



ENVIRONMENTAL STEWARDSHIP

The Role of Antimicrobials in Climate Change

By Jeanette Sams-Dodd, BSc, BScVet; and Frank Sams-Dodd, PhD, Dr med

Widespread antimicrobial use disrupts vital microbial communities that regulate Earth's biogeochemical cycles, driving antimicrobial resistance, worsening human and environmental health, and significantly contributing to climate change.

Climate change is usually attributed to the burning of fossil fuels; however, there is a general consensus that “the microbial world constitutes the life support system of the biosphere.”¹ It is bacteria and other microbes that, millions of years ago, created the unique atmosphere on Earth able to support higher life-forms, such as plants and animals. In fact, the microbes are the primary entities responsible for the biogeochemical cycles, including water, carbon, and nitrogen.¹⁻³ If the microbial world is harmed and altered substantially, it will lead to changes in the atmosphere and climate. The widespread use of antimicrobials, such as antibiotics, antifungals, antivirals, antiseptics, and disinfectants, is indeed causing such effects. A recent study⁴ estimated that one course of antibiotics, by impacting microbial soil environments, results in the release of 9.84 tonnes of CO₂ from soil storage, the equivalent of a car driving around Earth 1.5 times. This same study also found that over 7% of the current amounts of CO₂ in the atmosphere could be removed by avoiding the use of antibiotics.

Microbes are everywhere, in soil, waterways, air, and on all surfaces, eg, rocks, walls, and floors. In living organisms, they usually reside

both on and in surfaces in contact with the surroundings, eg, the skin, gut, and respiratory systems in mammals, and leaves and stems in plants. They typically organize themselves into communities, microbiomes, composed of a large variety of microbes, including bacteria, archaea, fungi, and viruses. These microbiomes are highly variable as they are characterized by all the microbial species they comprise, and they are constantly changing in response to external conditions. In these complex, collaborative communities, individual species often provide metabolites needed by other members or by their host, as exemplified in the gut-brain axis in mammals^{5,6} and the nitrogen conversion pathway in soil. The effectiveness of these vital processes depends on a balanced interspecies composition and high species diversity within the microbiome. Antimicrobials disrupt these microbiomes, changing the species composition and virulence level, and thereby the very function of the microbiome.

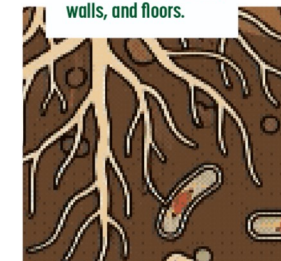
It has been the general assumption that the use of antimicrobials would sterilize the matter, leaving nothing behind and causing no lasting harm. However, new data show that microbes have several defense systems against antimicrobials, which, in many cases, leave them ineffective. This explains why health conditions involving infections often fail to respond to antimicrobials and why our efforts to control the level of infective agents, in, for example, hospital wards, fail.

The aim of this article is to provide a short introduction to these defense systems and to demonstrate that we need to integrate the diversified and ingenious ways in which microbes respond to antimicrobials into our health care approaches. Given that antimicrobials have developed and refined their defenses over millions of years, we are but novices, in comparison, lacking experience, speed, and a deeper understanding.

Bacterial Stages and Responses to Antimicrobial Stressors

Bacteria are typically viewed as single organisms floating aimlessly around, but this is a

Microbes are everywhere, in soil, waterways, air, and on all surfaces, eg, rocks, walls, and floors.



JEANETTE SAMS-DODD, BSC, BSC(VET)

With a background in veterinary medicine, she focuses on the role of microbial societies in the health of humans, animals, nature and the planet. A healthy microbial environment is essential for these systems and malfunction leads to disease, eg skin issues, cancer, deforestation, and climate change. The current focus is on the healing of severe wounds and on developing methods to create a healthy living environment in closed systems such as space crafts and submarines, where the current use of antimicrobials results in an aggressive environment causing the development of rashes, wounds, respiratory, and gastrointestinal problems, issues that are very likely to prevent deep-space exploration.

FRANK SAMS-DODD, PHD, DR.MED

With a background in biology, he focuses on the role of microbial societies in the health of humans, animals, nature and the planet. A healthy microbial environment is essential for these systems and malfunction leads to disease, eg skin issues, cancer, deforestation, and climate change. The current focus is on the healing of severe wounds and on developing methods to create a healthy living environment in closed systems such as space crafts and submarines, where the current use of antimicrobials results in an aggressive environment causing the development of rashes, wounds, respiratory, and gastrointestinal problems, issues that are very likely to prevent deep-space exploration.

highly simplistic view. Bacteria have 2 main life cycle-stages (Table): the planktonic, in which they are unattached, and the more common sessile state, in which they attach to a substrate and reside as part of a microbiome inside a collective biofilm, ie, a gelatinous layer created by the microbes, which acts like a shield to protect them, against, for example, antimicrobials. Biofilm usually has a negative connotation, but bacteria living naturally on and in our skin form biofilm, which offers us protection.⁷

Stressful conditions can induce bacteria to enter stages in which they can tolerate extreme environmental conditions. One is the persister cell, which occurs when the colony is exposed to stress, eg, antimicrobials. Here, bacteria in a certain stage of their development are induced to enter a prolonged low-metabolic state, which allows them to survive the stress.^{8,9} Once the stress is gone, they will return to their normal development. An example is the recurrence of an infection once the use of antimicrobials has ended. The spore is another stage in which the bacteria develop a protective shell and enter an inactive metabolic state, in which they can survive for thousands of years under extreme conditions.¹⁰ Bacteria can also release mobile genetic elements (MGEs), which are short sequences of DNA that, for example, in response to external conditions, can rapidly transfer traits horizontally between bacteria without the necessity of cell division.¹¹⁻¹³ Antimicrobial-resistant genes (ARGs) are a typical example of the traits carried by MGEs. This transfer can happen via cell-cell contact, or MGEs can be released into the environment and be taken up and used by other bacteria. Finally, when the cell wall breaks after a bacterium has died, MGEs and core DNA present inside the bacterium are released into the environment. This now extracellular DNA (eDNA) can remain viable for at least 3 months, during which it can be picked up, used, and replicated by other microbes.¹⁴

Bacteria respond to antimicrobials by activating their defense systems. This response includes increased development and sharing of ARGs, within and outside the microbiome.¹⁴⁻¹⁷ ARGs and other genetic solutions

that may benefit the bacteria can therefore spread swiftly, globally, and across microbial species. Additionally, genes affecting virulence are frequently positioned close to the ARGs on the MGEs, meaning that the spread of resistance will typically be accompanied by increased virulence, including accelerated mobility, and the ability to cross anatomical barriers more easily and to invade different tissue types.

Challenges in Identifying the Infective Agent

The most used diagnostic approach for bacterial infections is culturing, but this analysis has a number of downsides (see Table). The technique only focuses on a small set of preselected bacteria. Also, it is only suitable for bacteria in the planktonic state whereas the more common sessile bacteria enclosed in biofilm and anaerobic strains are only sampled to a limited extent. Furthermore, culturing is not effective for identifying persister cells, spores, MGEs or eDNA. Using proper sampling techniques and gene-based analysis methods, it is possible to identify all the strains present, but this is very rarely done in clinic. Therefore, in practice, we are unaware of what microbes are present and the defense responses we trigger.

Internal bodily regions are normally sterile, which means that the presence of a microbe can likely be linked to the infection. However, the majority of infections occur in surfaces in direct or indirect contact with the surroundings, and they are generally caused by commensals, ie, microbes that inhabit the body's natural microbiomes, such as the gut or skin. Over a thousand different species are typically present, and here infection is when the compositional balance of the microbiomes is severely disrupted. This presents a very different scenario from an internal infection, as they are typically polymicrobial, and demonstrating the presence or abundance of a specific species does not prove it to be the species causing the infection; it can easily be a desirable response by the commensals, aimed at restoring balance. Identifying the infective

TABLE. Comparison of the Sensitivity of Bacterial States and Extracellular DNA to Antimicrobials

	Bacterial state	Type of resistance	Identification using standard diagnostics	Antimicrobials
Planktonic bacteria	Free-living; metabolically fully active	Genetic ²⁹	Yes, with limitations	Effective unless resistant
Sessile bacteria	Embedded in wet or dry biofilm; metabolically fully active	Genetic and biofilm ³⁰	Limited	Highly limited efficacy
Persister cells	Phenotypic variant in a metabolically low-active, non-dividing, state with reduced cellular activity, induced by stressors ⁹	Prolonged tolerance to stressors, including antimicrobials, starvation, and low pH. Remain susceptible to same stressor after stress removal ³¹⁻³⁴	No	Not effective
Spores	Metabolically inactive state in which they can persist for many years ³⁵	Highly resistant state; tolerate heat, freezing, desiccation, radiation, chemicals, starvation	No	Not effective
Semi-autonomic DNA segments	Mobile genetic elements and extracellular DNA ³⁶	Resistance to antibiotics, disinfectants, UV light, heat ²⁷⁻²⁹	No	Not effective

agent requires understanding the role of individual species and their level of abundance in the infected area under "normal" noninfection conditions, which is practically never the case.

Impact of Antimicrobials on the Microbes, Infections, Patients, and Earth

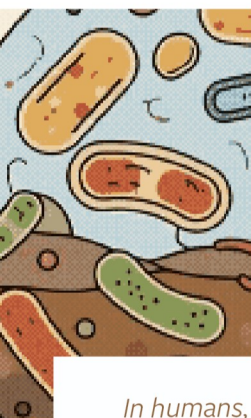
Using antimicrobials to treat an infection will, assuming the infective agent is not resistant, be able to impact bacteria in the planktonic stage. However, once the infection has become sessile and established within a protective biofilm adhering to the inside of a vessel or an organ, antimicrobials will have minimal effect, and antimicrobials will not affect persister cells, spores, MGEs, or eDNA (Table).

In internal body regions, the use of antibiotics will, when combined with the actions of the immune system, be able to treat many acute infections, because there will typically be very few infecting species. However, antibiotics are

ineffective against chronic adhering infections such as bone infection and endocarditis. They will change the infection but not eliminate it.

As the commensal community populating the body microbiomes will virtually always include at least one, and often more, resistant species, the use of antimicrobials will selectively support the resistant strains by harming the sensitive strains, thereby upsetting the microbial equilibrium established by the body. They typically induce an increased level of virulence, exacerbating the condition long-term and reinforcing the hardiness and robustness against antimicrobials. Antimicrobials, therefore, change the infection on microbiome-holding surfaces but do not eliminate it.

Antibiotics are excreted largely unmetabolized, and most antimicrobials are highly stable compounds.¹⁸ Around 50% typically escape water treatment plants and travel the water cycle, ie, from rivers, to oceans, to clouds, and return around the Earth as



In humans, epidemiological studies have found that antimicrobials are associated with increased prevalence of cancer, diabetes, asthma, obesity, immune dysfunction, depression, miscarriages, birth defects, and various developmental complications.

precipitation. Antimicrobials remaining in the water treatment plants contribute to the development and spread of antimicrobial resistance.¹⁹ These microbes will, eventually, release their antimicrobial resistance capabilities as eDNA into the extracellular realm for other microbes to pick up and use, for instance, 3 months later, thousands of miles downstream. Studies of the clearance of antimicrobial compounds in nature have usually focused on measuring their concentration in the water phase. However, data show that they often bind to organic material, where they remain active, and this typically extends their half-life considerably. Most studies of the speed at which antimicrobials are cleared from aquatic bodies, therefore, strongly underestimate their true impact.²⁰

Studies are increasingly showing long-term impacts of antimicrobials. In humans, epidemiological studies have found that antimicrobials are associated with increased prevalence of cancer, diabetes, asthma, obesity, immune dysfunction, depression, miscarriages, birth defects, and various developmental complications.^{5,6,21} In nature, antimicrobials, by damaging the microbial systems, cause deforestation and desertification by changing the structure of the soil. They further contribute to climate change by causing the release of carbon from soil storage^{22,23} and reducing the levels of CO₂ absorbed from the atmosphere into the oceans by 50% due to their effects on the microbial communities inhabiting the water-air interface.^{3,24-28}

Using Antimicrobials Optimally

The situation is, therefore, that microbes highly effectively and rapidly develop and share defenses against our antimicrobial compounds, and that these treatments and our sterilization approaches themselves lead to long-term complications in humans and animals, as well as to environmental damage. This strongly emphasizes the importance of limiting their use to conditions where they are demonstrably effective and beneficial.

Antimicrobials are frequently used to treat infections on microbiome-holding surfaces, eg,

the skin and gut, and for disinfection. However, studies show that the presence of MGEs, which carry ARGs and virulence factors, is higher on surfaces that are more frequently disinfected with antimicrobials,²⁹ showing that antimicrobial compounds, as expected, increase the hostility of the environment. This advocates for using nonantimicrobial approaches on skin and in wards, intensive care units, and other surfaces where sterility is unachievable³⁰ ie, using soap and water, to allow the area to establish a stable, diverse microbial environment that will be less aggressive.

Environmental Stewardship

Overall, the microbial world is stewarding Earth's air, water, and land resources to our benefit, and when we kill the microbes and disrupt their natural communities, we change vital biogeochemical cycles, thereby damaging the Earth. The microbes have had millions of years to develop and fine-tune their defense mechanisms. They hold vast advantages in speed and agility in raising their offense and defense capabilities. As we are learning, these capabilities are highly advanced and, when provoked, eg, by our synthetic antimicrobials, the microbes rapidly identify solutions. For that reason, antimicrobials—both antibiotics and antiseptics—should only be used when no alternative solution is available and where they demonstrably provide significant clinical benefit, in which case the one with the shortest total active half-life, when also considering metabolites, should be prioritized.

Health care is one of many sectors with an extensive use of antimicrobials in which such rules based on demonstrable necessity need to be applied. There are other sectors in which antimicrobials are used for basically nonessential purposes, eg, for textiles, clothing, paints, soaps, sanitation, plastics, feed, etc, and by banning their use, except where justified, it will be possible to assist the Earth to recover without a negative impact on our ways of living. **C**

REFERENCES ARE AVAILABLE AT
CONTAGIONLIVE.COM

PURPLE FLAG

A Rare Cutaneous Adverse Drug Reaction Associated With Linezolid

By Philip Messer, DO; Andrea Cadorette; Michael Vala, PharmD; Kenny Ha, PharmD; and Mariel del Rio-Cadorette, MD



Final Diagnosis

Purpuric drug eruption due to linezolid

History of Present Illness

A 63-year-old male patient with a history of type 2 diabetes mellitus and methicillin-resistant *Staphylococcus aureus* (MRSA) infections presented to the emergency department (ED) with a rash in the lower extremities after being discharged 3 days prior on linezolid for MRSA prepatellar bursitis.

During his previous admission, he was diagnosed with left lower extremity cellulitis and septic prepatellar bursitis by orthopedic surgery. A needle aspiration of the bursa was performed, and cultures grew MRSA sensitive to vancomycin, daptomycin, linezolid, trimethoprim-sulfamethoxazole, and tetracycline. Septic arthritis was excluded by physical exam and a CT scan of the lower extremity. The infectious disease service was consulted, and he was treated with intravenous vancomycin in the hospital for 5 days. The erythema, swelling, and pain of his knee improved slowly. He was discharged with oral linezolid 600 mg twice daily to complete 14 days of treatment. His platelet counts were normal at the time of discharge.

After discharge, the patient noticed small red purpura on both legs 2 days after beginning linezolid. The purpura coalesced into red patches in both lower extremities, worse on the left side (Figures 1 and 2), with associated leg swelling, so the patient returned to the ED. He denied chills, rigors, fever, arthralgias, myalgias, anorexia, malaise, or confusion.

Medical History

His medical history was remarkable for type 2 diabetes mellitus with neuropathy and a history of diabetic foot infections. He had multiple previous MRSA infections, including a groin abscess, heel osteomyelitis, and

bacteremia. He also had a history of gastroesophageal reflux disease, hypertension, obesity, obstructive sleep apnea, chronic kidney disease stage IIIA, paroxysmal atrial fibrillation on anticoagulation, and depression.

Key Medications

His medications at home included apixaban, atorvastatin, duloxetine, finasteride, gabapentin, insulin glargine, insulin lispro, metoprolol, nortriptyline, and topiramate.

Allergies

He reported allergies to amoxicillin-clavulanate, oxycodone, and shellfish.

Epidemiological History

He is married and lives with his wife, with whom he is monogamous. He is a former cigarette smoker and has no history of alcohol or drug use. He had no recent travel or exposure to insects or pets.

Physical Examination

His vitals were notable for normothermia with a temperature of 35.7 °C and were otherwise unremarkable. A nonpalpable purpuric rash was observed on his bilateral lower extremities, including the lateral left leg from mid thigh to mid calf and the right ankle, with swelling of the left leg. His left knee bursitis had improved. The rash appeared worse in an area of previous scarring on the left lateral leg and knee (Figures 1 and 2).

Studies

His laboratory studies were notable for creatinine 1.97 mg/dL (normal range, 0.7-1.2 mg/dL), which was at his baseline; glucose 143 mg/dL (normal range 70-115 mg/dL); lactic acid 1.2 mmol/L (normal range 0.5-2.2 mmol/L); leukocytes 8.9K/