

Impact of Antibiotic Resistance on Treatment of Pneumococcal Disease in Ethiopia: An Agent-Based Modeling Simulation

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Abstract. Antimicrobial resistance (AMR) is a growing threat to global health. Although AMR endangers continued effectiveness of antibiotics, the impact of AMR has been poorly estimated in low-income countries. This study sought to quantify the effect of AMR on treatments for pediatric pneumococcal disease in Ethiopia. We developed the DREAMR (Dynamic Representation of the Economics of AMR) model that simulate children younger than 5 years who acquire pneumococcal disease (pneumonia, meningitis, and acute otitis media) and seek treatment from various health facilities in Ethiopia over a year. We examined the AMR levels of three antibiotics (penicillin, amoxicillin, and ceftriaxone), treatment failures, and attributable deaths. We used the cost-of-illness method to assess the resulting economic impact of AMR from a societal perspective by estimating the direct and indirect treatment costs and productivity losses. Findings showed that AMR against antibiotics that were used to treat pneumococcal disease led to 195,763 treatment failures per year, which contributed to 2,925 child deaths annually in Ethiopia. Antimicrobial resistance resulted in a first-line treatment failure rate of 29.4%. In 1 year, the proportion of nonsusceptible *Streptococcus pneumoniae* bacteria increased by 2.1% and 0.5% for amoxicillin and penicillin, and reduced by 0.3% for less commonly used ceftriaxone. Annual costs of AMR to treat pneumococcal disease were around US\$15.8 million, including US\$3.3 million for ineffective first-line treatments, US\$3.7 million for second-line treatments, and US\$8.9 million for long-term productivity losses. Antibiotic stewardship to reduce misuse and overuse of antibiotics is essential to maintain the effectiveness of antibiotics, and lessen the health and economic burden of AMR.

INTRODUCTION

Antimicrobial resistance (AMR) poses a global public health threat by diminishing the effectiveness of existing antimicrobials, leaving individuals prone to prolonged hospitalization and mortality from simple bacterial infections.^{1,2} Antimicrobial resistance is the mechanism by which microbes such as bacteria deactivate the efficacy of antimicrobials from destroying them and stopping their growth.³ Bacteria become resistant through various mechanisms, including producing destructive enzymes that neutralize antimicrobials, modifying antimicrobial targets by mutation so that drugs cannot recognize them, or removing antimicrobial agents by pumping them out of cells.⁴ Resistant bacteria can also prevent antimicrobials from entering bacterial cells by modifying the outer cell membrane or creating bypasses that allow bacteria to function without the enzymes targeted by antimicrobials.⁴ Resistance to all first-line and last-resort antimicrobials is increasing globally, and only a few antimicrobials have been developed in recent decades, resulting in an overall decline in the total portfolio of antimicrobial effectiveness.^{5–7}

The global costs of AMR are estimated to increase to US\$100 trillion annually by 2050.⁸ Antimicrobial resistance results in longer treatment duration, greater side effects from second- or third-line treatment, higher mortality and morbidity, more treatment costs, as well as income losses.⁹ In the United States, antimicrobial-resistant bacterial pathogens are responsible for

more than two million infections and 23,000 deaths each year at a direct cost of US\$20 billion, and additional productivity losses of US\$35 billion.¹⁰ In Europe, in 2007, 25,000 deaths were attributable to resistant infections, costing €1.5 billion annually in direct and indirect costs, according to an estimate from the European Medicines Agency and European Center for Disease Prevention and Control.¹¹ Although efforts have been made to estimate the impact of AMR, most studies have focused on high-income countries.¹² Similar estimates of the impact of AMR are not currently available in low- and middle-income countries, although these countries are experiencing the greatest increase in antimicrobial use.⁵

The World Health Organization (WHO) lists *Streptococcus pneumoniae* as a community-acquired infection of high global concern for resistance.¹³ *Streptococcus pneumoniae* is the pathogen that causes pneumococcal infections resulting in several diseases, such as acute otitis media, pneumonia, and meningitis. *Streptococcus pneumoniae* is known to cause at least 18% of severe pneumonia episodes and 33% of pneumonia deaths worldwide.¹⁴ Understanding the influence of AMR on pneumococcal infections is particularly important in countries with high rates of pneumonia and child deaths. Ethiopia is among the top five countries globally with the highest number of child deaths. Estimates suggest that between 33,000 and 37,000 Ethiopian children younger than age 5 die annually from pneumonia.¹⁴ We examined the impact of AMR on treatments for pediatric pneumococcal infections in Ethiopia.

Agent-based modeling (ABM), a type of individual-based model, can aid in simulating complex interactions among agents and assess their effects on the system as a whole.^{15,16} In an ABM, agents such as bacteria and humans follow predetermined rules based on their heterogenous characteristics

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and the environment. Some characteristics change over time, whereas others do not, resulting in dynamic interactions between agents and the environment to produce aggregate results over a period.¹⁵ ABM can provide better insights into the dynamics of infectious diseases compared to previous studies which primarily used compartmental approaches to model AMR.^{12,16} In addition, most models that have examined the impact of AMR were deterministic and did not incorporate an economic perspective. We aimed to develop a stochastic ABM with humans and bacterial agents to examine the broader health and economic effect of AMR on treatment for pneumococcal disease.¹²

MATERIALS AND METHODS

We developed the DREAMR (Dynamic Representation of the Economics of AMR) model, an ABM with two interactive sub-models for bacteria and humans. Creating two submodels facilitated dynamic interactions between humans and bacteria. In the human submodel, children become infected by *S. pneumoniae*, seek care at varying health facilities, and use antibiotic treatment or do not seek care. When antibiotics are used, bacterial agents survive or die, and replicate based on the magnitude of antibiotic exposure. This results in a change in the proportion of resistant strains in the bacterial population also known as the AMR pattern. The AMR pattern subsequently affects treatment outcomes of pneumococcal diseases among human agents, where treatment failures increase as the proportion of resistant strains increase. As a consequence, adverse health outcomes and costs of treatment increase because of prolonged treatment duration and use of second-line antibiotics. We used NetLogo 6.0.2 for our simulation, a freely available and widely used educational software with a multi-agent programmable modeling environment.¹⁷ We

estimated the annual health and economic impact of AMR on treatment of pediatric pneumococcal infections in Ethiopia.

Bacteria submodel. The DREAMR bacteria submodel included two types of bacteria: resistant strains and susceptible strains (Figure 1). The initial proportions of each type of bacteria were set based on resistance patterns for three antimicrobials (amoxicillin, penicillin, and ceftriaxone) that are commonly used to treat pneumococcal diseases in Ethiopia.¹⁸ The model simulated 5,000 bacterial agents, including both resistant and susceptible strains for each antibiotic. Each agent represented 0.02% of bacteria in the entire bacterial reservoir. The bacterial reservoir consisted of *S. pneumoniae* that colonized on either symptomatic or asymptomatic individuals, where bacterial agents could move randomly over the entire bacteria space. This represented disease transmission, where bacteria can move from one individual to another.

Every bacterial agent was categorized as either resistant or susceptible against each antibiotic. Individual bacterium was also assigned a minimum inhibitory concentration (MIC) value based on its resistant/susceptible characteristic, where resistant bacteria were more likely to be assigned higher MIC values. The MIC values were obtained from the gamma distribution to represent the highly right-skewed nature of MIC values, with variance obtained from the literature.^{19,20} The method of moments approach was applied to derive parameters needed for gamma distributions.²¹ The 90th percentile of MIC values in the gamma distribution for susceptible strains was set to be equal to the breakpoint for susceptible strains determined by the Clinical Laboratory Standards Institute (CLSI). The 90th percentile of MIC values for resistant strains was set to the breakpoint of intermediate strains by combining intermediate and resistant strains from CLSI as a resistant category.²⁰

We simulated pharmacokinetics based on currently approved dosages for the three antibiotics and evaluated the

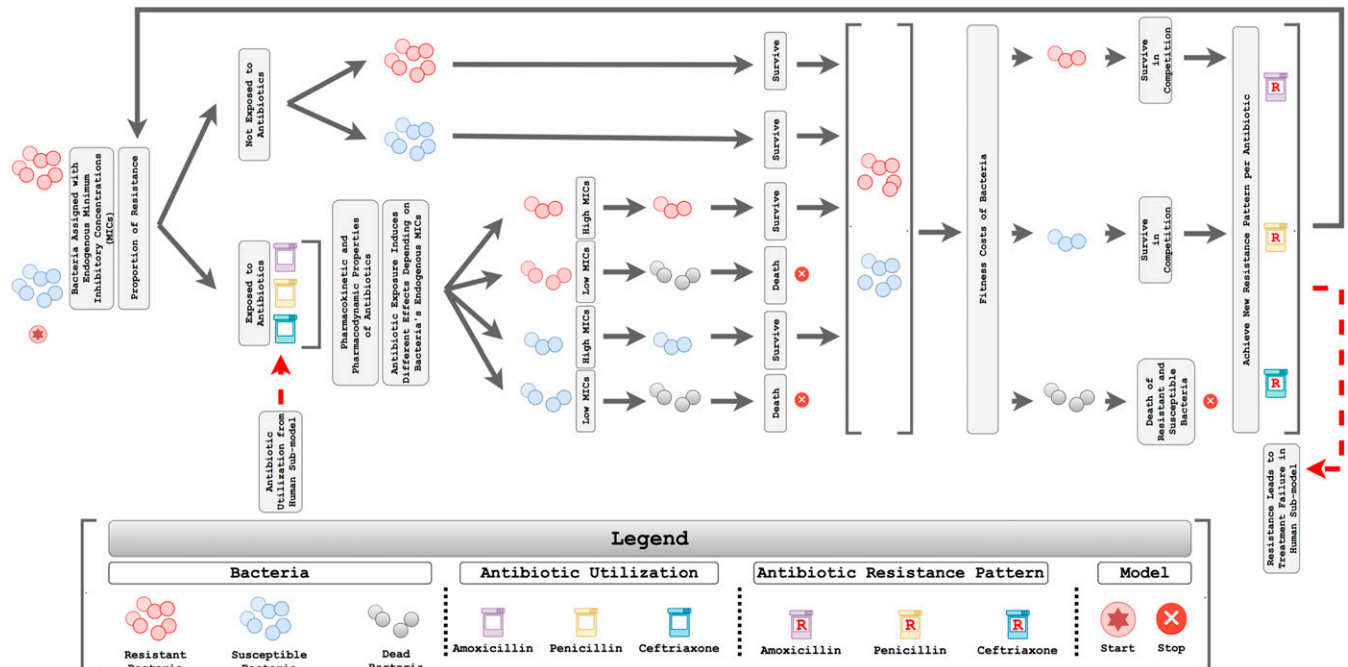


FIGURE 1. Conceptual framework of the bacteria submodel to estimate the accumulation of antimicrobial resistance. This figure appears in color at www.ajtmh.org.

resulting pharmacodynamics to determine whether each bacterium will die under antibiotic exposure (see Supplemental Appendix). Pharmacokinetic parameters, including the volume of distribution, total body clearance, and elimination rate constant were retrieved from the literature and product information for each antibiotic.^{22–25} The probability that adequate antibiotic exposure was achieved depended on defined daily doses (DDD) and the proportion of children colonized with *S. pneumoniae*.²⁶ DDDs were derived from the human submodel, which represented the proportion of children using antibiotics. Because not every individual carries *S. pneumoniae*, the model divided DDDs by the proportion of humans colonized to estimate the probability that bacterial agents would be exposed to antibiotics. As a result, larger DDDs led to a greater propensity for bacteria to be exposed to antibiotics.²⁷ Antibiotic utilization was simulated based on common recommended dosages and intervals.^{28,29}

Because the three beta-lactam antibiotics are time dependent, we simulated the effectiveness of antibiotics based on the percentage of time the exposed antibiotic concentration is above the MIC, where time is highly correlated with the overall dose, dosing frequency, and other pharmacokinetic characteristics.^{30,31} Bacteria would die if the percentage of time the antibiotic concentration is above its MIC value became larger than the threshold, under the assumption that all bacteria are susceptible to the antibiotic if adequate exposure is achieved.^{30,32,33} Susceptible strains of bacteria were more likely to be killed compared with resistant strains because of a lower MIC distribution.

The model also included fitness costs of resistant bacteria to counter the AMR growth.^{12,34} Fitness of resistant bacteria depended on several factors, such as host immune status, the environment in which the bacteria are growing, and prior drug pressure.³⁵ As the literature suggests that resistant strains are often less fit than susceptible strains when not facing antibiotic pressure, resistant bacteria in the model faced a lower probability of survival when competing with susceptible strains.³⁴ Survived bacteria, either resistant or susceptible strains, subsequently had an identical chance to reproduce regardless of their MIC values, until the total number of bacteria in the population reached 5,000 (100%) again to obtain a new resistant pattern. The rate of change in the proportion of resistant bacteria was associated with antibiotic pressure. We examined the AMR patterns for each antibiotic, which were dynamically updated over time in accordance with antibiotic exposure.

Human submodel. We also developed the DREAMR human submodel, where we simulated 8,000 child agents between 0 and 5 years of age with incidence of pneumococcal infections resulting in pneumonia, meningitis, and acute otitis media (Figure 2). Each child agent in the model represented 10 children, where the study modeled a total of 80,000 children over a 1-year horizon. A similar method was also used in other ABMs to overcome the computational burden.³⁶ Every child agent in the model was assigned an immunization status for receiving the pneumococcal conjugate vaccine (PCV) based on the WHO and United Nations Children's Fund (UNICEF) immunization coverage estimates in 2017 for Ethiopia.³⁷ Daily pneumococcal disease incidence rates were applied based on the child agent's PCV immunization status and vaccine efficacy.^{26,38–40} Disease incidence also took into consideration the effects of herd immunity, where immunization can

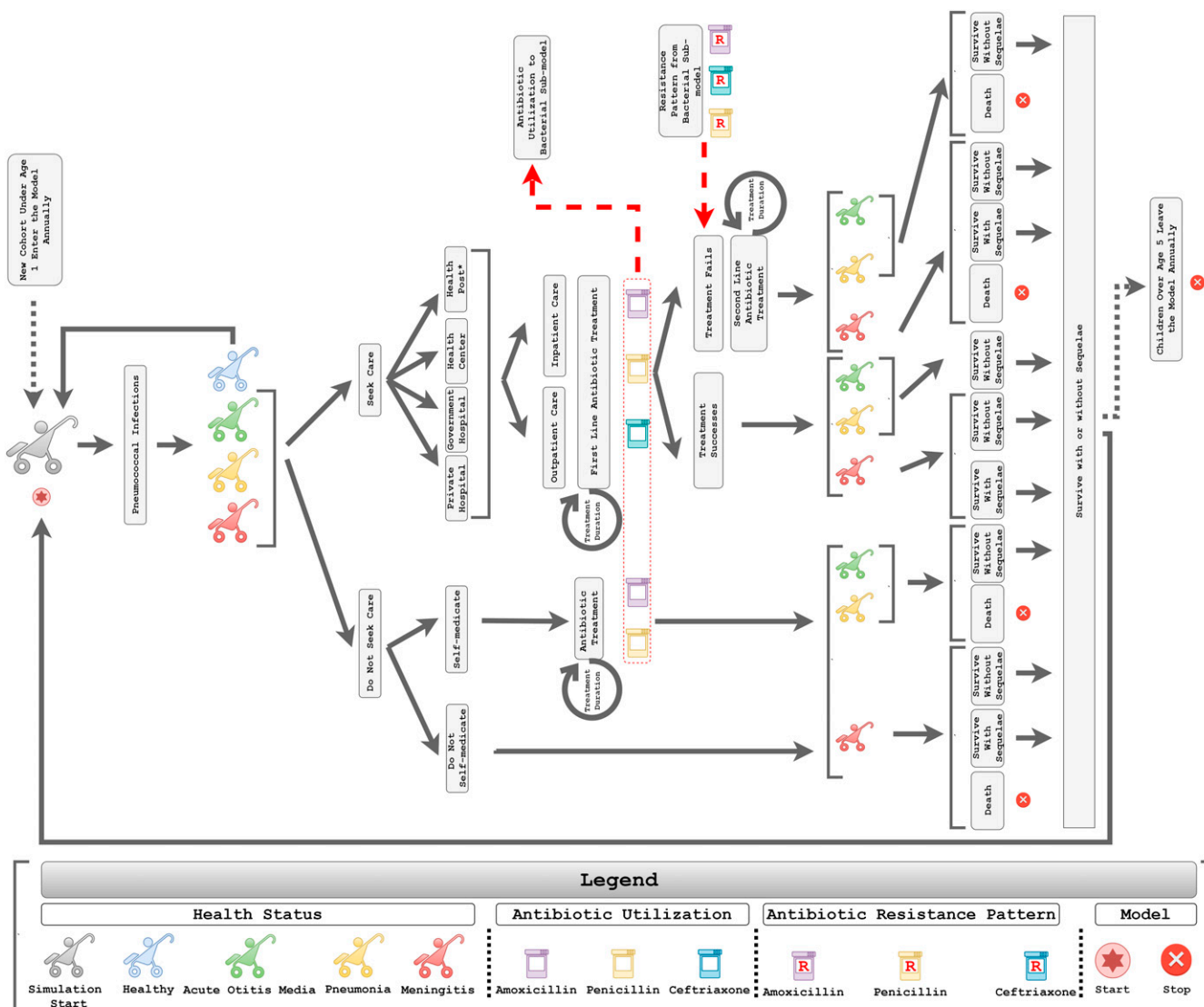
protect unvaccinated populations by preventing disease transmission when immunization coverage reaches a high threshold.^{41,42} This study simulated herd immunity at the vaccine coverage rate of 2017 (i.e., 68%), where lower disease incidence rates were applied to unvaccinated populations to simulate the indirect effect of immunization.⁴³

Care-seeking and treatment of pneumococcal disease in children was modeled over 1 year. Specifically, caregivers of children with pneumococcal infections chose whether or not to seek care from healthcare facilities based on care-seeking rates from the Ethiopian Demographic and Health Survey.⁴⁴ Children who did not receive treatment from healthcare facilities would either receive self-medication or remain untreated. Untreated children faced a greater propensity to develop adverse health outcomes (i.e., disability and death).⁴⁵ Children who sought health care received treatment from one of the following facilities: 1) health post, 2) health center, 3) government hospital, or 4) private hospital/clinic.⁴⁶ All children with acute otitis media were treated as outpatients, whereas all meningitis cases were treated as inpatients.^{47,48} Hospitalization rates from the literature were applied to children with pneumococcal pneumonia who sought care at the health center, government hospital, or private hospital/clinic.⁴⁶ At each health facility, we simulated child agents receiving first-line antibiotic treatment in an outpatient or inpatient setting. Three antibiotics were commonly used as first-line treatment for acute otitis media, pneumonia, and meningitis in Ethiopia: 1) amoxicillin, 2) penicillin, and 3) ceftriaxone.⁴⁹ In outpatient/self-medication scenarios, the model assumed that patients received antibiotics that could be given through oral route (i.e., amoxicillin and penicillin). Antibiotic regimens and treatment durations were extracted from the literature.^{28,29,50,51} The study also included the effects of noncompliance, where noncompliant individuals who received antibiotics from a healthcare facility or self-medication faced a 2-fold higher risk of adverse health outcomes.⁵²

The proportion of child agents using antibiotics was derived on a daily basis to estimate the magnitude of antibiotic utilization in the bacteria submodel. When exposed to antibiotics, the chance of bacterial survival depended on the antibiotic dose and dosing frequency, pharmacokinetic characteristics, the MIC values of bacteria, and fitness costs. In addition, the model assumed a 75% effective dose in the bacteria submodel when bacteria encountered antibiotic exposure caused by non-compliant patients. The proportion of antibiotic resistance in the bacteria submodel subsequently affected the rates of treatment failures observed among child agents receiving the third day of first-line treatment in the human submodel.⁵³

Child agents encountered first-line treatment failure if they used an antibiotic and had a susceptibility value (between 0 and 1 from a uniform distribution) lower than that of the same antibiotic's AMR pattern. Child agents with treatment failure would then be switched to second-line therapy, and the overall treatment duration was prolonged. Child agents with illness either died or recovered from the disease episode based on overall case fatality rates.³⁹ Child agents who survived meningitis faced a chance of developing neurological sequelae.⁵⁰

Outcomes and uncertainty analyses. We estimated the annual change in the AMR pattern, measuring the proportion of bacterial strains for each antibiotic. We simulated the number and percentage of treatment failures for pneumonia,



Note: * Every patient in health post will receive outpatient treatment.

FIGURE 2. Conceptual framework of the human submodel to estimate the health and economic impact of antimicrobial resistance. This figure appears in color at www.ajtmh.org.

meningitis, and acute otitis media for children under 5 in Ethiopia.⁵⁴ Other predicted health outcomes included average DDDs, cumulative incidence of disease episodes, and annual numbers of death and disability due to pneumococcal disease in Ethiopia. We also assessed the number of deaths due to treatment failures. All outcomes are based on an average of 10,000 simulation runs in the model.

In addition to health effects, we also assessed the economic impact using the cost-of-illness method taking a societal perspective, where we estimated direct medical costs, direct nonmedical costs, productivity losses for caregivers, and productivity losses due to death/disability.^{55,56} Direct medical costs combined costs of 1) registration/consultation, 2) laboratory diagnosis, 3) medicines and supplies, 4) hospital bed, and 5) traditional healer visits.⁴⁶ Direct nonmedical costs included costs of transportation and others such as food and lodging. Unit costs for each health facility were obtained from the literature.⁴⁶ We also estimated the productivity losses for caregivers and productivity losses due to death and disability

based on Ethiopia's gross domestic product per capita using a human capital approach.^{57,58} A disability-adjusted life year weight was applied to productivity losses for children living with disability due to meningitis.⁵⁹ All costs are expressed in U.S. dollars (2018), and future costs such as productivity losses due to premature death and disability were discounted back to 2018 using a 3% discount rate.

We performed sensitivity analyses to capture uncertainties resulting from 1) the stochastic nature of ABM, 2) model inputs, and 3) model assumptions. Some uncertainty is inherent in the ABM, as the modeling approach uses random processes to capture heterogeneities across individual agents and environments. For instance, MIC values were randomly assigned to bacterial agents, and bacterial agents stochastically encountered antibiotic exposure resulting in different AMR patterns across simulations. This uncertainty from ABM was minimized by running the base case simulation 10,000 times and taking an average across iterations (see Supplemental Appendix). In addition, we conducted a

multivariate probabilistic sensitivity analysis (PSA) to incorporate the additional uncertainty due to model inputs. By using Monte Carlo simulations, the PSA randomly drew parameter values from underlying distributions 10,000 times and estimated variances.²¹ Parameters that were varied during the PSA along with their underlying distributions and uncertainty ranges are listed in Table 1. The 95% confidence intervals were derived from taking the 2.5th and 97.5th percentiles of PSA results. We observed that the variance largely came from the stochastic nature of ABM rather than from key input parameters. Lastly, uncertainties from key model assumptions were also examined. Specific sensitivity analyses were carried out to assess the effect of the number of child agents in the model (i.e., simulating 2,500, 5,000, and 8,000 child agents) and to examine the outcome without herd immunity (see Supplemental Appendix). Estimated outcomes were comparable between 5,000 and 8,000 child agents, confirming that the model simulated adequate numbers of agents. Sensitivity analyses were also conducted to examine the impact of the multiplication approach, where one child agent represented 10 children. Similar results were obtained between simulating 800 child agents of 10 children compared with 8,000 child agents of one child, showing that the underlying model was robust.

Model validation. We assessed the model by examining its face and empirical validities. Face validity ensured that the model contained important elements, while empirical validity examined whether generated data are in line with other existing data. We validated the model at different levels, including at the micro-level representing fundamental rules the agents must follow, and at the macro-level reflecting the outcomes produced by complex interactions between agents and environments. Because this study included two sub-models that had within- and between-model interactions, we looked into the validity of each submodel and the composite outcomes that resulted from their interactions. For face validity, we consulted with several experts in microbiology and pharmacokinetics to ensure that the bacteria submodel included important components of AMR, such as antibiotic selection pressure and fitness costs. For the human submodel, we included factors that impacted disease incidence and treatment patterns, including direct and indirect effects of immunization and care-seeking behaviors. The aggregate results met face validity where the magnitude of emerging resistance against different antibiotics was correlated with antibiotic utilization, where antibiotics prescribed frequently were more likely to gain resistance, and those used less could reverse the trend of accumulating resistance due to fitness costs. For empirical validation, we compared our results with several outcomes of interest, such as disease incidence and burden of malaria. We calibrated the model so that the study outputs aligned with available data from Ethiopia.

RESULTS

Over the 1-year simulation, the proportion of non-susceptible *S. pneumoniae* increased by 2.14% (95% CI: 0.11–4.99%) and 0.47% (95% CI: –0.23–1.08%) for amoxicillin and penicillin, respectively, and reduced by 0.30% (95% CI: –1.22–0.67%) for less commonly used ceftriaxone (Figure 3). Increase in resistance was greater among antibiotics prescribed frequently such as amoxicillin. For antibiotics

not commonly prescribed for pneumococcal diseases such as ceftriaxone, our simulation showed that resistance in the bacteria population could be reduced because of the effect of fitness costs.

We observed a large disease burden from *S. pneumoniae* infections among children under age 5 in Ethiopia (Table 2). We estimated that pneumococcal infections were annually associated with 227,834 (95% CI: 215,319–240,656) cases of pneumonia, 2,405 (95% CI: 1,118–3,725) cases of meningitis, and 2,230,302 (95% CI: 2,191,006–2,269,795) cases of acute otitis media. Pneumococcal infections in children under 5 were responsible for 26,979 (95% CI: 17,695–36,698) deaths per year, with many individuals not seeking care. Meningitis and acute otitis media were simulated to result in 1,310 (95% CI: 373–2,235) disabilities annually. We projected that pediatric pneumococcal infections resulted in 666,370 (95% CI: 636,459–696,435) first-line antibiotic treatments per year in Ethiopia. On average, 1.43 (95% CI: 1.33–1.53) in every 1,000 children under 5 used antibiotics per day to treat pneumococcal diseases.

Antimicrobial resistance resulted in an overall first-line treatment failure rate of 29.38% (95% CI: 28.06–31.28%), where patients needed to switch to second-line therapy, endured a longer duration of illness, and incurred greater costs. Antimicrobial resistance against antibiotics led to 195,763 (95% CI: 180,856–215,506) treatment failures annually, where the majority of treatment failures in the model came from AMR against amoxicillin and penicillin to treat acute otitis media (183,604, 95% CI: 170,984–197,811). Resistance-related treatment failures contributed to 519 (95% CI: 0–1,490) and 2,406 (95% CI: 745–4,843) deaths through treatment at healthcare facilities and from self-medication, respectively.

On average, costs for each successful first-line treatment and treatment failure due to resistance were US\$16.69 (95% CI: US\$12.05–\$25.89) and US\$35.39 (95% CI: US\$25.90–\$52.97), respectively (Table 3). Annual overall costs related to AMR were around 15.8 million, including US\$3,267,389 (95% CI: US\$2,149,468–US\$5,658,071) for ineffective first-line treatments, US\$3,661,208 (95% CI: US\$2,547,098–\$5,741,984) for second-line treatments, and \$8,872,107 (95% CI: US\$2,158,860–US\$20,017,281) for long-term productivity losses due to premature deaths and disabilities. Costs of ineffective first-line treatments included US\$1,432,290 (95% CI: US\$478,281–US\$3,609,864) in direct costs and US\$1,835,099 (95% CI: \$1,671,186–\$2,048,206) in productivity losses for the caregiver. Second-line treatments, which could have been prevented by effective first-line treatments, added US\$1,391,019 (95% CI: US\$416,648–US\$3,275,032) in direct costs and US\$2,270,190 (95% CI: US\$2,130,451–US\$2,466,952) in productivity losses to the caregiver.

DISCUSSION

This is the first study to estimate the health and economic impact of AMR on treatment of pediatric pneumococcal infections in Ethiopia by developing an agent-based model. Our simulation estimated that among all first-line antibiotics used to treat pneumococcal disease, around 30% resulted in ineffective treatments because of AMR, necessitating the need to switch to second-line antibiotic therapy. Treatment failures led to significant child deaths, higher costs due to prolonged

TABLE 1
DREAMR model inputs

Parameter variable	Unit	Value	Standard error or uncertainty range	Underlying distribution	Source
Demographics					
Total population	Thousands	99,873	–	–	UN DESA ⁵⁴
Population, age 0–4 years	Thousands	14,901	–	–	UN DESA ⁵⁴
Population growth rate	%	2.60	–	–	UN DESA ⁵⁴
Gross domestic product per capita	USD	706.76	–	–	World Bank ⁵⁸
Life expectancy at birth	Years	65.97	–	–	UNDESA ⁵⁴
Vaccine characteristics					
Vaccine effectiveness	%	60.20	–	–	Moore et al. ³⁸
Pneumococcal conjugate vaccine 13 coverage rate	%	68	–	–	WHO/UNICEF ³⁷
Pneumococcal disease incidence					
<i>Streptococcus pneumoniae</i> colonization prevalence	%	43.78	2.61	Beta	Gabre et al. ²⁶
Pneumonia					
Incidence	Per 100,000	3,397	–	–	O'Brien et al. ³⁹
Case fatality rate	%	11	7–18	Beta	O'Brien et al. ³⁹
Meningitis					
Incidence	Per 100,000	38	–	–	O'Brien et al. ³⁹
Case fatality rate	%	73	18–94	Beta	O'Brien et al. ³⁹
Neurological sequelae	%	32	3.80	Beta	Arditi et al. ⁵⁰
Acute otitis media					
Incidence	Per 100	60.99	–	–	Monasta et al. ⁴⁰
AOM caused by <i>S. pneumoniae</i>	%	40.00	–	–	Gebre et al. ²⁶
Clinical resolution rate, amoxicillin	%	92.80	–	–	Le Saux et al. ⁴⁷
Clinical resolution rate, no treatment	%	84.17	–	–	Le Saux et al. ⁴⁷
Probability of hearing loss	Per 10,000	22.84	–	–	Monasta et al. ⁴⁰
Herd immunity (indirect effect)					
Pneumonia (in months)					
< 12	%	33	–	–	Blank et al. ⁴³
12–23	%	42	–	–	Blank et al. ⁴³
24–35	%	37	–	–	Blank et al. ⁴³
36–47	%	37	–	–	Blank et al. ⁴³
48–59	%	54	–	–	Blank et al. ⁴³
Meningitis (months)					
< 12	%	48	–	–	Blank et al. ⁴³
12–23	%	56	–	–	Blank et al. ⁴³
24–35	%	43	–	–	Blank et al. ⁴³
36–47	%	43	–	–	Blank et al. ⁴³
48–59	%	41	–	–	Blank et al. ⁴³
Acute otitis media (months)					
< 12	%	22	–	–	Blank et al. ⁴³
12–23	%	27	–	–	Blank et al. ⁴³
24–35	%	0	–	–	Blank et al. ⁴³
36–47	%	0	–	–	Blank et al. ⁴³
48–59	%	0	–	–	Blank et al. ⁴³
Care-seeking behaviors					
Facility types					
Health post	%	12.17	1.76	Dirichlet	Memirie et al. ⁴⁶
Health center	%	53.33	2.69	Dirichlet	Memirie et al. ⁴⁶
Government hospital	%	24.06	2.30	Dirichlet	Memirie et al. ⁴⁶
Private clinic/hospital	%	10.43	1.65	Dirichlet	Memirie et al. ⁴⁶
Hospitalization rate					
Pneumonia					
Health post	%	0	0	Beta	Memirie et al. ⁴⁶
Health center	%	1.63	0.93	Beta	Memirie et al. ⁴⁶
Government hospital	%	31.33	5.09	Beta	Memirie et al. ⁴⁶
Private clinic/hospital	%	36.11	8.01	Beta	Memirie et al. ⁴⁶
Meningitis	%	100	–	–	Assumption
Acute otitis media	%	0	–	–	Assumption
Sought care at formal healthcare facilities					
Pneumonia	%	27.00	0.43	Beta	EDHS ⁴⁴
Meningitis	%	35.30	0.47	Beta	EDHS ⁴⁴
Acute otitis media	%	27.00	0.43	Beta	EDHS ⁴⁴
Mortality rate of nonseeking vs. seeking care	Odds ratio	7.56	3.77–15.10	Normal	Reyes et al. ⁴⁵

(continued)

TABLE 1
Continued

Parameter variable	Unit	Value	Standard error or uncertainty range	Underlying distribution	Source
Proportion of self-medication	%	30.93	2.34	Beta	Gebeyehu et al. ⁷³
Noncompliance					
Formal healthcare facility	%	17.11	3.05	Beta	Yadesa et al. ⁷⁴
Self-medication	%	27.31	2.26	Beta	Gebeyehu et al. ⁷³
Antibiotic' effective dose	%	75.00	–	–	Assumption
Risk of adverse health outcomes	Fold	2	–	–	WHO ⁵²
Pneumonia					
Inpatient	Days	10	–	–	Harris et al. ⁵³
Outpatient	Days	7	–	–	Assumption
Meningitis	Days	15	–	–	Arditi et al. ⁵⁰
Acute otitis media	Days	7	–	–	Klein et al. ⁴⁸
Follow-up since onset of therapy	Days	3	–	–	Harris et al. ⁵³
Initial antibiotics resistance rate					
Amoxicillin	%	29.00	–	–	Anagaw et al. ¹⁸
Penicillin	%	31.30	–	–	Anagaw et al. ¹⁸
Ceftriaxone	%	9.80	–	–	Anagaw et al. ¹⁸
Antibiotic utilization					
Pneumonia					
Amoxicillin	%	20.80	3.63	Dirichlet	Achalu et al. ⁴⁹
Penicillin	%	25.60	3.90	Dirichlet	Achalu et al. ⁴⁹
Ceftriaxone	%	53.60	4.46	Dirichlet	Achalu et al. ⁴⁹
Meningitis					
Ceftriaxone	%	100.00	–	–	Achalu et al. ⁴⁹
Acute otitis media					
Amoxicillin	%	57.14	18.70	Beta	Achalu et al. ⁴⁹
Penicillin	%	42.86	18.70	Beta	Achalu et al. ⁴⁹
Costs					
Direct medical costs					
Outpatient	%	78.95	–	–	Memirie et al. ⁴⁶
Inpatient	%	79.52	–	–	Memirie et al. ⁴⁶
Mean medical expenditure					
Outpatient					
Health post	USD	1.61	2.71	Gamma	Memirie et al. ⁴⁶
Health center	USD	4.06	5.91	Gamma	Memirie et al. ⁴⁶
Government hospital	USD	12.08	12.05	Gamma	Memirie et al. ⁴⁶
Private clinic/hospital	USD	28.12	8.85	Gamma	Memirie et al. ⁴⁶
Inpatient					
Health post	USD	–	–	–	Memirie et al. ⁴⁶
Health center	USD	12.13	8.80	Gamma	Memirie et al. ⁴⁶
Government hospital	USD	47.89	28.81	Gamma	Memirie et al. ⁴⁶
Private clinic/hospital	USD	139.66	71.97	Gamma	Memirie et al. ⁴⁶
DALY weight for hearing loss	DALYs	0.158	–	–	Salomon et al. ⁵⁹
Pharmacokinetics and pharmacodynamics characteristics					
Penicillin G					
Dose	IU/kg/day	50,000	–	–	AAP ²⁹
Dosing interval	Hours	6	–	–	AAP ²⁹
mg to IU conversion	IU/mg	1,670	–	–	Humphrey et al. ²²
Volume of distribution	L/kg	1.39	–	–	Bolme et al. ²³
Clearance	mL/min/kg	22.2	–	–	Bolme et al. ²³
Ceftriaxone					
Dose	mg/kg/day	50–100	–	–	Bradley et al. ²⁸
Dosing interval	Hours	12	–	–	Bradley et al. ²⁸
Vd	mL/kg	387	–	–	Steele et al. ²⁵
Elimination half-life ($t_{1/2}$)	Hours	5.4	–	–	Steele et al. ²⁵
Amoxicillin					
Dose	mg/kg/dose	30	–	–	Bradley et al. ²⁸
Dosing interval	Hours	8	–	–	Bradley et al. ²⁸
Vd	mL/kg	764	–	–	Schaad et al. ²⁴
Elimination half-life ($t_{1/2}$)	Hours	1.17	–	–	Schaad et al. ²⁴
Fitness cost (resistant vs. susceptible)	Relative fitness	0.86	–	–	Maher et al. ³⁴

AAP = American Academy of Pediatrics; AOM = acute otitis media; DALY = disability-adjusted life year; EDHS = Ethiopia Demographic and Health Survey; IU = international units; UN DESA = United Nations Department of Economic and Social Affairs; UNICEF = United Nations Children's Fund; USD = United States dollars; Vd = volume of distribution; WHO = World Health Organization.

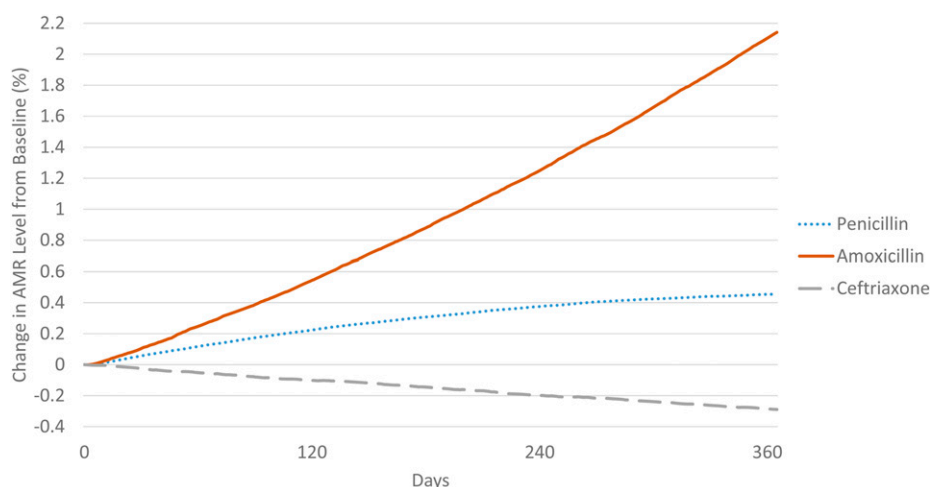


FIGURE 3. Changes in proportions of antimicrobial resistance over time. This figure appears in color at www.ajtmh.org.

treatment duration, greater loss in productivity, and societal costs associated with increasing AMR. Treatment failures and resulting impact can be averted by controlling the development of AMR associated with antibiotics.

Our results demonstrate the substantial impact of AMR in low-income countries such as Ethiopia, with a large infectious disease burden and high rates of antibiotic utilization. Although this study focused specifically on antibiotic utilization for treatment of pediatric pneumococcal diseases over a year, we still observed sizable treatment failures and avertable costs to patients and healthcare systems. These results are important because the impact of AMR is often not measured and underappreciated in developing countries, although

antibiotics are commonly available without prescriptions, and widespread antibiotic misuse is prevalent in these settings.

Our results revealed that resistance tended to increase for amoxicillin and penicillin, whereas it tended to decrease for ceftriaxone to treat pediatric pneumococcal infections in Ethiopia over a year. Differences in these estimates can be explained by two influential factors: the magnitude of antibiotic utilization and fitness costs. Whereas higher antibiotic exposure increased AMR, fitness costs provided negative feedback as resistant bacteria often have reduced fitness to compete with susceptible strains.³⁴ In most cases, the effects of antibiotic exposure overrode the opposite influence of bacteria fitness, resulting in an increase in resistance.⁶⁰

TABLE 2
Annual impact of antibiotic resistance on health outcomes due to pneumococcal infections in Ethiopia

Outcomes	Value*	95% Probabilistic sensitivity analysis CI†	
Incremental change in resistance			
Amoxicillin, %	2.14	0.11	4.99
Penicillin, %	0.47	-0.23	1.08
Ceftriaxone, %	-0.30	-1.22	0.67
Average defined daily dose, per 1,000 patient days	1.43	1.33	1.53
Disease cases			
Pneumococcal pneumonia, <i>n</i>	227,834	215,319	240,656
Pneumococcal meningitis, <i>n</i>	2,405	1,118	3,725
Pneumococcal AOM, <i>n</i>	2,230,302	2,191,006	2,269,795
Adverse health outcomes			
Overall death, <i>n</i>	26,979	17,695	36,698
Death during formal treatment, <i>n</i>	2,979	1,118	5,774
Death due to resistance in formal treatment, <i>n</i>	519	0	1,490
Death due to resistance in self-medication, <i>n</i>	2,406	745	4,843
Death due to not seeking care, <i>n</i>	21,075	15,832	24,591
Disability, <i>n</i>	1,310	373	2,235
Treatment behaviors			
Overall treatments, <i>n</i>	666,370	636,459	696,435
Overall treatment failures, <i>n</i>	195,763	180,856	215,506
Treatment failures (pneumonia), <i>n</i>	11,931	9,872	16,950
Treatment failures (meningitis), <i>n</i>	228	0	745
Treatment failures (AOM), <i>n</i>	183,604	170,984	197,811
Proportion of treatment failures, %	29.38	28.06	31.28

AOM = acute otitis media; CI = confidence interval.

* Point estimates were derived by taking average values across 10,000 base case simulations.

† The 2.5th and 97.5th percentiles across 10,000 probabilistic sensitivity analyses were used to derive 95% CIs.

TABLE 3
Annual impact of antibiotic resistance on costs of pneumococcal infections in Ethiopia

Outcomes	Value*	95% Probabilistic sensitivity analysis CI†	
Average costs per successful first-line treatment			
Overall costs, USD	16.69	12.05	25.89
Direct medical costs, USD	5.78	2.13	13.03
Direct nonmedical costs, USD	1.54	0.56	3.46
Productivity losses for caregiver, USD	9.37	9.36	9.40
Average costs per treatment failure due to resistance			
Overall costs, USD	35.39	25.90	52.97
Direct medical costs, USD	11.39	3.91	25.23
Direct nonmedical costs, USD	3.03	1.03	6.72
Short-term productivity losses (caregiver), USD	20.97	20.95	21.02
Long-term productivity losses (death/disability), USD	45.32	11.94	92.89
Annual costs incurred by first-line treatments due to resistance			
Overall costs, USD	3,267,389	2,149,468	5,658,071
Direct medical costs, USD	1,131,602	378,609	2,851,699
Direct nonmedical costs, USD	300,688	99,672	758,165
Productivity losses for caregiver, USD	1,835,099	1,671,186	2,048,206
Annual costs incurred by second-line treatments due to resistance			
Overall costs, USD	3,661,208	2,547,098	5,741,984
Direct medical costs, USD	1,098,473	329,218	2,585,820
Direct nonmedical costs, USD	292,546	87,430	689,212
Short-term productivity losses (caregiver), USD	2,270,190	2,130,451	2,466,952
Long-term productivity losses (death/disability), USD	8,872,107	2,158,860	20,017,281

CI = confidence interval; USD = United States dollars.

* Point estimates were derived by taking average values across 10,000 base case simulations.

† The 2.5th and 97.5th percentiles across 10,000 probabilistic sensitivity analyses were used to derive 95% CIs.

However, when antibiotics were less frequently used, the effect of fitness costs was observed where the proportion of resistant strains reduced.⁶⁰ By incorporating antibiotic exposure and fitness costs of bacteria in the DREAMR model, we demonstrated the dynamic relationship between antibiotic use and AMR.

Because AMR growth is directly attributable to antibiotic use, antibiotic stewardship to improve appropriate use of antibiotics is essential. Antimicrobial stewardship can be defined as “a coherent set of actions which promote using antimicrobials in ways that ensure sustainable access to effective therapy for all who need them.”⁶¹ On the supply side, this may entail making antibiotics obtainable by prescription, making accurate diagnosis, following antimicrobial guidelines, monitoring antimicrobial use and resistance, and investing in a clinical decision support system to improve prescribing and responsible use. On the demand side, interventions may involve education and community engagement programs to ensure that patients understand issues surrounding rational medication use, including how and when to take antimicrobials, and ensuring that patients do not store and use leftover antimicrobials. Improved antimicrobial stewardship involving both demand- and supply-side initiatives are critical to mitigating the global impact of AMR. Increased PCV immunization coverage could also reduce pneumococcal disease incidence, thereby reducing antibiotic utilization and curbing AMR. Future research should demonstrate the impact of antibiotic stewardship and vaccination on controlling the development of AMR.

Our findings are consistent with previous studies on the impact of pneumococcal disease, antibiotic utilization, and

proportion of resistance. Global estimate of the burden of *S. pneumoniae* in children under 5 has projected approximately 57,000 pneumococcal deaths in Ethiopia in 2000, before the country introduced PCV.³⁹ In our model, overall deaths due to pneumococcal diseases reflect current vaccine coverage and vaccine efficacy, resulting in fewer deaths. As for the rate of antibiotic utilization, previous studies report average overall antibiotic use at around 10 to 35 DDDs across countries.^{13,62} We believe our lower antibiotic utilization rate is reasonable, given that our model only focused on utilization of three antibiotics for pediatric pneumococcal infections, and because Ethiopia has a relatively low proportion of individuals seeking care. Although data about antibiotic resistance in Africa are scarce due to the lack of susceptibility testing and weak surveillance systems, previous studies have reported the proportion of nonsusceptible *S. pneumoniae* to be between 9% and 69%, which aligns with our analysis.^{13,63} Our results on the positive correlation between antibiotic use and AMR are consistent with previous findings in other countries.^{64,65}

There are a number of potential limitations to our study. First, our model relies on the quality of data reported in the existing literature. Although we conducted an extensive literature search to incorporate the most recent and best quality published data, results of the study are subject to the quality of data inputs. Data availability also limits the model's ability to capture heterogeneity across the population. For example, data on antibiotic utilization for people who do not seek care at formal health facilities were limited, including those who obtain antibiotics from pharmacies or from other individuals.

Real-world antibiotic utilization may also differ from recommended treatment guidelines, and antibiotic prescription behaviors may change in accordance with resistance patterns. While previous studies have suggested the examination of *S. pneumoniae* serotypes, this study was not able to incorporate the impact of different serotypes, assuming no difference in bacterial virulence and response to PCV13. This limited our ability to examine how heterogeneous serotypes of infected patients may affect vaccine effectiveness and disease severity. Further studies should be conducted to examine the heterogeneity of several crucial data, including bacterial susceptibility, effect of serotypes, antibiotic quality, antibiotic utilization, impact of vaccination, bacterial fitness, and treatment costs, in order to provide a more comprehensive insight on the impact.

Secondly, the bacteria submodel does not incorporate the effects of genetic mutation and substandard and falsified antibiotics. Although bacteria can develop resistance under strong selection pressure due to antibiotic exposure, they may also become resistant as a result of genetic mutation under subinhibitory concentrations or acquire resistant genes from other strains.^{66–68} Mutation-related resistance may play an important role among countries with high prevalence of substandard and falsified medicines, where bacteria are more likely to be exposed to subinhibitory antibiotic concentrations.⁶⁹ Future studies should examine these impacts.

Third, our model examined the AMR impact across three antibiotics commonly used in Ethiopia to treat pediatric pneumococcal disease. These results may not be easily generalizable to other antibiotics and contexts. In addition, antibiotic treatment might not be optimized, where the route of administration and studies of combination therapies that can result in more realistic exposures among patients were not available for the study. Finally, our model results focused solely on the use of antibiotics for human health. Further studies should estimate the impact of AMR from a broader scope under a One Health approach, taking into account antibiotic use across animal and environmental health sectors and additional stakeholders influenced by the accumulation of AMR.⁷⁰

The DREAMR model is the first to estimate the annual impact of AMR in a low-income country by estimating the impact of antibiotic resistance on treatment of pediatric pneumococcal disease in Ethiopia. The results can inform in-country stakeholders, international donors, and national and regional child health experts to recognize the burden of AMR and examine interventions to improve appropriate antibiotic use. Reducing the impact of AMR is essential to achieve the Sustainable Development Goals (SDGs)—to achieve access to safe, effective, quality, and affordable essential medicines for all.⁷¹ Maintaining the effectiveness of current antibiotics is also vital to meet the Global Health Security Agenda (GHSA) to help create a world safe and secure from infectious disease threats.⁷²

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