



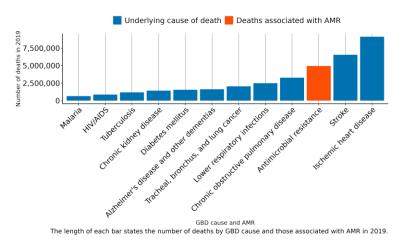


# The burden of antimicrobial resistance (AMR) in United States Virgin Islands

## 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.

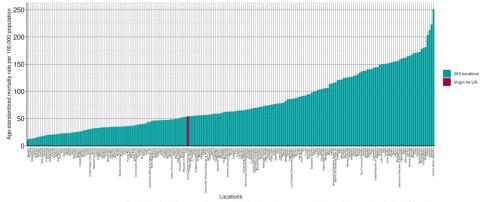
Figure 1 Global number of deaths by GBD cause and those associated with AMR



### AMR burden in United States Virgin Islands

- In United States Virgin Islands in 2019, there were 22 deaths attributable to AMR and **90** deaths associated with AMR.
- United States Virgin Islands has the 81st 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



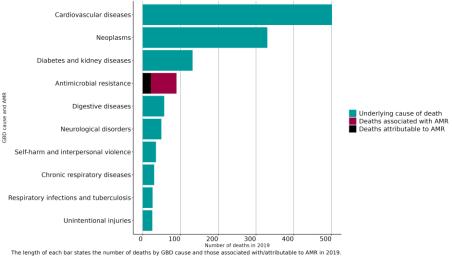
In the GBD region of Caribbean, United States Virgin Islands has the 5th lowest age-standardized mortality across 19 countries.





The number of AMR deaths in **United States Virgin Islands** is higher than deaths from digestive diseases, neurological disorders, self-harm and interpersonal violence, chronic respiratory diseases, and respiratory infections and tuberculosis.

Figure 3. Placing AMR in context with other causes of death in 2019, United States Virgin Islands



- There are five pathogens to be aware of in **United States Virgin Islands** (number of deaths associated with AMR in parentheses): Escherichia coli (18), Staphylococcus aureus (15), Klebsiella pneumoniae (12), Acinetobacter baumannii (9), and Pseudomonas aeruginosa (8).
- These commonly caused peritoneal and intra-abdominal infections, bloodstream infections, lower respiratory infections and all related infections in the thorax, and urinary tract infections and pyelonephritis.
- According to the Tracking AMR Country Self-Assessment Survey (TrACSS)1, "there is no information on a National AMR action plan" in United States Virgin **Islands**. The next step is that the plan should be "developed" in the short term, and this data should be used to ensure this progression.









# Addendum: A summary of data sources for United States Virgin Islands

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 <a href="https://www.healthdata.org/antimicrobial-resistance">https://www.healthdata.org/antimicrobial-resistance</a>.

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 Global Health Data Exchange (GHDx).

#### Contact us:

- For inquiries about the analysis and questions from government officials, health departments, or research institutions: <a href="mailto:engage@healthdata.org">engage@healthdata.org</a>
- For media-related inquiries: media@healthdata.org
- IHME social media handles:
- Twitter: @IHME UW
- Facebook: <a href="https://www.facebook.com/IHMEUW">https://www.facebook.com/IHMEUW</a>
- LinkedIn: <a href="https://www.linkedin.com/company/institute-for-health-metrics-and-evaluation">https://www.linkedin.com/company/institute-for-health-metrics-and-evaluation</a>