Antibiotic use and population ecology: How you can reduce your “resistance footprint”

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A 8-year-old child presents with a 4-day history of a nonproductive cough and a temperature of 38°C. Her chest is clear, except for some wheezing. She has no tachypnea or tachycardia. The child’s father is deeply concerned. What would be the harm of prescribing antibiotics for acute bronchitis? Even when antimicrobials are used wisely, they create selective pressure that may increase the prevalence of antibiotic-resistant organisms. Every antibiotic prescription represents a balance of benefits and risks, both to the individual and the population. The risks of antimicrobial therapy to the patient are well-known and include hypersensitivity, drug interaction and disruption of normal flora. However, not all of the risks associated with antibiotic use may be directly experienced by the patient receiving treatment.

The metaphor “tragedy of the commons” describes an event in which individuals acting locally to benefit themselves inadvertently contribute to catastrophe at the ecological level. This metaphor originates from a description of the tragic collective outcome incurred by farmers who individually contributed to over-grazing of common lands. Indeed, this metaphor can be applied to antimicrobial use. What is the parallel problem with prescribing antibiotics for acute bronchitis in a previously healthy child? Just as the farmer who attempts to benefit from adding 1 more animal to the pasture does not consider the impending catastrophe of complete consumption of the pasture, the prescribing physician or patient who attempts to benefit from antibiotic therapy likely does not consider the impending catastrophe of antibiotic resistance. Examination of the population perspective of antibiotic resistance requires consideration of how our individual actions sum over a population to create a measurable effect on antibiotic resistance. Our collective prescriptions constitute an ecological problem that may reduce the success of future therapy. In this article, we provide an overview of some of the abundant literature that documents this effect.

During a clinical encounter is not the easiest time to stand back and consider our relationship with microorganisms in the broadest sense, along the evolutionary time scale and across the entire population. Microbes and multicellular animals have engaged in a co-evolutionary tango for hundreds of millions of years. Natural selection is the engine that created our current pageant of diverse and complex life on earth. This dance has refined our immune system, led to the establishment of highly beneficial normal flora and caused remarkable adaptation of man to microbe and vice versa. The process of natural selection also ensures that if a substance affects an organism’s chance of survival, those best able to reproduce and thrive in the presence of the substance will propagate, increase in prevalence and become dominant in the microbial population over time. Chemical competition among microorganisms has occurred for hundreds of millions of years and has resulted in the production of antimicrobial compounds long before humans walked the earth. Because many antimicrobials in use by humans were derived from these substances, bacteria were endowed with rich genetic machinery to resist the effects of antibiotics long before Domagk and Fleming discovered sulfa drugs and penicillin in the first half of the twentieth century. Many of these genes can be transferred horizontally between species and are important evolutionary elements in their own right. Such mechanisms are well documented in the laboratory where it is possible to observe natural selection at the level of the microbial population.

Physicians depend on antibiotics for many good reasons, even while knowing that their use carries a long-term evolutionary cost. Drawing a parallel to carbon emissions and global warming, we may well ask how we may reduce our “resistance footprint” without causing harm by withholding

Key points

- Genes that determine resistance to antibiotics were widespread in nature even before humans discovered the use of these drugs.
- There is ecological, observational and experimental evidence to suggest that populations with lower rates of antibiotic use will generally experience a lower burden of colonization by antibiotic-resistant organisms.
- Physicians can contribute to decreased antibiotic prescribing and prevalence of resistant organisms by carefully following evidence-based guidelines.
- The medical profession needs to engage governments to assist in striking the best balance between controlling antibiotic use through formulary restrictions and making antibiotics available to those who can truly benefit.

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antibiotics from those that need them. There is plenty of room for action based on current knowledge, but we are going to have to make a concerted effort to change our own habits.

Population-level effect of prescribing on antibiotic resistance

In recent years, there has been a vast improvement in our ability to measure antibiotic use at the population level, track trends in resistance and identify the relations between use, resistance and outcomes of disease.10,11 As an example, Figure 1 illustrates the remarkable concordance between the increasing rate of fluoroquinolone use and the increasing rate of ciprofloxacin resistance among uropathogens in British Columbia. Table 1 summarizes the studies supporting the hypothesis that increasing levels of population use of many drugs and drug classes corresponds with increasing resistance among pathogens. These papers were identified by searching MEDLINE for the keywords “antibiotic,” “resistance” and “utilization” and by screening the results for articles that focused on observations of association in entire populations rather than on individual institutions. Ecological studies, which measure both resistance and use at the population level rather than the individual or institutional level, demonstrate that the spatial and temporal distribution of resistance is strongly associated with the rate of use of specific classes of antibiotics in human populations.12–20 These studies show a clear association between the use of penicillins, macrolides and fluoroquinolones and drug resistance in common human pathogens like *Streptococcus pneumoniae* and between the use of fluoroquinolones and resistance in *Escherichia coli*.

There are also increasing data from individual-level cross-sectional studies, cohort studies and randomized controlled trials that strengthen the causal inference of this relation (Table 1).12,21,22 The relation between antibiotic use and resistance satisfies a logical temporal sequence, has been demonstrated in individuals and has even been demonstrated in a randomized controlled trial.21 An interesting finding of such studies is that not all drugs of a particular class are equal with respect to their potency in selecting for resistance.23 For example, azithromycin has a long terminal elimination half-life.24 Although this has proven convenient from a dosing point of view, the downside is that there is a great deal of time at the end of a course of therapy when the drug is still present in the body at subinhibitory concentrations. Several studies indicate that azithromycin is far more likely to select for macrolide resistance than drugs of the same class with shorter half-lives.19,21

Finally, there is also strong ecological evidence that deliberate efforts to control antimicrobial use can have beneficial effects on the trajectory of resistance.12,18,23 It appears that populations that use antibiotics prudently are benefiting.

The question of whether the relation between antibiotic use and resistance at the population level is causal is not straightforward or easily answered. However, application of the 9 Bradford Hill criteria for the evaluation of environmental factors and relation to disease strongly supports causality.26 There is a reasonably strong statistical association and consistency across populations. There is relative specificity of the effect, a logical temporal sequence of exposure followed by effect, and evidence of a biological gradient or a dose–response relation. This relation is entirely plausible based on known biological mechanisms and is coherent with other observed effects of evolution. Probably most important, there is experimental evidence both in laboratories and in groups of humans that helps assure that our view is not entirely obscured by unknown confounders.

Reasonable goals in antimicrobial therapy

Most bacteria are neutral or beneficial from the standpoint of our health. Even those with the potential to be pathogens do not always cause disease. Yet all bacteria are able to mutate or acquire new genetic material. If the resulting genetic modification is heavily favoured by the abundant presence of antimicrobials, our ability to treat disease will be impaired. Some drug advertising and much of our medical training wrongly conceptualizes infectious disease treatment as a “war of extermination.” In reality, the purpose of antimicrobial therapy is rarely, if ever, the complete eradication or elimination of microbes from the patient or their environment. Because we are fated to co-exist with microbes, the purpose of antimicrobial therapy must be more appropriately and realistically considered to be the prevention or favourable modification of the course of infectious disease.

Gaps, complexities and confounders

There remain key gaps in our knowledge about population-level effects of antibiotic prescribing on resistance. Most existing studies focus only on widely cultured human pathogens where data are available from diagnostic laboratories in sufficient quantity to allow statistical inference. There is a shortage of observations in large populations for some important
categories of human pathogens. However, data available from hospitals and institutions, although not reflective of the general population in all respects, strongly broaden the argument that our practice affects our future with respect to a wide array of antibiotic-resistant organisms.27–29

Just as the natural relations between microorganisms, animals, humans and the environment are exquisitely complicated, it is expected that the effects of wide-scale, man-made distribution of antimicrobials on microorganisms will be complicated. We are just beginning to understand the contribution to evolution of horizontally transferable genetic elements. Because these movable pieces of DNA may confer resistance to more than 1 class of antibiotic, the use of 1 class of antimicrobials may select for resistance to another, and resistance could conceivably increase even with declining use of some classes of drugs.30

Host factors will also prove important. The density of human populations is an independent predictor of the prevalence of antibiotic resistance.31 With or without antimicrobial selection, some newly introduced microbial clones are very fit to spread through human populations. Although antibiotic use may contribute to an environment that favours the spread of community-associated methicillin-resistant Staphylococcus aureus in North America, the strains also possess other attributes that facilitate their spread and pathogenicity.32

Although it is a legitimate focus for physicians, use of antimicrobials by humans is not the only source of selective pressure. The massive use of antibiotics for both therapeutic purposes and growth promotion in the agriculture and food industries and in veterinary medicine are reasons for concern.33 Such use could result in colonization of humans by re-

| Table 1: Evidence that antibiotic use at the population level is associated with antibiotic resistance |
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| **Level; study design** | **What the studies show** | **Example** |
| Community level | Antibiotic use is associated with the prevalence of carriage of resistant organisms in the community. | Resistance of Streptococcus pneumoniae to penicillin was correlated with the use of β-lactam antibiotics and macrolides across Europe.12 |
| Ecological studies | | In Europe, the prevalences of resistance to macrolides and β-lactams in S. pneumoniae, macrolide resistance in Streptococcus pyogenes and resistance to quinolones and co-trimoxazole in Escherichia coli were significantly correlated with the use of relevant classes of antibiotics.13 |
| | Antibiotic use (within the past year) and population density were independent predictors of S. pneumoniae antibiotic resistance in Belgium.14 |
| | Antibiotic use correlates with carriage of resistant S. pneumoniae in Iceland.15 |
| | Associations are most consistent if a 2-year lag is observed between antibiotic use and resistance trends (multiple countries).16 |
| | Decreased sales of β-lactams in the United Kingdom were followed by a decrease in penicillin-resistant pneumococci. A similar relation was not observed with decreased rates of macrolide prescription.17 |
| | Pneumococcal strains resistant to multiple antibiotics disappeared from areas with the lowest rates of antimicrobial use but prevalence remained unchanged in the area with the highest rates of use. The prevalence of erythromycin-resistant strains increased in association with increased use of macrolides, especially azithromycin.18 |
| | The prevalence of fluoroquinolone-resistant E. coli was associated with the rate of fluoroquinolone use in Spain.19 |
| | The prevalence of fluoroquinolone-resistant E. coli in hospitals was associated with the rate of fluoroquinolone use in the surrounding community.20 |
| Individual level | Use of antibiotics by individuals is associated with a higher rate of carriage of resistant organisms. | Individual-level antibiotic use correlates with the carriage of antibiotic-resistant S. pneumoniae in Iceland.13 |
| Cross-sectional study | Use of antibiotics by individuals is associated with a higher risk of subsequent infection by resistant strains. | Use of longer-acting macrolides was associated with increasing resistance to macrolides.21 |
| Prospective cohort study | Use of antibiotics by individuals is associated with a higher risk of subsequent colonization by resistant strains. | Use of macrolides increased the carriage of macrolide-resistant S. pneumoniae by 50%. The effect was larger with longer-acting macrolides than with shorter-acting macrolides.22 |
| Randomized controlled trial | Use of antibiotics by individuals is associated with a higher risk of subsequent colonization by a resistant strain. | |
sistant organisms through a variety of routes. Residual antibiotics in meat eaten by humans may lead to selection for resistance among colonizing organisms in the human host. Most food producers in developed countries observe a wash-out period to reduce this risk. Selection of resistant organisms in food animals and pets can lead to colonization of humans with the same strain.36 Resistant commensal organisms may not carry disease but they may carry horizontally transferable genetic elements that could transfer resistance to other human commensal organisms or pathogens. Finally, excess antibiotics spilt into the environment from the agriculture and food industries could lead to selection of resistant organisms in the environment that could subsequently spread disease or resistance to humans.

What are the major reasons for prescribing antibiotics?

None of us would feel very comfortable managing pneumonia, pyelonephritis or spreading cellulitis without antibiotics. Fortunately, prescriptions for these serious infections account for a small fraction of overall antibiotic use. The majority of outpatient prescribing is for the acute respiratory conditions pharyngitis, sinusitis and acute bronchitis, as well as for otitis media.35 These syndromes are almost entirely caused by viruses. Several systematic reviews and meta-analyses have concluded that there is little or no benefit of antibiotics to these conditions.36 Rates of antibiotic prescriptions for these indications need to drop. A key example of early success in this area has been that, in recent years, it has been shown that most cases of otitis media are adequately managed symptomatically, with consideration of antibiotic prescription only with failure to improve over 48 hours.17,38 Pediatricians and family physicians have shown that by using such an approach, antibiotic prescriptions can safely be reduced.35

| Table 2: Impact of population-level interventions on antimicrobial use and resistance |
|------------------|-----------------|--------------------------|------------------|
| Intervention                  | Study design                                      | Result                                      | Example                                      |
| Public education              | • Polling before and after a public campaign       | • Effects may be small from public education alone | • Public education initiative in the United Kingdom did not change knowledge very much.24 |
| Physician and other health professional education | • Ecological evaluation of educational programs or professional education | • Programs can be associated with moderate reduction in prescribing and may be associated with lower prevalence of antibiotic-resistant organisms | • Various interventions following a national consensus conference were associated with declining use in Canada after 1997.26 |
|                              | • Noncontrolled before- after community interventions. |                                        | • Modest impact of professional education efforts at reducing prescribing.44,51 |
|                              | • Controlled trial of physician education intervention |                                        | • Reduced prescribing in areas with education for prescription reduction (one study also associated this with lower carriage of resistant pneumococci).45,46 |
| Mixed public and professional education | • Cluster randomized trial                           | • Programs can be associated with moderate reduction in prescribing | • Moderate reductions in prescribing in study communities in the United States.28 |
| Change to drug formulary      | • Ecological time series evaluation of formulary changes | • Policy change is associated with change in the rate of use and this may be associated with moderating the prevalence of antibiotic-resistant organisms | • Policy change in a provincial formulary reduced fluoroquinolone use with no excess of admissions to hospital.49 |
|                              |                                                |                                        | • Policy changes in a provincial formulary affected clarithromycin use.54 |
|                              |                                                |                                        | • Control of population use was associated with relatively low and stable rates of resistance in pneumococci.25 |
Strategies to reduce your antibiotic resistance footprint

With the availability of many detailed guidelines for the treatment of specific infections, it is sometimes difficult to identify actions to take at the practice level. A few simple suggestions for reducing unnecessary antimicrobial use are presented in Box 1. In general, a higher threshold for antimicrobial prescriptions should be developed. Such actions do not lie outside of the mainstream literature and are compatible with guidelines for the management of individual infections.46-49

Physicians may be concerned that avoidance of antibiotic use could lead to more infectious complications or hospital admissions. Indeed, when surveyed, some physicians and patients do not agree with strategies such as delayed prescribing.41 This is why it is critical to construct guidelines for use and nonuse based on rigorous reviews of the evidence. However, there are some reassuring data available. Restriction of fluoroquinolones on the prescription drug formulary for Ontario was not associated with a net change in hospital admissions, and the rate of hospital admissions for gastrointestinal complications was actually decreased.44 Re-evaluation of patients during the course of an infection to look for improvement or deterioration is an important aspect of medical care as we improve the judicious use of antimicrobials.

Physicians and the regulation of antibiotics

Evaluations of efforts to curb prescribing and resistance at population level are summarized in Table 2. These articles were identified by searching MEDLINE for the keywords “antibiotic,” “resistance” and “utilization,” and by screening the results for articles that focused on interventions in entire populations rather than individual institutions. Reductions in antibiotic use through such programs, although measurable, are modest.45-50 One review concluded that there was some benefit from social marketing, practice guidelines, authorization systems and peer review with feedback, and speculated that online systems that provide clinical information, structured order entry, and decision support may be the most promising approach.51

A book developed as part of the “Do Bugs Need Drugs?” program in British Columbia and Alberta represent important initiatives to create understanding among doctors and patients about the importance of reducing antimicrobial misuse.52-53 Evaluation of the approach suggested by this program is underway.

Conclusion

It is becoming clear that the greatest successes in changing patterns of antibiotic use have resulted from administrative decisions, such as changes to drug formularies at the institutional level or changes in reimbursement from provincial drug plans.44,45 Such approaches will prove necessary to adequately steward the use of the drugs that we have. As a profession, we would do well to thoroughly engage in discussions and processes in this area as a means of assuring the best interests of our patients while reducing our collective resistance footprint.

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