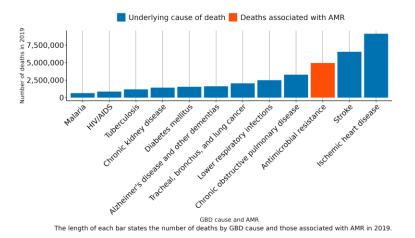


The burden of antimicrobial resistance (AMR) in Antigua and Barbuda

AMR represents a global challenge

- **4.95 million** people who died in 2019 suffered from drug-resistant infections.
- AMR directly caused **1.27 million** of those deaths.
- 1 in 5 of those deaths occurred among children under 5 years old.

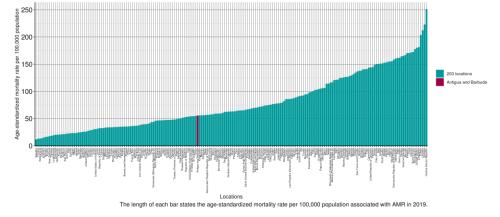
Figure 1 Global number of deaths by GBD cause and those associated with AMR



AMR burden in Antigua and Barbuda

- In Antigua and Barbuda in 2019, there were 12 deaths attributable to AMR and 49 deaths associated with AMR.
- Antigua and Barbuda has the 85th lowest age-standardized mortality rate per 100,000 population associated with AMR across 204 countries.

Figure 2. Age-standardized mortality rate associated with AMR in 2019 for 204 locations

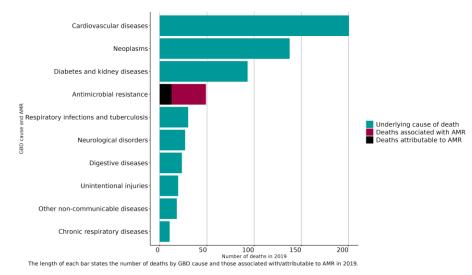


In the GBD region of **Caribbean**, **Antigua and Barbuda has the 7th lowest** agestandardized mortality across **19** countries. •



The number of AMR deaths in **Antigua and Barbuda** is higher than deaths from respiratory infections and tuberculosis, neurological disorders, digestive diseases, unintentional injuries, and other non-communicable diseases.

Figure 3. Placing AMR in context with other causes of death in 2019, Antigua and Barbuda



- There are five pathogens to be aware of in Antigua and Barbuda (number of deaths associated with AMR in parentheses): Acinetobacter baumannii (9), Escherichia coli (7), Staphylococcus aureus (7), Streptococcus pneumoniae (6), and Klebsiella pneumoniae (5).
- These commonly caused bloodstream infections, lower respiratory infections and all related infections in the thorax, and peritoneal and intraabdominal infections.
- According to the Tracking AMR Country Self-Assessment Survey (TrACSS)1, "there is no National AMR action plan" in Antigua and Barbuda. The next step is that the plan should be "developed" in the short term, and this data should be used to ensure this progression.

¹ World Health Organization (WHO). TrACSS 2018-2019 [Internet]. [cited 2023 Feb 16]. Available from: <u>https://amrcountryprogress.org/download/AMR-self-assessment-survey-country-responses-2018-19.xls</u>



Addendum: A summary of data sources for Antigua and Barbuda

In total, 471 million individual records or isolates covering 7,585 study-location-years were used as input data to our estimation process to develop the most comprehensive set of AMR estimates to date. There was no input data accessible that could be used to inform these country estimates. Estimates were informed by results from the Global Burden of Disease study and data from the surrounding region. Any datasets that could be used to improve these estimates in the future are welcome. Going forward, new strategies for data preparation, implementation of more usable data, and inclusion of new systematic literature reviews will result in an enhanced overall analysis. Specific policies that will improve AMR surveillance and link outcomes with resistance data will help us improve this research endeavor. If we expand the quantity and quality of data in this country (but also worldwide), we are confident that future iterations of these estimates (but also those of other research groups) will be able to assess the effect of AMR even more precisely and help tailor optimal approaches to ever-increasing threat of antibiotic resistance.



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, the Wellcome Trust, and the Bill & Melinda Gates Foundation.

All resources:

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

Further details are available on our FAQ page

Data sources:

To download the list of data input sources by country, and AMR results by region, visit the <u>Global Health Data Exchange (GHDx)</u>.

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