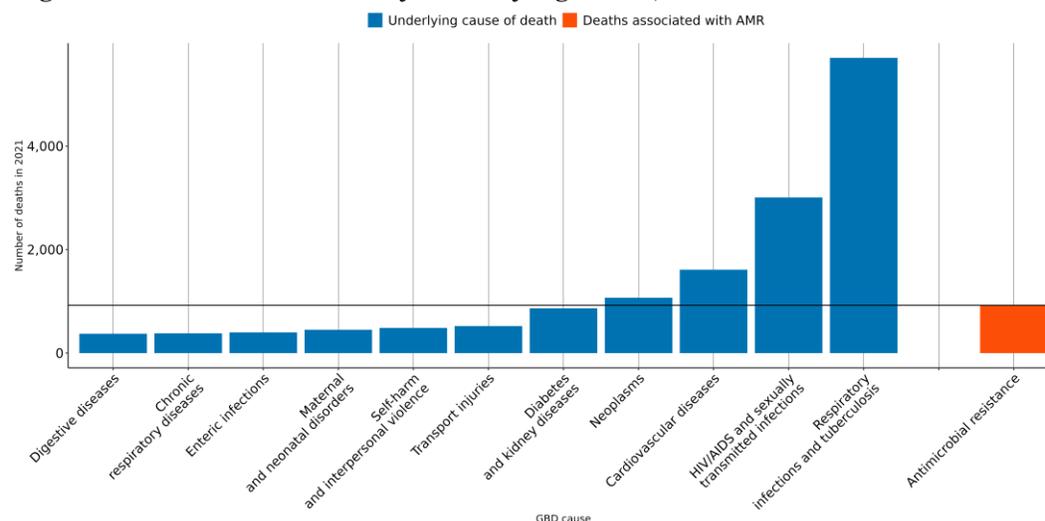


The burden of antimicrobial resistance (AMR) in Eswatini

Executive summary

- Antimicrobial Resistance (AMR) is a major global health threat, over **300 lives** have been lost each year since 1990 in Eswatini due to AMR.
- In 2021, there were an estimated **224 UI (114-334)** deaths attributable to AMR and **925 UI (623-1,230)** deaths associated with AMR in this location.
- The largest number of deaths associated with AMR in 2021 occurred among those aged **50 to 69** in the country.
- Among the most deadly pathogen-drug combinations in 2021 were multi-drug resistant *Mycobacterium tuberculosis* (excluding extensive drug-resistance), *Acinetobacter baumannii* resistant to carbapenems and *Streptococcus pneumoniae* resistant to carbapenems.

Figure 1 Number of deaths by underlying cause, and those associated with AMR in 2021



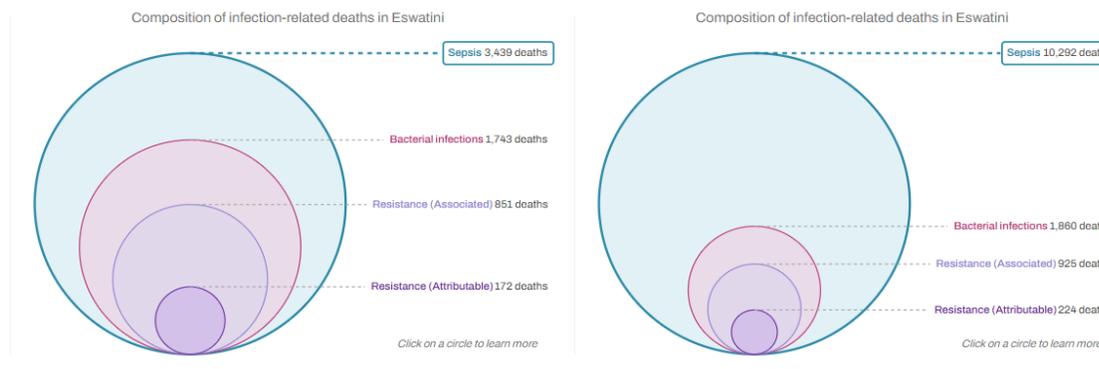
- In 2021, the number of deaths associated with AMR (orange bar in *figure 2*) were high compared to the most relevant underlying causes of death (depicted in blue) in the country. AMR associated deaths occur within multiple Global Burden of Disease (GBD) causes of death and AMR is not an underlying cause of death by itself.
- At the [2024 United Nations General Assembly high level meeting on antimicrobial resistance](#), country members agreed to aim for a **10% reduction** compared to 2019 baseline (**from 4.95 to 4.45 million**) in the global number of deaths associated with AMR by 2030. But [our forecast](#) indicates that in absence of concerted action, deaths associated with AMR could reach **5.5 million** (UI 4.8 - 6.2) if current trends continue. For Eswatini, a 10% reduction means to decrease the number of deaths associated with AMR to **941**, but currently the trend for this country could reach up to **964 UI [619-1,450]** AMR-associated deaths in 2030.

AMR in Eswatini

Key takeaways

- Antimicrobial Resistance (AMR) is a major global health threat, over *a million lives* have been lost each year since 1990.
- Globally, 4.71 (95% Uncertainty Interval (UI) 4.2-5.2) million deaths were associated with bacterial drug-resistant infections in 2021.
- And 1.14 (UI 1 - 1.3) million deaths were attributable to bacterial drug-resistant infection in the same year.
- *39 (UI 33 - 46) million deaths* directly attributable to bacterial AMR are projected to occur between 2025-2050 unless concerted action is taken. This equates to three deaths every minute.

Figure 2 Comparing 30 years of infection related deaths, and those associated with and attributable to AMR in Eswatini between 1990 and 2019.



- To look at these and more visualization interactively visit [Measuring Infectious Causes and Resistance Outcomes for Burden Estimation \(MICROBE\)](#)
- In **Eswatini** in 2021, there were an estimated **224 UI (114-334)** deaths attributable to AMR and **925 UI (623-1,230)** deaths associated with AMR. Here “*attributable deaths*” are considered to be those that would have been prevented had the drug-resistant bacteria causing the infections not been drug-resistant. “*Associated deaths*” are considered to be those that would not have occurred had the infections been prevented entirely.
- Across 204 countries, **Eswatini has the 15th highest** age-standardized mortality rate associated with AMR in 2021.
- *Table 1* shows the bacteria which caused most deaths in 2021 (↑ indicates an increasing estimated annual rate between 1990-2021, ↓ indicates a decreasing annual trend), and *table 2* shows the pathogen-drug combinations which caused most deaths in 2021.

Table 1. Bacteria which cause most deaths in 2021 (Number of deaths in parenthesis)

	Overall susceptible and resistant	Associated	Attributable
Burden rank	Mycobacterium tuberculosis 669 UI (398-939) ↑	Streptococcus pneumoniae 170 UI (115-225) ↓	Mycobacterium tuberculosis 47 UI (0-139) ↑
	Streptococcus pneumoniae 220 UI (159-281) ↓	Klebsiella pneumoniae 136 UI (94-178) ↑	Streptococcus pneumoniae 36 UI (22-51) ↓
	Klebsiella pneumoniae 185 UI (133-236) ↑	Mycobacterium tuberculosis 125 UI (36-272) ↑	Klebsiella pneumoniae 32 UI (22-42) ↑
	Staphylococcus aureus 144 UI (104-183) ↑	Escherichia coli 119 UI (84-154) ↓	Acinetobacter baumannii 30 UI (23-37) ↑
	Pseudomonas aeruginosa 135 UI (98-172) ↑	Staphylococcus aureus 95 UI (59-131) ↑	Escherichia coli 23 UI (15-31) ↑
	Escherichia coli 129 UI (93-165) ↓	Acinetobacter baumannii 73 UI (52-94) ↑	Pseudomonas aeruginosa 16 UI (9-23) ↑
	Acinetobacter baumannii 81 UI (58-103) ↑	Pseudomonas aeruginosa 65 UI (42-88) ↑	Staphylococcus aureus 16 UI (9-22) ↑
	Group B Streptococcus 39 UI (27-50) ↑	Serratia spp. 21 UI (15-27) ↓	Serratia spp. 6 UI (4-8) ↓
	Shigella spp. 36 UI (16-55) ↓	Shigella spp. 20 UI (7-33) ↓	Enterobacter spp. 4 UI (3-6) ↓
	Serratia spp. 30 UI (21-38) ↑	Enterobacter spp. 18 UI (13-23) ↑	Shigella spp. 2 UI (0-4) ↓

Annualized rate of change (1990-2021):
 <-3% (dark blue), -1.5% to 0% (light blue), 1.5% to 3% (red), >5.0% (dark red)
 -3% to -1.5% (medium blue), 0% to 1.5% (pink), 3% to 5% (orange)

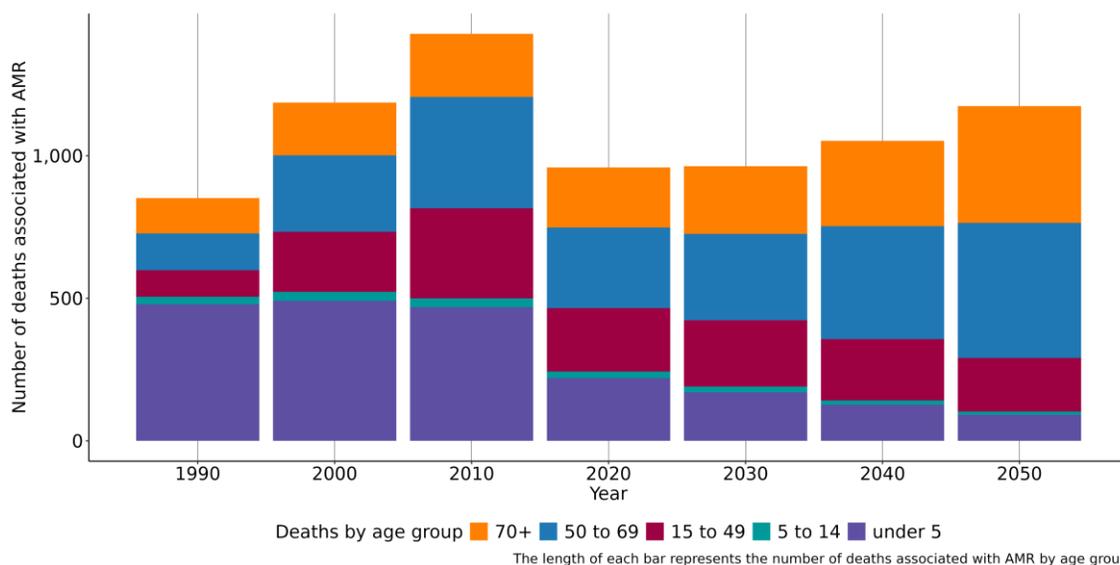
Table 2. Combinations which cause most deaths in 2021 (Number of deaths in parenthesis)

	Associated	Attributable
Burden Rank	Mycobacterium tuberculosis MDR excluding XDR 124 UI (36-269) ↑	Mycobacterium tuberculosis MDR excluding XDR 46 UI (0-137) ↑
	Streptococcus pneumoniae Macrolides 119 UI (78-160) ↓	Streptococcus pneumoniae Carbapenems 21 UI (11-31) ↓
	Streptococcus pneumoniae TMP-SMX 119 UI (68-169) ↓	Acinetobacter baumannii Carbapenems 19 UI (14-24) ↑
	Escherichia coli Aminopenicillin 117 UI (79-155) ↓	Klebsiella pneumoniae 3GC 10 UI (5-14) ↑
	Klebsiella pneumoniae 3GC 111 UI (78-144) ↑	Acinetobacter baumannii Fluoroquinolones 7 UI (6-9) ↑
	Klebsiella pneumoniae Beta-Lactam/Lactamase Inhib. 105 UI (66-145) ↑	Klebsiella pneumoniae Aminoglycosides 7 UI (4-10) ↓
	Klebsiella pneumoniae Aminoglycosides 101 UI (69-133) ↑	Klebsiella pneumoniae Fluoroquinolones 7 UI (4-10) ↑
	Klebsiella pneumoniae TMP-SMX 99 UI (66-133) ↓	Staphylococcus aureus TMP-SMX 6 UI (4-9) ↑
	Klebsiella pneumoniae Fluoroquinolones 95 UI (61-128) ↑	Pseudomonas aeruginosa Carbapenems 6 UI (3-9) ↑
	Streptococcus pneumoniae Beta-Lactam/Lactamase Inhib. 89 UI (46-132) ↑	Escherichia coli Fluoroquinolones 5 UI (2-8) ↑

Annualized rate of change (1990-2021):
 <-3% (dark blue), -1.5% to 0% (light blue), 1.5% to 3% (red), >5.0% (dark red)
 -3% to -1.5% (medium blue), 0% to 1.5% (pink), 3% to 5% (orange)

- Independently of antimicrobial resistance, the infectious syndromes accounting for the most deaths in 2021 were as follows (estimated thousands of deaths in parenthesis) lower respiratory infection (excl. COVID) (829 UI (601-1,060)), tuberculosis (669 UI (398-939)), bloodstream infections (582 UI (413-751)), diarrhea (396 UI (245-547)) and meningitis (83 UI (55-111)).

Figure 3. Number of deaths associated with AMR by age group between 1990-2020 and 2050 projection



- In Eswatini, people aged under 5 experienced the largest number of deaths associated with AMR in 1990 but this changed by 2021 as the largest number of deaths occurred among the 50 to 69. This indicates that prevention of infections among the under 5 has contributed to the reduction in the number of AMR associated deaths. In 2021, the number of deaths associated with AMR among the 50 to 69 was 277 UI (181-373), whereas the mortality rate per 100,000 was 875 UI (652-1,100).

Data sources for Eswatini

In total, 520 million individual records or isolates covering 19,513 study-location-years were used as input data to our estimation process. The subset of input data for this country is shown below.

Table 3. Data inputs for Eswatini by source type

Source type	Years	Sample size	Sample size units
Antibiotic use	1990-2021	648	Study-year datapoints
Microbial or laboratory data without outcome	1990-2021	4,701	Isolates

More information

About GRAM:

The purpose of the Global Research on AntiMicrobial resistance (GRAM) project is to **generate accurate and timely estimates of the magnitude and trends in antimicrobial resistance (AMR) burden** across the world, which can be used to inform treatment guidelines and agendas for decision-making and research, detect emerging problems and monitor trends to inform global strategies, as well as facilitate the assessment of interventions over time.

GRAM is the flagship project of the University of Oxford–IHME Strategic Partnership. GRAM was launched with support from the United Kingdom Department of Health and Social Care’s Fleming Fund, and the Wellcome Trust.

All resources:

For all resources on AMR analysis at IHME, visit <https://www.healthdata.org/antimicrobial-resistance>.

To look at these and more visualization interactively visit [Measuring Infectious Causes and Resistance Outcomes for Burden Estimation \(MICROBE\)](#).

Data sources:

To download the list of data input sources by country, and AMR results by region, visit the [Global Health Data Exchange \(GHDx\)](#).

Contact us:

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