reducing selection pressures that drive resistance<sup>36</sup>. The challenge, however, lies in deploying alternative strategies such as enhanced biosecurity, vaccination programs, and improved husbandry practices to maintain animal health and productivity without exacerbating AMR risks<sup>37,38</sup>. Coordinated efforts by international organizations such as the WOAH, the FAO, and the WHO are pivotal in driving policy harmonization and capacity building at the animal-human-environment interface<sup>19,23</sup>.

### Education, Awareness, and Behavioral Change

Technological interventions, while vital, cannot be a substitute for behavioral transformation. Awareness campaigns—such as the World Antimicrobial Awareness Week—seek to empower both consumers and professionals to act responsibly<sup>39</sup>. Education must modify the patient's insistence on antibiotics, the farmer's reliance on prophylactic feeds, and the policymaker's short-term economic calculations. AMR is sustained not only by ignorance but also by ingrained practices and incentives that outpace regulation<sup>40,41</sup>.

### The Innovation Gap: Where Are the New Drugs?

While the microbial evolution gains pace, human innovation in antibiotic development has slowed dramatically. Of the 32 new antimicrobials under development in hospitals in 2019, only six were truly innovative. Pharmaceutical economics discourages investment in drugs that may quickly become obsolete<sup>42</sup>. Thus, global mechanisms such as the Global Innovation Fund and public-private partnerships (e.g., the Global Alliance for Antibiotic Research and Development) aim to reignite the discovery pipeline<sup>43</sup>. However, new antibiotics alone will not be able to handle the

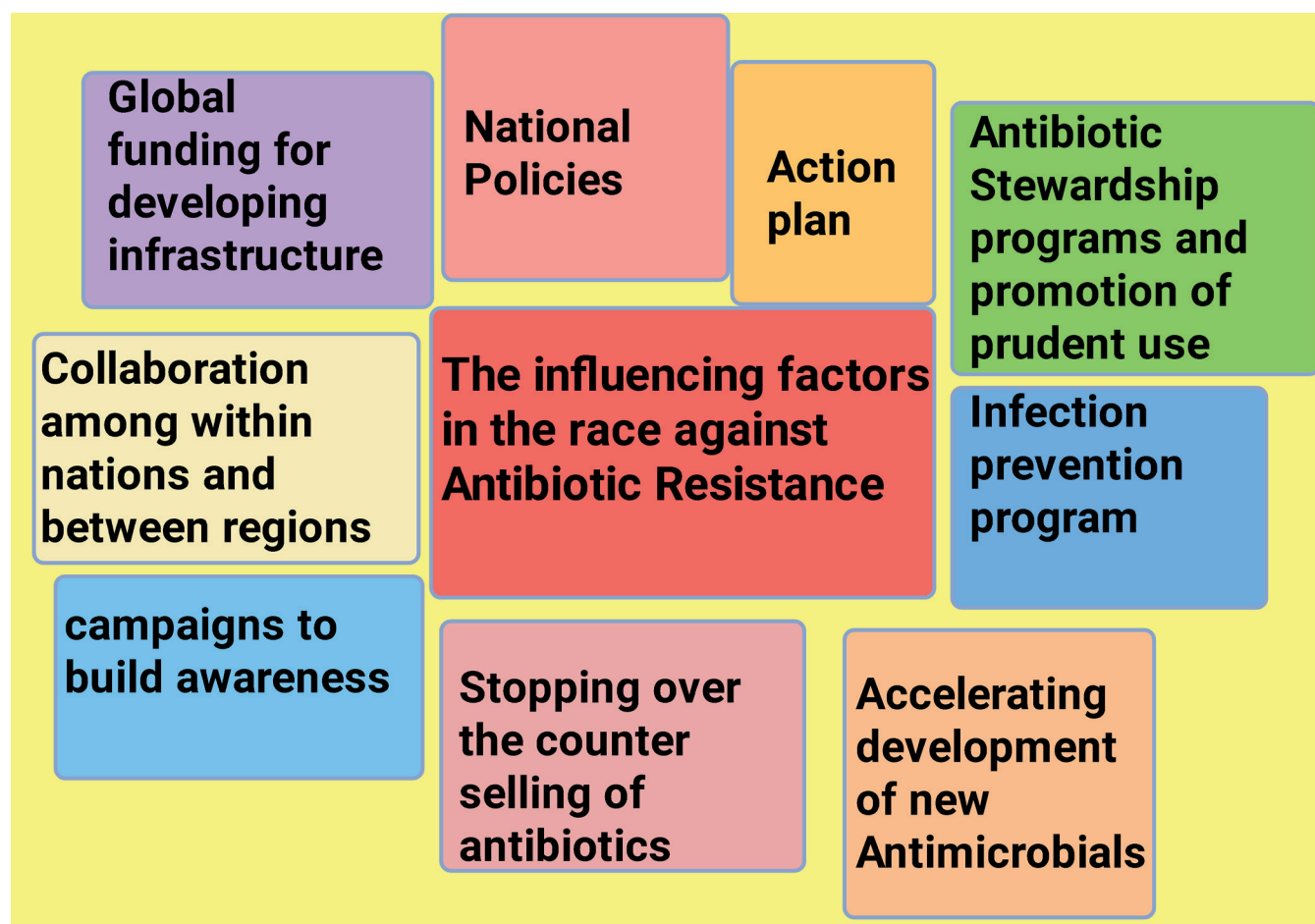
current AMR trend. Without stewardship, even novel agents like ceftazidime-avibactam or meropenem-vaborbactam will soon lose their edge. The challenge lies not only in creation but in conservation <sup>44</sup>.

### Emerging Alternatives and Future Directions

Central to this approach is the acknowledgment that successful AMR mitigation cannot rely solely on the development of new antibiotics. The diminishing pipeline of novel antimicrobials, reducing the overall demand for these drugs, emerges as a critical pillar of intervention. Woolhouse identified effective means to achieve this goal, including expanding Universal Health Coverage (UHC), promoting the widespread adoption of water, sanitation, and hygiene (WASH) protocols, and strengthening infection control measures within healthcare settings <sup>45</sup>. For example, improved WASH infrastructure lowers infection rates among

communities, which in turn diminishes the necessity for antibiotic administration. This domino effect not only preserves the efficacy of current antimicrobials but also contributes to global health equity and sustainability <sup>46</sup>.

Critically, the environmental dimension of AMR is gaining recognition. Woolhouse has emphasized in his study how environmental reservoirs—through wastewater, manure, and run-off—facilitate the dissemination of resistant determinants across sectors <sup>45</sup>. Implementing robust surveillance, waste stream management, and environmental monitoring is essential not only for curbing the spread of AMR genes but also for preserving ecological integrity <sup>47</sup>. Recent global action plans, such as the One Health Joint Plan of Action (OH JPA), stress the need for policy alignment, cross-sector research, and interdisciplinary collaboration to tackle these intertwined challenges <sup>47,48</sup> [Figure 2].



**Figure 2:** Display of the factors that influence the race against Antibiotic Resistance. This figure was drawn via the premium version of BioRender (<https://biorender.com/>), accessed on November 1<sup>st</sup>, 2025, with license number VV28Y1VLY7 <sup>17</sup>. (Illustration Credit: Rahnuma Ahmad).



Expanding vaccine development, bacteriophage therapy, and immunotherapeutic strategies as complements to antimicrobial drugs needs to be considered. These alternatives aim to bypass the resistance hurdle altogether by preventing infections or exploring natural microbial antagonism. Additionally, the integration of rapid diagnostics could curb overprescribing. Machine learning-based diagnostic tools and genomic sequencing for resistance profiling promise to improve clinical decision-making, tailoring antibiotic use with precision<sup>49,50</sup>.

### A Call for Global Solidarity

Ultimately, the AMR crisis is not a distant scientific problem but an existential one. For the One Health Approach to succeed, it is necessary to recognize that humanity cannot separate itself from the ecological networks that it disrupts<sup>51</sup>. Collaboration must be transnational and transdisciplinary. Governments must see antimicrobial stewardship as a security policy and not just medical ethics<sup>52</sup>. The COVID-19 pandemic provided both a warning and an opportunity. It revealed how interconnected global health is, and how resource prioritization can shift rapidly under existential threat<sup>53</sup>. Now, this vigilance must be redirected toward AMR, which kills silently but persistently.

### Towards a Common Future

One Health is not an abstraction—it is a survival strategy. Mitigating AMR requires humility, collaboration, and sustained action across all domains of life. Human well-being, animal welfare, and environmental integrity are not competing interests but interdependent pillars of existence<sup>54</sup>. Advances in surveillance and data-sharing frameworks are now enabling real-time tracking of

resistant strains, antimicrobial consumption patterns, and intervention outcomes. Such systems foster greater accountability, inform targeted stewardship programs, and facilitate evidence-based policy-making that is adaptable to regional contexts and evolving threats<sup>55</sup>. The time for incremental reforms has passed. What remains is the collective determination to redefine what responsible medicine means in a globalized world. The cure for AMR is a mindset.

### Consent for Publication

The author has reviewed and approved the final version and agrees to be accountable for all aspects of the work, including any accuracy or integrity issues.

### DISCLOSURE

The authors declare that they do not have any financial involvement or affiliations with any organization, association, or entity directly or indirectly related to the subject matter or materials presented in this review paper.

### Data Availability

Information for this review paper is taken from freely available sources.

### Authorship Contribution

All authors contributed significantly to the work, whether in the conception, design, utilization, collection, analysis, or interpretation of data, or all these areas. They also participated in the paper's drafting, revision, or critical review, gave their final approval for the version that would be published, decided on the journal to which the article would be submitted, and made the responsible decision to be held accountable for all aspects of the work.

## REFERENCE

1. Salam MA, Al-Amin MY, Salam MT, Pawar JS, Akhter N, Rabaan AA, Alqumber MAA. Antimicrobial Resistance: A Growing Serious Threat for Global Public Health. *Healthcare (Basel)*. 2023;**11**(13):1946. doi: 10.3390/healthcare11131946.
2. Danasekaran R. One Health: A Holistic Approach to Tackling Global Health Issues. *Indian J Community Med*. 2024;**49**(2):260-263. doi: 10.4103/ijcm.ijcm\_521\_23.
3. Leonardi GS, Zeka A, Ashworth M, Bouland C, Crabbe H, Duarte-Davidson R, Etzel RA, Giuashvili N, Gökdemir Ö, Hanke W, van den Hazel P, Jagals P, Khan EA, Martin-Olmedo P, Pett J, Ruadze E, Santamaria MG, Semenza JC, Sorensen C, Vardoulakis S, Yip F, Lauriola P. A new environmental public health practice to manage current and future global health challenges through education, training, and capacity building. *Front Public Health*. 2024;**12**:1373490. doi: 10.3389/fpubh.2024.1373490.
4. Sambaza SS, Naicker N. Contribution of wastewater to antimicrobial resistance: A review article. *J Glob Antimicrob Resist*. 2023;**34**:23-29. doi: 10.1016/j.jgar.2023.05.010.
5. Velazquez-Meza ME, Galarde-López M, Carrillo-Quiróz B, Alpuche-Aranda CM. Antimicrobial resistance: One Health approach. *Vet World*. 2022;**15**(3):743-749. doi: 10.14202/vetworld.2022.743-749.
6. Iwu CD, Korsten L, Okoh AI. The incidence of antibiotic resistance



- within and beyond the agricultural ecosystem: A concern for public health. *Microbiologyopen*. 2020;**9**(9):e1035. doi: 10.1002/mbo3.1035.
7. Oliveira M, Antunes W, Mota S, Madureira-Carvalho Á, Dinis-Oliveira RJ, Dias da Silva D. An Overview of the Recent Advances in Antimicrobial Resistance. *Microorganisms*. 2024;**12**(9):1920. doi: 10.3390/microorganisms12091920.
  8. Cella E, Giovanetti M, Benedetti F, Scarpa F, Johnston C, Borsetti A, Ceccarelli G, Azarian T, Zella D, Ciccozzi M. Joining Forces against Antibiotic Resistance: The One Health Solution. *Pathogens*. 2023;**12**(9):1074. doi: 10.3390/pathogens12091074.
  9. Odey TOJ, Tanimowo WO, Afolabi KO, Jahid IK, Reuben RC. Antimicrobial use and resistance in food animal production: food safety and associated concerns in Sub-Saharan Africa. *Int Microbiol*. 2024;**27**(1):1-23. doi: 10.1007/s10123-023-00462-x.
  10. Muteeb G, Rehman MT, Shahwan M, Aatif M. Origin of Antibiotics and Antibiotic Resistance, and Their Impacts on Drug Development: A Narrative Review. *Pharmaceuticals (Basel)*. 2023;**16**(11):1615. doi: 10.3390/ph16111615.
  11. Tian M, He X, Feng Y, Wang W, Chen H, Gong M, Liu D, Clarke JL, van Eerde A. Pollution by Antibiotics and Antimicrobial Resistance in Livestock and Poultry Manure in China, and Countermeasures. *Antibiotics (Basel)*. 2021;**10**(5):539. doi: 10.3390/antibiotics10050539.
  12. Wang C, Song Y, Liang J, Wang Y, Zhang D, Zhao Z. Antibiotic resistance genes are transferred from manure-contaminated water bodies to the gut microbiota of animals through the food chain. *Environ Pollut*. 2024;**363**(Pt 1):125087. doi: 10.1016/j.envpol.2024.125087.
  13. Elbehiry A, Marzouk E, Abalkhail A, Edrees HM, Ellethy AT, Almuzaini AM, Ibrahim M, Almujaidel A, Alzaben F, Alqmi A, Abu-Okail A. Microbial Food Safety and Antimicrobial Resistance in Foods: A Dual Threat to Public Health. *Microorganisms*. 2025;**13**(7):1592. doi: 10.3390/microorganisms13071592.
  14. Endale H, Mathewos M, Abdeta D. Potential Causes of Spread of Antimicrobial Resistance and Preventive Measures in One Health Perspective-A Review. *Infect Drug Resist*. 2023;**16**:7515-7545. doi: 10.2147/IDR.S428837.
  15. Castañeda-Barba S, Top EM, Stalder T. Plasmids, a molecular cornerstone of antimicrobial resistance in the One Health era. *Nat Rev Microbiol*. 2024;**22**(1):18-32. doi: 10.1038/s41579-023-00926-x.
  16. McKernan C, Benson T, Farrell S, Dean M. Antimicrobial use in agriculture: critical review of the factors influencing behavior. *JAC Antimicrob Resist*. 2021;**3**(4):dlab178. doi: 10.1093/jacamr/dlab178.
  17. BioRender. 2025. Available from <https://biorender.com> [Accessed October 31, 2025]
  18. Cassidy A. Humans, Other Animals and 'One Health' in the Early Twenty-First Century. In: Woods A, Bresalier M, Cassidy A, et al. *Animals and the Shaping of Modern Medicine: One Health and Its Histories*. London (UK): Palgrave Macmillan; 2017 Dec. Chapter 6. doi: 10.1007/978-3-319-64337-3\_6 Available from <https://www.ncbi.nlm.nih.gov/books/NBK481748/> [Accessed November 2, 2025]
  19. Frenk J, Gómez-Dantés O, Knaul FM. Globalization and infectious diseases. *Infect Dis Clin North Am*. 2011;**25**(3):593-9, viii. doi: 10.1016/j.idc.2011.05.003.
  20. Hernandez A, Lee J, Kang H. Navigating the Interconnected Web of Health: A Comprehensive Review of the One Health Paradigm and Its Implications for Disease Management. *Yonsei Med J*. 2025;**66**(4):203-210. doi: 10.3349/ymj.2024.0108.
  21. Ajulo S, Awosile B. Global antimicrobial resistance and use surveillance system (GLASS 2022): Investigating the relationship between antimicrobial resistance and antimicrobial consumption data across the participating countries. *PLoS One*. 2024;**19**(2):e0297921. doi: 10.1371/journal.pone.0297921.
  22. Nazir A, Nazir A, Zuhair V, Aman S, Sadiq SUR, Hasan AH, Tariq M, Rehman LU, Mustapha MJ, Bulimbe DB. The Global Challenge of Antimicrobial Resistance: Mechanisms, Case Studies, and Mitigation Approaches. *Health Sci Rep*. 2025;**8**(7):e71077. doi: 10.1002/hsr2.71077.
  23. United Nations Food and Agriculture Organization (FAO), the World Health Organization (WHO), and the World Organization for Animal Health (WOAH). The Tripartite's Commitment: Providing multi-sectoral, collaborative leadership in addressing health challenges. FAO: Rome, WHO: Geneva, WOAH: Paris. 2017. Available from <https://www.woah.org/app/uploads/2018/05/tripartite-2017.pdf> [Accessed on October 26th, 2025]
  24. El-Sayed Ahmed MAE, Zhong LL, Shen C, Yang Y, Doi Y, Tian GB. Colistin and its role in the Era of antibiotic resistance: an extended review (2000-2019). *Emerg Microbes Infect*. 2020;**9**(1):868-885. doi: 10.1080/22221751.2020.1754133.
  25. Das N, Madhavan J, Selvi A, Das D. An overview of cephalosporin antibiotics as emerging contaminants: a serious environmental concern. *3 Biotech*. 2019;**9**(6):231. doi: 10.1007/s13205-019-1766-9.
  26. Hayer SS, Casanova-Higes A, Paladino E, Elnekave E, Nault A, Johnson T, Bender J, Perez A, Alvarez J. Global Distribution of Fluoroquinolone and Colistin Resistance and Associated Resistance Markers in *Escherichia coli* of Swine Origin - A Systematic Review and Meta-Analysis. *Front Microbiol*. 2022;**13**:834793. doi: 10.3389/fmicb.2022.834793.
  27. Valiakos G, Kapna I. Colistin Resistant *mcr* Genes Prevalence in Livestock Animals (Swine, Bovine, Poultry) from a Multinational Perspective. A Systematic Review. *Vet Sci*. 2021;**8**(11):265. doi: 10.3390/vetsci8110265.
  28. Nagy Á, Székelyhidi R, Hanczné Lakatos E, Kapcsándi V. Review on the occurrence of the *mcr-1* gene causing colistin resistance in cow's milk and dairy products. *Heliyon*. 2021;**7**(4):e06800. doi: 10.1016/j.heliyon.2021.e06800.
  29. Collignon P, Aarestrup FM, Irwin R, McEwen S. Human deaths and third-generation cephalosporin use in poultry, Europe. *Emerg Infect Dis*. 2013;**19**(8):1339-40. doi: 10.3201/eid1908.120681.
  30. Smith JL, Fratamico PM. Fluoroquinolone resistance in *Campylobacter*. *J Food Prot*. 2010;**73**(6):1141-52. doi: 10.4315/0362-028x-73.6.1141.
  31. Institute of Medicine (US) Committee on Emerging Microbial Threats to Health in the 21st Century; Smolinski MS, Hamburg MA, Lederberg J, editors. *Microbial Threats to Health: Emergence, Detection, and Response*. Washington (DC): National Academies Press (US); 2003. 3. Factors in Emergence. Available from <https://www.ncbi.nlm.nih.gov/books/NBK221497/> [Accessed November 2, 2025]
  32. Poudel AN, Zhu S, Cooper N, Little P, Tarrant C, Hickman M, Yao G. The economic burden of antibiotic resistance: A systematic review and meta-analysis. *PLoS One*. 2023;**18**(5):e0285170. doi: 10.1371/journal.pone.0285170.
  33. Weets CM, Katz R. Global approaches to tackling antimicrobial resistance: a comprehensive analysis of water, sanitation and hygiene policies. *BMJ Glob Health*. 2024;**9**(2):e013855. doi: 10.1136/bmjgh-2023-013855.
  34. Oliveira M, Antunes W, Mota S, Madureira-Carvalho Á, Dinis-Oliveira RJ, Dias da Silva D. An Overview of the Recent Advances in Antimicrobial Resistance. *Microorganisms*. 2024;**12**(9):1920. doi: 10.3390/microorganisms12091920.
  35. Sağlam MK, Yanaşoğlu E, Sav NM, Kurt F, Kocabay K, Akbaş E, Sungur MA. The rising threat of antibiotic and multidrug resistance in neonatal urinary tract infections. *BMC Pediatr*. 2025;**25**(1):824. doi: 10.1186/s12887-025-06210-6.

36. Wallinga D, Smit LAM, Davis MF, Casey JA, Nachman KE. A Review of the Effectiveness of Current US Policies on Antimicrobial Use in Meat and Poultry Production. *Curr Environ Health Rep.* 2022;**9**(2):339-354. doi: 10.1007/s40572-022-00351-x.
37. Elbehiry A, Marzouk E. From Farm to Fork: Antimicrobial-Resistant Bacterial Pathogens in Livestock Production and the Food Chain. *Vet Sci.* 2025;**12**(9):862. doi: 10.3390/vetsci12090862.
38. Pinto Ferreira J, Battaglia D, Dorado García A, Tempelman K, Bullon C, Motriuc N, Caudell M, Cahill S, Song J, LeJeune J. Achieving Antimicrobial Stewardship on the Global Scale: Challenges and Opportunities. *Microorganisms.* 2022;**10**(8):1599. doi: 10.3390/microorganisms10081599.
39. Wu D, Walsh TR, Wu Y. World Antimicrobial Awareness Week 2021 - Spread Awareness, Stop Resistance. *China CDC Wkly.* 2021;**3**(47):987-993. doi: 10.46234/ccdcw2021.241.
40. National Research Council (US) Committee to Study the Human Health Effects of Subtherapeutic Antibiotic Use in Animal Feeds. The Effects on Human Health of Subtherapeutic Use of Antimicrobials in Animal Feeds. Washington (DC): National Academies Press (US); 1980. Appendix K, Antibiotics In Animal Feeds. Available from <https://www.ncbi.nlm.nih.gov/books/NBK216502/> [Accessed November 2, 2025]
41. Rocha V, Estrela M, Neto V, Roque F, Figueiras A, Herdeiro MT. Educational Interventions to Reduce Prescription and Dispensing of Antibiotics in Primary Care: A Systematic Review of Economic Impact. *Antibiotics (Basel).* 2022;**11**(9):1186. doi: 10.3390/antibiotics11091186.
42. Sertkaya A, Berlind A, McGeeney JD, et al. Analysis of Market Challenges for Antimicrobial Drug Development in the United States: Final Report [Internet]. Washington (DC): Office of the Assistant Secretary for Planning and Evaluation (ASPE); 2022. Executive Summary. Available from: <https://www.ncbi.nlm.nih.gov/books/NBK602555/> [Accessed November 2, 2025]
43. Piddock LJV, Alimi Y, Anderson J, de Felice D, Moore CE, Røttingen JA, Skinner H, Beyer P. Advancing global antibiotic research, development and access. *Nat Med.* 2024;**30**(9):2432-2443. doi: 10.1038/s41591-024-03218-w.
44. Gaibani P, Lombardo D, Bussini L, Bovo F, Munari B, Giannella M, Bartoletti M, Viale P, Lazzarotto T, Ambretti S. Epidemiology of Meropenem/Vaborbactam Resistance in KPC-Producing *Klebsiella pneumoniae* Causing Bloodstream Infections in Northern Italy, 2018. *Antibiotics (Basel).* 2021;**10**(5):536. doi: 10.3390/antibiotics10050536.
45. Woolhouse MEJ. One Health approaches to tackling antimicrobial resistance. *Sci One Health.* 2024 ;**3**:100082. doi: 10.1016/j.soh.2024.100082.
46. Ercumen A, Mertens AN, Butzin-Dozier Z, Jung DK, Ali S, Achando BS, Rao G, Hemlock C, Pickering AJ, Stewart CP, Tan ST, Grembi JA, Benjamin-Chung J, Wolfe M, Ho GG, Rahman MZ, Arnold CD, Dentz HN, Njenga SM, Meerkerk T, Chen B, Nadimpalli M, Islam MA, Hubbard AE, Null C, Unicom L, Rahman M, Colford JM Jr, Luby SP, Arnold BF, Lin A. Water, sanitation, handwashing, and nutritional interventions can reduce child antibiotic use: evidence from Bangladesh and Kenya. *Nat Commun.* 2025 ;**16**(1):556. doi: 10.1038/s41467-024-55801-x.
47. Hart A, Warren J, Wilkinson H, Schmidt W. Environmental surveillance of antimicrobial resistance (AMR), perspectives from a national environmental regulator in 2023. *Euro Surveill.* 2023;**28**(11):2200367. doi: 10.2807/1560-7917.ES.2023.28.11.2200367.
48. Milazzo A, Liu J, Multani P, Steele S, Hoon E, Chaber AL. One Health implementation: A systematic scoping review using the Quadripartite One Health Joint Plan of Action. *One Health.* 2025;**20**:101008. doi: 10.1016/j.onehlt.2025.101008.
49. Girma A. Bacteriophages as an alternative strategy for the treatment of drug-resistant bacterial infections: Current approaches and future perspectives. *Cell Surf.* 2025;**14**:100149. doi: 10.1016/j.tcsu.2025.100149.
50. Yarahmadi A, Najafiyani H, Yousefi MH, Khosravi E, Shabani E, Afkhami H, Aghaei SS. Beyond antibiotics: exploring multifaceted approaches to combat bacterial resistance in the modern era: a comprehensive review. *Front Cell Infect Microbiol.* 2025;**15**:1493915. doi: 10.3389/fcimb.2025.1493915.
51. Zou Z, Tang F, Qiao L, Wang S, Zhang H. Integrating sequencing methods with machine learning for antimicrobial susceptibility testing in pediatric infections: current advances and future insights. *Front Microbiol.* 2025;**16**:1528696. doi: 10.3389/fmicb.2025.1528696.
52. Alatawi AD, Hetta HF, Ali MAS, Ramadan YN, Alaqyli AB, Alansari WK, Aldhaheri NH, Bin Selim TA, Merdad SA, Alharbi MO, Alatawi WAH, Algammal AM. Diagnostic Innovations to Combat Antibiotic Resistance in Critical Care: Tools for Targeted Therapy and Stewardship. *Diagnostics (Basel).* 2025;**15**(17):2244. doi: 10.3390/diagnostics15172244.
53. Taghizade S, Chattu VK, Jaafaripooyan E, Kevany S. COVID-19 Pandemic as an Excellent Opportunity for Global Health Diplomacy. *Front Public Health.* 2021;**9**:655021. doi: 10.3389/fpubh.2021.655021.
54. Hitchcock MM, Markley JD, Tassone D, Kamath M, Lee KB, Greenfield A, Rittmann B, Sastry S. Collaboration on antimicrobial stewardship practices amongst university health systems, Veterans Affairs medical centers, and other affiliates: opportunities for greater harmony. *Antimicrob Steward Healthc Epidemiol.* 2023;**3**(1):e220. doi: 10.1017/ash.2023.495.
55. Singh S, Kumar M, Sarma DK, Kumawat M, Verma V, Kriti M, Tiwari R. Advancing AMR Surveillance: Confluence of One Health and Big Data Integration: Converging One Health and Big Data for AMR. *Ecohealth.* 2025;**22**(3):403-414. doi: 10.1007/s10393-025-01724-y.