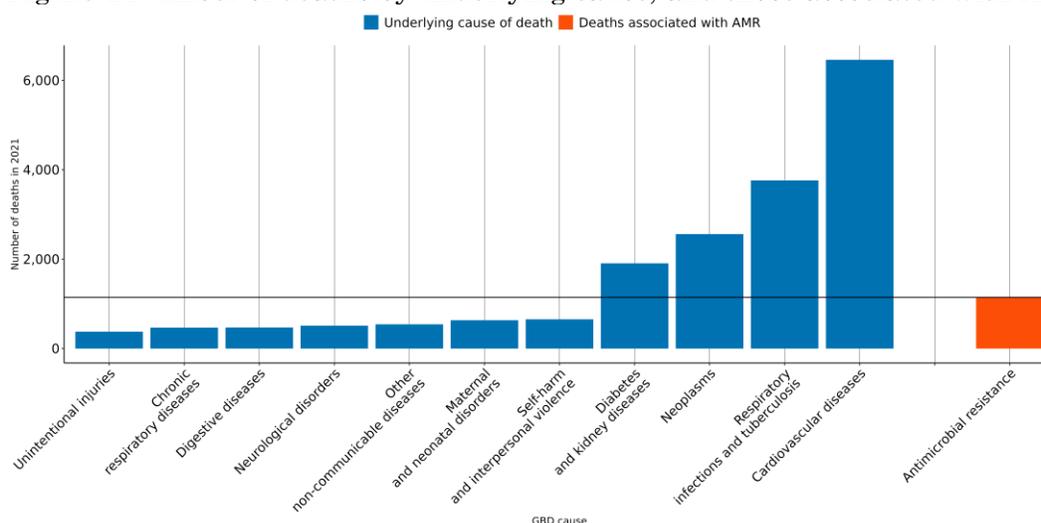


The burden of antimicrobial resistance (AMR) in Palestine

Executive summary

- Antimicrobial Resistance (AMR) is a major global health threat, over **400 lives** have been lost each year since 1990 in Palestine due to AMR.
- In 2021, there were an estimated **300 UI (241-360)** deaths attributable to AMR and **1,150 UI (956-1,340)** deaths associated with AMR in this location.
- The largest number of deaths associated with AMR in 2021 occurred among those aged **70+** in the country.
- Among the most deadly pathogen-drug combinations in 2021 were *Staphylococcus aureus* resistant to methicillin, *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



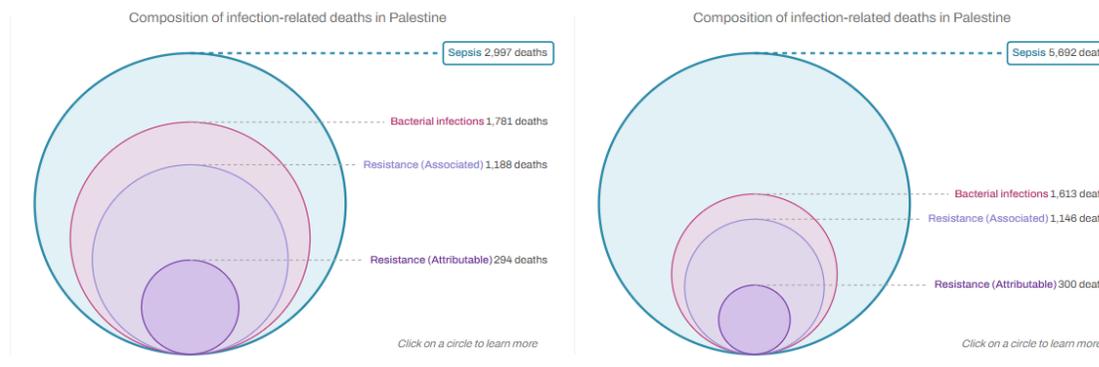
- In 2021, the number of deaths associated with AMR (orange bar in *figure 2*) were high compared to the most relevant underlying causes of death (depicted in blue) in the country. AMR associated deaths occur within multiple Global Burden of Disease (GBD) causes of death and AMR is not an underlying 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 Palestine, a 10% reduction means to decrease the number of deaths associated with AMR to **1,060**, but currently the trend for this country could reach up to **1,570 UI [1,230-2,130]** AMR-associated deaths in 2030.

AMR in Palestine

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 Palestine between 1990 and 2019.



- To look at these and more visualization interactively visit [Measuring Infectious Causes and Resistance Outcomes for Burden Estimation \(MICROBE\)](#)
- In **Palestine** in 2021, there were an estimated **300 UI (241-360)** deaths attributable to AMR and **1,150 UI (956-1,340)** 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, **Palestine has the 90th lowest** 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	Staphylococcus aureus 325 UI (287-363) ↑	Staphylococcus aureus 244 UI (193-294) ↑	Staphylococcus aureus 72 UI (55-89) ↑
	Streptococcus pneumoniae 218 UI (193-242) ↓	Streptococcus pneumoniae 167 UI (138-196) ↓	Streptococcus pneumoniae 44 UI (32-56) ↓
	Pseudomonas aeruginosa 188 UI (167-209) ↑	Escherichia coli 155 UI (128-181) ↑	Acinetobacter baumannii 43 UI (37-49) ↓
	Klebsiella pneumoniae 185 UI (163-206) ↓	Klebsiella pneumoniae 145 UI (124-165) ↓	Pseudomonas aeruginosa 34 UI (26-42) ↑
	Escherichia coli 178 UI (156-199) ↑	Pseudomonas aeruginosa 130 UI (108-151) ↑	Klebsiella pneumoniae 34 UI (28-41) ↓
	Acinetobacter baumannii 107 UI (95-120) ↓	Acinetobacter baumannii 106 UI (94-119) ↓	Escherichia coli 30 UI (23-38) ↑
	Group A Streptococcus 45 UI (39-51) ↑	Enterococcus faecium 34 UI (30-38) ↑	Enterococcus faecium 10 UI (9-12) ↑
	Enterobacter spp. 43 UI (38-48) ↑	Enterobacter spp. 33 UI (28-37) ↓	Enterobacter spp. 8 UI (5-10) ↓
	Group B Streptococcus 42 UI (36-49) ↓	Proteus spp. 25 UI (19-30) ↑	Serratia spp. 5 UI (4-6) ↓
	Serratia spp. 40 UI (36-45) ↑	Serratia spp. 21 UI (18-24) ↓	Haemophilus influenzae 4 UI (2-6) ↑

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

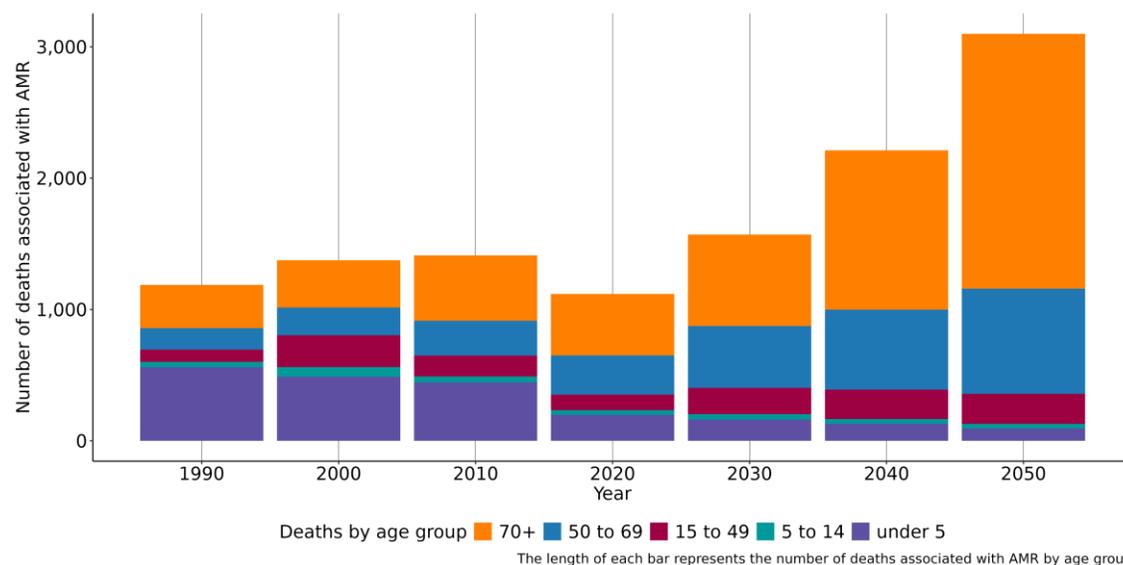
Table 2. Combinations which cause most deaths in 2021 (Number of deaths in parenthesis)

	Associated	Attributable
Burden Rank	Staphylococcus aureus Methicillin 221 UI (161-282) ↑	Staphylococcus aureus Methicillin 56 UI (42-70) ↑
	Staphylococcus aureus Macrolides 152 UI (121-182) ↑	Streptococcus pneumoniae Carbapenems 36 UI (25-46) ↓
	Escherichia coli Aminopenicillin 149 UI (113-185) ↑	Acinetobacter baumannii Carbapenems 25 UI (20-30) ↓
	Streptococcus pneumoniae Carbapenems 123 UI (93-154) ↓	Pseudomonas aeruginosa Carbapenems 16 UI (11-22) ↑
	Klebsiella pneumoniae Beta-Lactam/Lactamase Inhib. 123 UI (96-149) ↓	Acinetobacter baumannii Fluoroquinolones 11 UI (9-13) ↑
	Klebsiella pneumoniae TMP-SMX 120 UI (101-139) ↓	Escherichia coli 3GC 9 UI (6-12) ↑
	Streptococcus pneumoniae Macrolides 120 UI (93-146) ↓	Klebsiella pneumoniae 3GC 8 UI (5-11) ↑
	Acinetobacter baumannii Carbapenems 105 UI (93-117) ↓	Pseudomonas aeruginosa Fluoroquinolones 8 UI (6-10) ↑
	Acinetobacter baumannii 3GC 104 UI (92-116) ↓	Enterococcus faecium Vancomycin 8 UI (7-9) ↑
	Acinetobacter baumannii 4GC 104 UI (92-116) ↓	Klebsiella pneumoniae Fluoroquinolones 7 UI (5-10) ↑

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

- Independently of antimicrobial resistance, the infectious syndromes accounting for the most deaths in 2021 were as follows (estimated thousands of deaths in parenthesis) bloodstream infections (947 UI (830-1,070)), lower respiratory infection (excl. COVID) (908 UI (799-1,020)), peritoneal and intra-abdominal infections (160 UI (135-185)), urinary tract infections and pyelonephritis (107 UI (86-127)) and infections of the skin and subcutaneous systems (78 UI (62-94)).

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



- In Palestine, 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 70+. 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 70+ was 478 UI (391-565), whereas the mortality rate per 100,000 was 460 UI (377-544).

Data sources for Palestine

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 Palestine by source type

Source type	Years	Sample size	Sample size units
Antibiotic use	1990-2021	5,042	Study-year datapoints
Microbial or laboratory data with outcome	2010-2021	2,881	Isolates
Literature studies	1990-2021	4,265	Cases/isolates/susceptibility tests
Single drug resistance profile data	2010-2021	50,731	Antibiotic susceptibility test

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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