



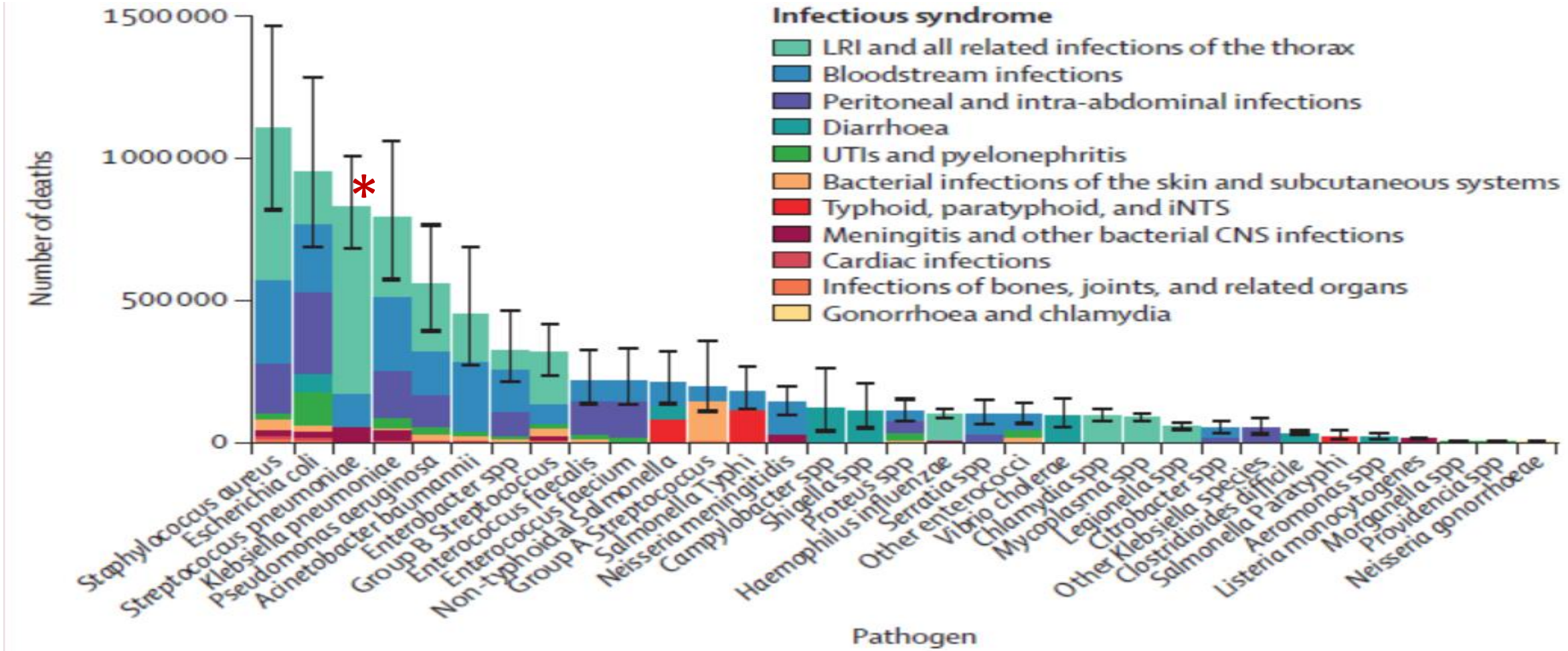
Laboratory Diagnosis of Pneumonia

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Global number of deaths by pathogen and infectious syndrome, 2019



GBD 2019 Antimicrobial Resistance Collaborators. Global mortality associated with 33 bacterial pathogens in 2019: a systematic analysis for the Global Burden of Disease Study 2019. Lancet. 2022 Dec 17;400(10369):2221-2248. doi: 10.1016/S0140-6736(22)02185-7. Epub 2022 Nov 21. PMID: 36423648; PMCID: PMC9763654.

Preventing and effectively treating infections

Achieving Sustainable Development Goal (SDG) 3

THE 17 GOALS | **169** Targets | **3581** Events | **1342** Publications | **6752** Actions

1 NO POVERTY
Icon: Family of four

2 ZERO HUNGER
Icon: Bowl with steam

3 GOAL 3: Ensure healthy lives and promote well-being for all at all ages.
13 Targets, 44 Publications, 25 Events, 1021 Actions
More Info

4 QUALITY EDUCATION
Icon: Open book and pencil

5 GENDER EQUALITY
Icon: Gender symbols

6 CLEAN WATER AND SANITATION
Icon: Water tap

7 AFFORDABLE AND CLEAN ENERGY
Icon: Sun with gear

8 DECENT WORK AND ECONOMIC GROWTH
Icon: Bar chart with upward arrow

9 INDUSTRY, INNOVATION AND INFRASTRUCTURE
Icon: Three cubes

10 REDUCED INEQUALITIES
Icon: Scales of justice

11 SUSTAINABLE CITIES AND COMMUNITIES
Icon: Buildings

12 RESPONSIBLE CONSUMPTION AND PRODUCTION
Icon: Infinity symbol

13 CLIMATE ACTION
Icon: Eye with globe

14 LIFE BELOW WATER
Icon: Fish

15 LIFE ON LAND
Icon: Tree and bird

16 PEACE, JUSTICE AND STRONG INSTITUTIONS
Icon: Dove and gavel

17 PARTNERSHIPS FOR THE GOALS
Icon: Interlocking circles

SUSTAINABLE DEVELOPMENT GOALS
See all
<https://sdgs.un.org>



**TARGET
3.2**

By 2030, end preventable deaths of newborns and children under 5 years of age, with all countries aiming to reduce neonatal mortality to at least as low as 12 per 1,000 live births and under-5 mortality to at least as low as 25 per 1,000 live births

Accurate diagnosis and attribution of the causes of pneumonia

- Measuring the burden of disease,
- Implementing appropriate preventive or treatment strategies, and
- Developing more effective interventions

Pathogenesis of Pneumonia

Former **thought**

- Invasion of a sterile LRT by a single pathogenic organism

Recent evidence

- Healthy lung is not sterile,
- Normal lung microbiome exists in a dynamic state
- Dysbiosis in the normal microbial ecology
- Overgrowth of a single pathogen or multiple pathogens
- it is hard to distinguish colonizing flora from pathogenic organisms in respiratory samples.

Global childhood immunization programs of vaccines against bacterial pathogens have had an impact on the causes of childhood pneumonia

Current guidelines for microbiological diagnosis

- British Thoracic Society ((BTS) guidelines for community acquired pneumonia (CAP) recommend microbiological testing only in:
 - Children with severe pneumonia who need to be admitted to intensive care or those with complications.
 - In such cases, a variety of diagnostic tests are suggested including;
 - blood culture;
 - nasal specimens for viral detection;
 - acute and convalescent serology for viruses and atypical organisms; and
 - pleural fluid (if present) for microscopy, culture, antigen detection, and polymerase chain reaction.

- Guidelines from those of the Pediatric Infectious Diseases Society and the Infectious Diseases Society of America (IDSA), which include:
 - rapid diagnostic testing for influenza virus and other respiratory viruses in children with CAP who are being treated in the community or in hospital,
 - For children with CAP who are in hospital,
 - blood culture in those with complicated or moderate to severe disease and
 - sputum for Gram stain and culture in those who can produce a specimen.
 - tracheal aspirates for culture and detection of viral pathogens are recommended in children who have been intubated,
 - bronchoscopy, bronchoalveolar lavage, or lung biopsy are recommended only in those with severe pneumonia who have negative diagnostic tests and
 - targeted testing may be warranted in outbreak situations, or if infection with specific pathogens, such as *M tuberculosis* or *Bordetella pertussis*, is suspected.

Value of different specimen types and associated diagnostic tests

- Optimal specimen collection is key to accurate identification of the cause of pneumoniae.
- Diagnostic yield depends on:
 - the type and quality of the specimen collected,
 - timing of collection,
 - previous use of antimicrobials,
 - transport, storage, and laboratory processes for testing.
- It may be difficult to obtain a representative sample from the lower respiratory tract, and contamination with organisms such as *Streptococcus pneumoniae* or *Haemophilus influenzae*, which colonize the upper airways but are also important pathogens, is likely. This may lead to overestimation or underestimation of the contribution of these bacteria to pneumonia.

Specimen types and laboratory tests for the causes of pneumonia in children

- **Nasopharyngeal swab or aspirate**
 - Bacterial culture, molecular or antigen detection of bacteria and viruses
 - Sputum or induced sputum
 -
- **Bronchoalveolar lavage or aspirate**
 - Bacterial culture, molecular or antigen detection of bacteria and viruses
- **Transthoracic lung aspirate**
 - Bacterial culture, molecular or antigen detection of bacteria and viruses
- **Blood** (blood culture is indicated only in children who are in hospital, in intensive care, or have complicated pneumonia.)
 - Culture for bacterial pathogens
 - Molecular testing
 - Serology
 - Biomarker detection
- **Urine** (high sensitivity (close to 100%) for invasive pneumococcal infection but lacks specificity (60-80%), particularly in children with nasopharyngeal pneumococcal carriage)
 - Antigen detection

Laboratory tests in children with ambulatory or severe pneumonia

| Test | Indications | Limitations |
|--|--|--|
| <i>Children with mild or ambulatory pneumonia</i> | | |
| Serum for C reactive protein or procalcitonin | Limited evidence that use in an algorithm for children with mild or uncomplicated pneumonia may reduce antibiotic exposure | Data are very limited; studies include longitudinal monitoring of kinetics of response to treatment; requires a blood specimen; not currently recommended to distinguish bacterial disease |
| Upper respiratory tract samples for rapid viral detection of influenza, for example | During outbreaks of viral illness, rapid testing may reduce antibiotic prescription; guidelines differ on recommendation | Bacterial coinfection cannot be excluded; vigilance for coexisting bacterial disease required |
| <i>Children with severe disease, complications, and those in hospital</i> | | |
| Blood culture | All children | Low yield but higher in HIV infected children; cost effectiveness questionable |
| Upper or lower respiratory tract samples for rapid viral testing (PCR or antigen detection) | Children with severe or complicated pneumonia | Difficult to infer causality for many pathogens; most useful for influenza virus and RSV |
| Induced or expectorated sputum for mycobacterial culture and PCR (Xpert MTB/RIF) for <i>Mycobacterium tuberculosis</i> | Children in whom pulmonary tuberculosis is clinically suspected | Repeated specimens needed to optimize microbiologic yield |
| Induced or expectorated sputum or upper respiratory tract sample for PCR for <i>Bordetella pertussis</i> | Children in whom pertussis is suspected | Upper respiratory samples have a lower yield than sputum |
| Induced or expectorated sputum for PCR for <i>Pneumocystis jirovecii</i> | Children in whom pneumocystis pneumonia is suspected | May not distinguish colonizing from disease causing organisms |
| Acute and convalescent serum for respiratory pathogens and mycoplasma and chlamydia | Children with severe or complicated pneumonia | Clinical utility questionable, given need for comparison of acute and convalescent sera |
| Pleural fluid microscopy, culture, and pneumococcal antigen detection or PCR | If pleural effusion present | |
| Tracheal aspirate for bacterial culture; PCR for pneumocystis and viral pathogens | Intubated children | May not distinguish colonizing from disease causing organisms |
| Bronchoscopy, bronchoalveolar lavage, or lung biopsy for bacterial culture; PCR for pneumocystis and viral pathogens; culture and PCR for <i>M tuberculosis</i> (in epidemiological context) | Children with severe illness in whom no pathogen has been detected or those who have not responded to treatment | Invasive and requires specific expertise |

Diagnosis pneumococcal disease

- Suspect serious pneumococcal disease, like meningitis or bloodstream infections, collect samples of cerebrospinal fluid or blood.
- Growing the bacteria in a laboratory helps identify the specific type of bacteria causing the infection.
- Possible use a urine test to help make a diagnosis of **pneumococcal pneumonia** in adults.

Diagnosis *Mycoplasma pneumoniae*

- CDC uses multiplex real-time polymerase chain reaction (PCR) as the primary laboratory procedure for *M. pneumoniae* identification.
- CDC does not use culture or serological testing as routine diagnostic methods. Specimens identified as positive for *M. pneumoniae* by previous lab testing and/or upon testing at CDC will be subsequently tested for macrolide susceptibility using molecular test methods.
 - A Dutch study found that *Mycoplasma pneumoniae* DNA was not detected any more often in children with any respiratory tract infection than in control children.
 - Two recent studies, a birth cohort in South Africa and a multicenter study in the US have reported similar findings. RSV, influenza virus, and *B pertussis* have been consistently and strongly associated with pneumonia in many studies.

There is considerable heterogeneity with regard to the causative role of many other viruses and bacteria detected in respiratory samples, which may relate to case definitions, sampling, and the inclusion of a relevant control group.

Antimicrobial resistance in
S. pneumoniae and
M. pneumoniae



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Burden of *Streptococcus pneumoniae* and *Haemophilus influenzae* type b disease in children in the era of conjugate vaccines: global, regional, and national estimates for 2000–15

[Brian Wahl, PhD](#)   • [Prof Katherine L O'Brien, MD](#) • [Adena Greenbaum, MD](#) • [Anwasha Majumder, MHS](#) •

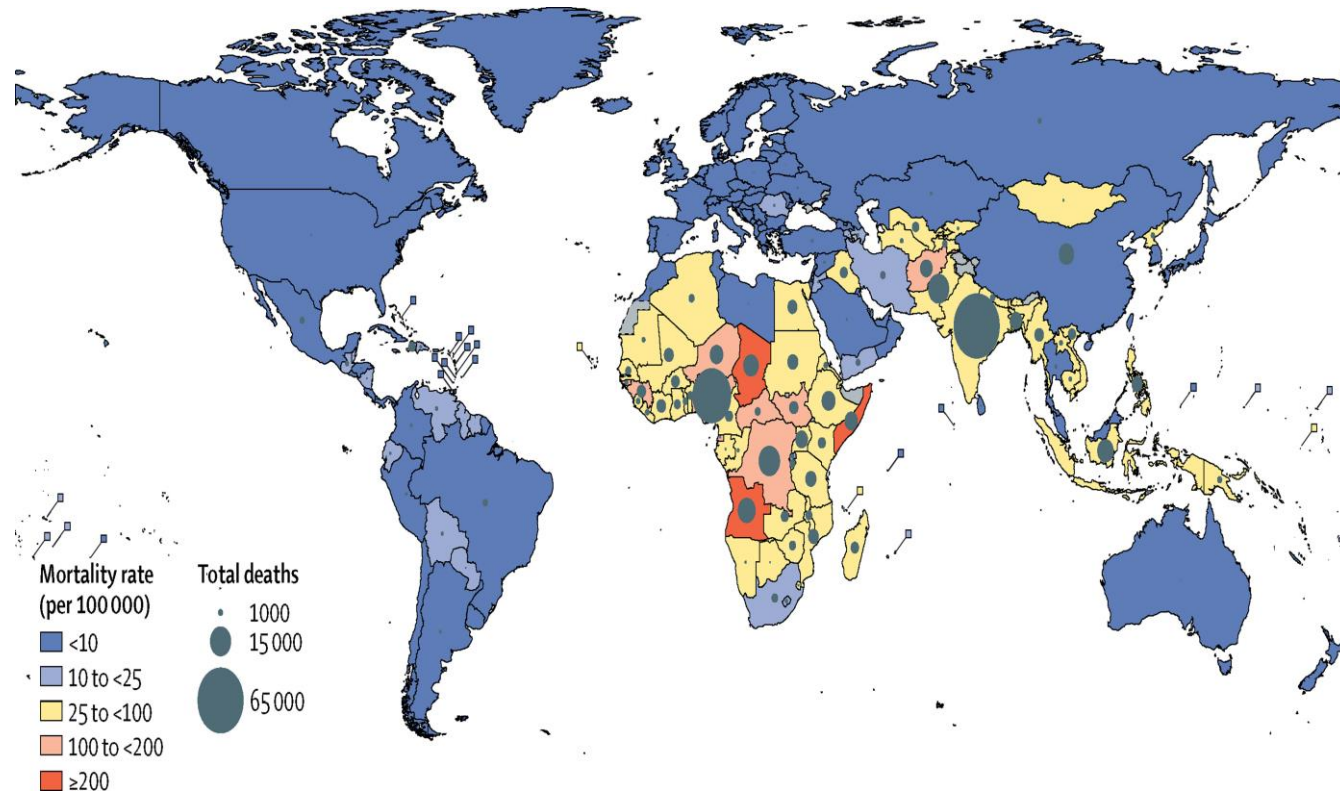
[Li Liu, PhD](#) • [Yue Chu, MSPH](#) • et al. [Show all authors](#)

[Open Access](#) • Published: July, 2018 • DOI: [https://doi.org/10.1016/S2214-109X\(18\)30247-X](https://doi.org/10.1016/S2214-109X(18)30247-X) •



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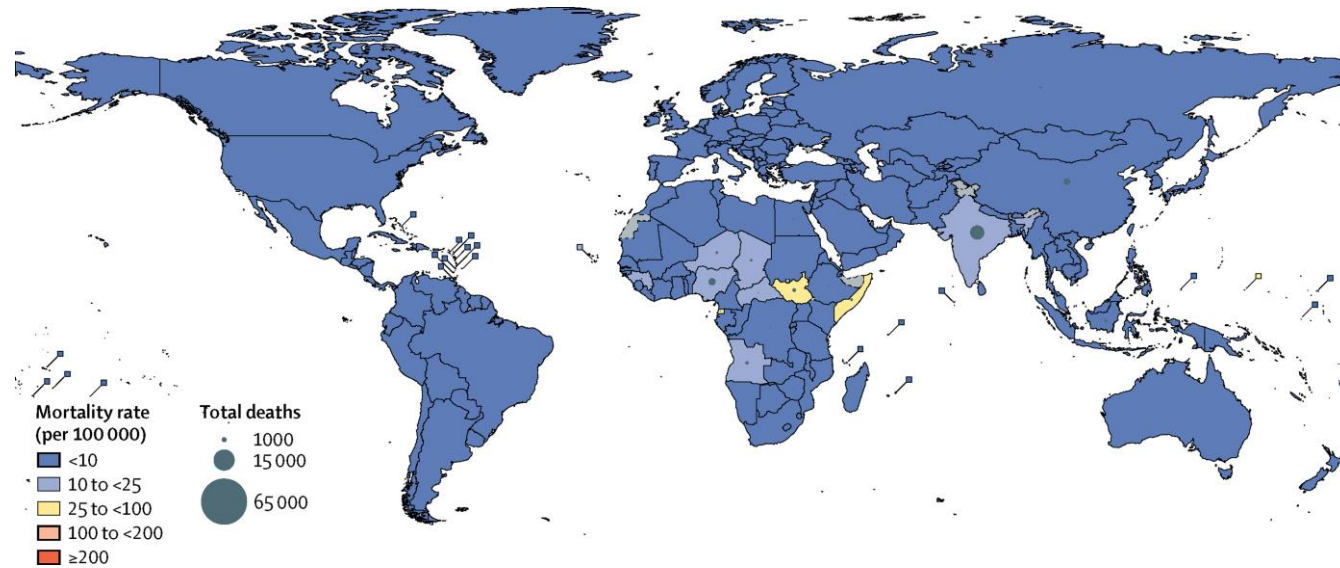




Country-specific mortality rates and deaths attributable to pneumococcus in 2015.

Mortality rates and deaths in children aged 1–59 months are HIV-negative deaths only. Mortality rates are deaths per 100 000 children aged 1–59 months. Pneumococcus=*Streptococcus pneumoniae*.





Country-specific mortality rates and deaths attributable to Hib in 2015.

Antimicrobial resistance in paediatric *Streptococcus pneumoniae* isolates amid global implementation of pneumococcal conjugate vaccines: a systematic review and meta-regression analysis

Kristin Andrejko, BS • Buddhika Ratnasiri, BA • William P Hausdorff, PhD • Ramanan Laxminarayan, PhD

Joseph A Lewnard, PhD

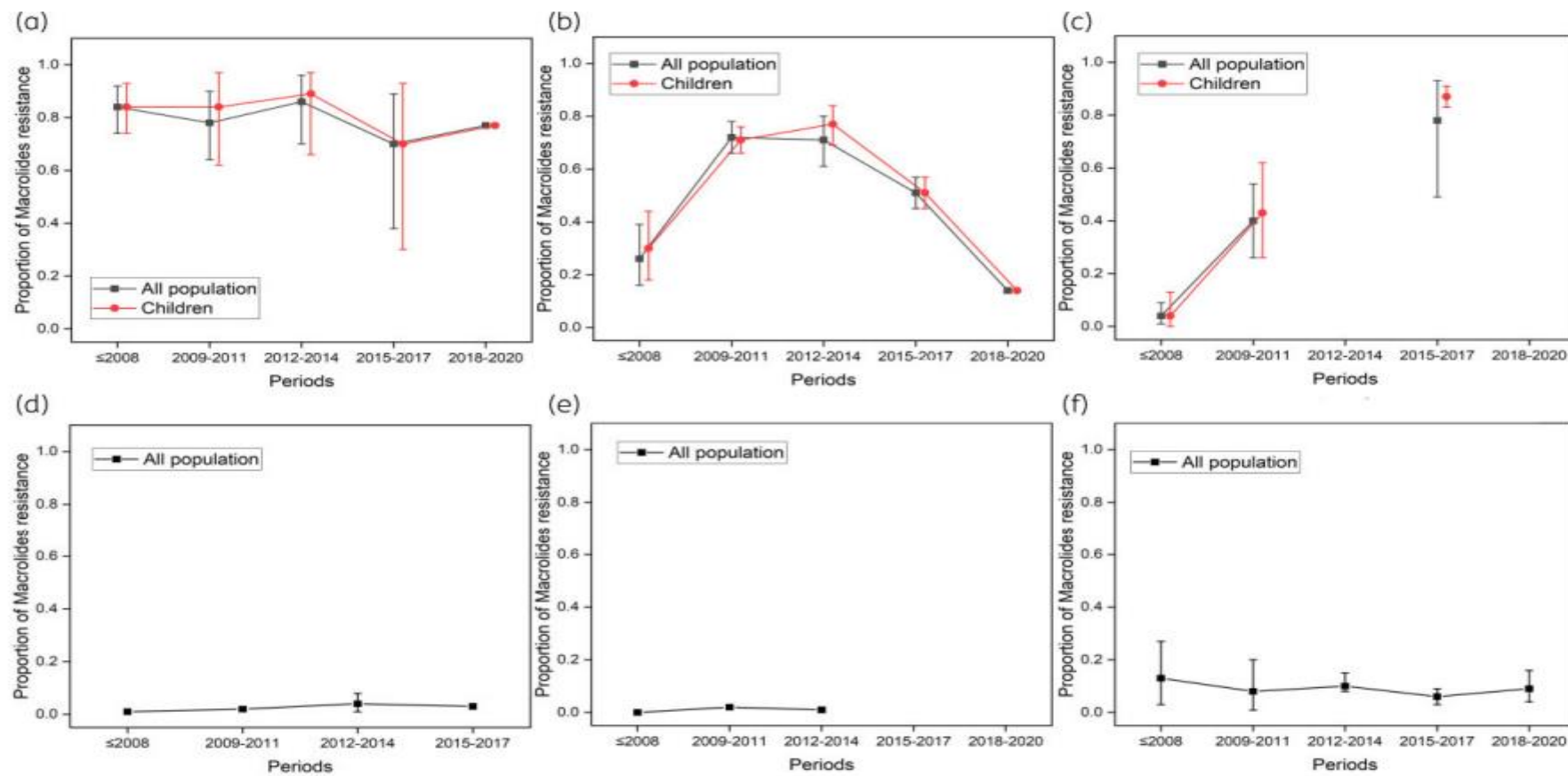


Figure 3. Prevalence change over time in (a) China, (b) Japan, (c) Korea, (d) Germany, (e) Slovenia and (f) America. This figure appears in colour in the online version of *JAC* and in black and white in the print version of *JAC*.

- Circulating penicillin-resistant *S. pneumoniae* are much more heterogeneous.
- Internationally, penicillin and macrolide resistance were increasing at the turn of the 21st century.
- However, infection rates overall and with penicillin-resistant serotypes significantly decreased following the introduction of pneumococcal vaccines.
- Early U.S. studies noted an initial decrease in pneumococcal activity after the introduction of 7-valent vaccine (PCV7), with residual high resistance by serotypes not covered by PCV7. The 13-valent vaccine (PCV13) coverage of serotype 19A led to a decrease in one of the most pathogenic (and resistant) circulating serotypes, and since 2010, there has been an overall downward trend in non-susceptible invasive pneumococcal infections.
- Regional differences in circulating antibiotic-resistant *S. pneumoniae* remain; however, this resistance does not appear related to a dominant non-vaccine serotype.

Antibiotic resistance of *S. pneumonia* in Iran

- The 44 pneumococcal invasive isolates, 39 (89%) were isolated from children and 5 (11%) from adults.
 - All pneumococcal isolates were susceptible to linezolid but had varying resistance to trimethoprim-sulfamethoxazole (86%), erythromycin (73%), tetracycline (66%), clindamycin (43%), penicillin (16%), chloramphenicol (14%) and levofloxacin (2%).
 - All of the penicillin resistant isolates were multidrug resistant (MDR) and in addition to penicillin were resistant to tetracycline, erythromycin and trimethoprim-sulfamethoxazole.
 - The most common capsular types detected in 64% of the pneumococcal isolates was 6A/B, 19A, 15A, 23F.
 - The co resistance to erythromycin and clindamycin (the constitutive phenotype) was observed in 43% (19/44) of the isolates and resistance to erythromycin, but not to clindamycin (the M phenotype) was observed in 29.5% (13/44) of the isolates.

- Evaluating the level and type of *S. pneumoniae* resistance to macrolides in Iran 2000–2017:
 - The results of the present study showed that resistance to macrolides is considerable in different parts of Iran.
 - Half of previous studies have reported prevalence rates of more than 50%.
 - Similar to other countries around the world, especially European countries, 23S rRNA methylation was the dominant mechanism in the development of macrolide resistance.
 - In addition, efflux-mediated resistance (independent or in combination with *ermB* gene) was considerable, as well.
- A study by Haghi Ashtiani et al., evaluating the trend of changes in resistance over 10 years, showed that levels of antibiotic resistance are increasing rapidly. For instance, resistance to macrolides, such as erythromycin, has increased from 25 to 65%.

Global prevalence of resistance to macrolides in *Mycoplasma pneumoniae*: a systematic review and meta-analysis

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Table 1. Characteristics of included studies

| | Macrolide resistance |
|---|-------------------------------|
| Sampling years | 2000–20 |
| Geographical regions | |
| Asia | * 63% [56, 69], (8628/13 721) |
| Europe | 3% [2, 7], (71/2285) |
| North America | 8.6% [6, 11], (136/1795) |
| South America | 0% (0/42) |
| Oceania | 3.3% (1/30) |
| Annual data ^a | |
| ≤2008 | 27% [15, 40], (491/2090) |
| 2009–11 | 50% [39, 61], (3209/5630) |
| 2012–14 | 51% [35, 68], (2551/4540) |
| 2015–17 | 39% [24, 57], (1626/3545) |
| 2018–20 | 23% [2, 54], (218/427) |
| Mutation (Total resistant sample detected for point mutation: 8257) | |
| A2063G | 96% [94, 98], (7565) |
| A2063C | 0.2% (17) |
| A2063T | 5% [2, 10], (257) |
| A2064G | 4% [2, 6], (213) |
| A2064C | 0.01% (1) |
| C2617A | 0.04% (4) |
| C2617G | 0.13% (11) |
| C2617T | 0.04% (3) |
| No mutation | 2% (186) |

The data are presented as mean prevalence [95% CI].

^aNot all studies reported year-stratified prevalence data.

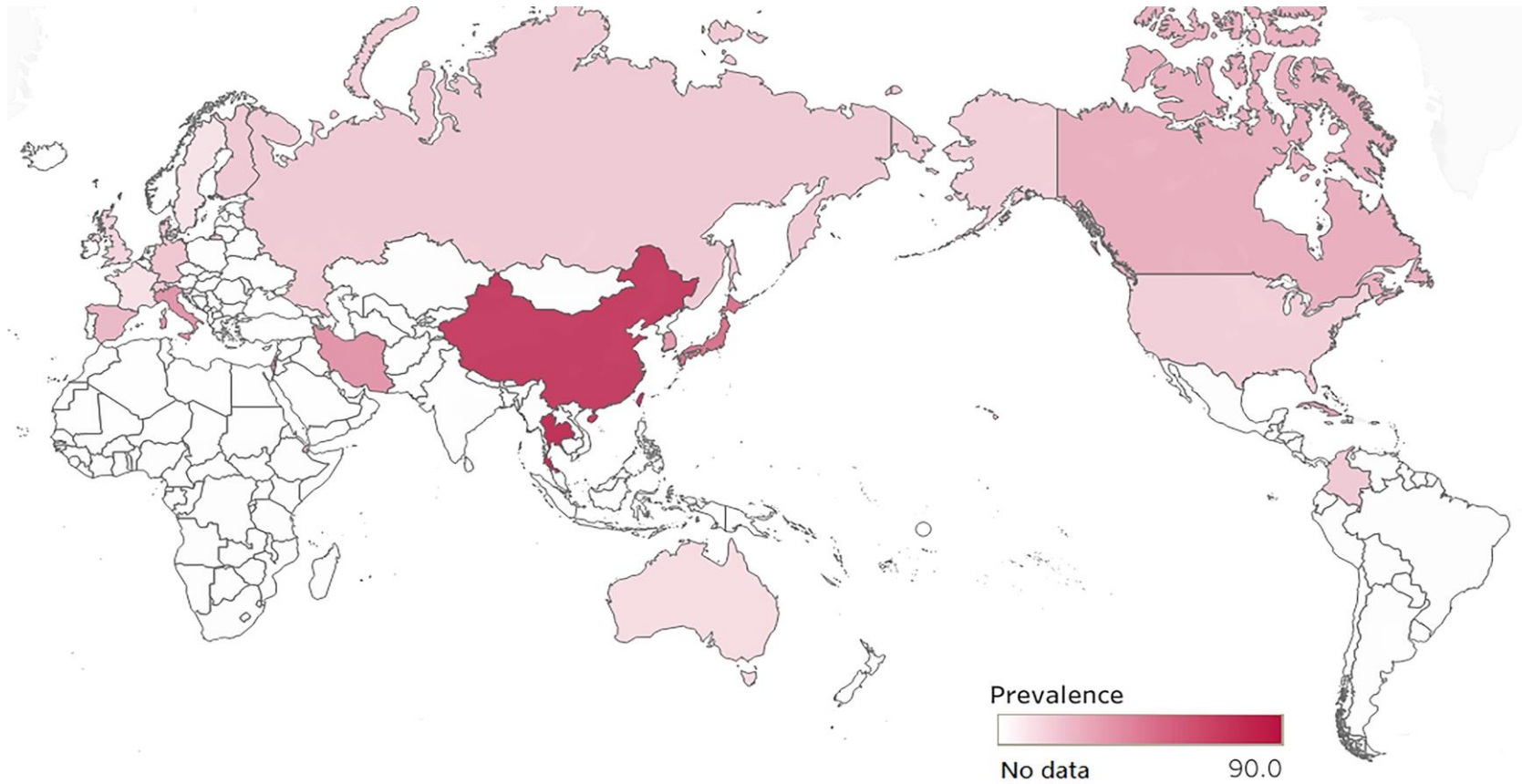


Figure 1. The mean prevalence map of macrolide resistance all over the world (from ≤ 2008 to 2020).

MIC (mg/L) range of isolated *Mycoplasma pneumoniae*

| Study | Mutation in 23S rRNA | ERY (MIC of reference strain) | CLR | AZM | TET | DOX | MIN | CIP | LVX | MXF | GAT | JOS |
|-------------|----------------------|----------------------------------|--------------------------|--------------------------|----------------------|----------------------|----------------------|-------------------|-------------------|-----------------------|------------------------|----------------------|
| China | | | | | | | | | | | | |
| Zhao F 2013 | A2063G | 128~>256 | 64~>256 | 2~64 | 0.032~0.5 | | | 0.125~2 | 0.125~2 | | 0.016~0.125 | |
| | A2064G | 256~>256 | 256~>256 | 4~32 | 0.125~0.25 | | | 0.5~1 | 0.25~1 | | 0.016~0.125 | |
| | A2063T | 32 | 16 | 0.064 | 0.25 | | | 0.5 | 0.25 | | 0.064 | |
| | None (M129)* | 0.008~0.016 (0.016) | <0.008~0.008 (<0.008) | <0.008~0.008 (<0.008) | 0.016~0.5 (0.125) | | | 0.008~1 (1) | 0.008~1 (1) | | 0.008~0.125 (0.125) | |
| Xin D 2009 | Not specified (FH) | 0.01~512 (0.01) | | 0.001~256 (0.001) | | | | | | | | 0.01~64 (0.01) |
| Liu Y 2009 | Not specified | ≤0.007~>128 (≤0.007) | ≤0.007~>128 (≤0.007) | ≤0.007~>128 (≤0.007) | 0.015~0.25 (0.06) | ≤0.007~125 (0.03) | ≤0.007~125 (0.06) | 0.015~1 (0.25) | 0.015~1 (0.25) | ≤0.007~0.06 (0.06) | | ≤0.007~8 (≤0.007) |

MIC ≤0.5 mg/L was considered as the breakpoint to discriminate erythromycin susceptible or resistant in CLSI guideline.

All *M. pneumoniae* with the **A2063G** mutation were resistant to **erythromycin** (32-1024 mg/L) and **clarithromycin** (32-256 mg/L), with some also resistant to **azithromycin** (≤0.007.64 mg/L);

The similar finding was suggested for the **A2064G** mutation (**erythromycin**: 64-256 mg/L; **clarithromycin**: 16-256 mg/L; **azithromycin**: 2-32 mg/L). Additionally, the **A2063T** mutation might not be associated with resistance to **azithromycin** (0.0642 mg/L).

- Two main challenges: a relatively high carriage in asymptomatic children, and a worldwide increase in macrolide-resistant *M. pneumoniae* (MRMP). This review focuses on the scientific and clinical implications of these crucial issues.
- Asia is the most common location for macrolide-resistant *M. pneumoniae*, and both erythromycin and clarithromycin are no longer effective in treating resistant pathogens.
- Azithromycin may be an option for treating susceptible *M. pneumoniae*, but its use should also be restricted to avoid building resistance via widespread use.

