

## INVASIVE STREPTOCOCCAL DISEASE IN GERMANY DURING THE SARS-COV-2 PANDEMIC

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### BACKGROUND

The GNRCS monitors invasive streptococcal disease in Germany. This includes *Streptococcus pneumoniae*, *Streptococcus pyogenes* (GAS), *Streptococcus agalactiae* (GBS), *Streptococcus dysgalactiae* (SD) and all other validly published streptococcal species. Here, we report on the species-specific reduction of invasive streptococcal disease during the SARS-CoV-2 pandemic in Germany.

### METHODS

Species identification was performed using a combination of microbiological, biochemical, immunological and molecular-biological techniques, including hemolysis-assessment, catalase-test, optochine- and bile-susceptibility, pyrrolidonyl-arylamidase-test, leucine-aminopeptidase-test, Lancefield-typing, *emm*-typing, serotyping and multiple PCR-sequence analyses.

### RESULTS

SARS-CoV-2 reached Germany beginning of March 2020. A strong reduction in case numbers of invasive pneumococcal disease (IPD), both among children (Fig. 1) as well as among adults (Fig. 2) was reported shortly after. This effect continued till October 2021, and resulted in a reduction of 75%. With retraction of the non pharmaceutical intervention measures in the autumn of 2021, IPD case numbers increased to pre-pandemic levels. For children case numbers remained on pre-pandemic levels in 2022, For adults IPD case numbers were once again much lower than pre-pandemic levels in the first months of 2022.

A comparable reduction of 67% was observed for invasive GAS infections (Fig. 3), accompanied with a disproportional decrease of *emm1* (-85%) and *emm12* (-73%). However, no effect was seen for GBS (Fig. 4) and SD (Fig. 6).

Among viridans streptococci, a 40% reduction was observed for oral-streptococci of the Mitis-, Salivarius- and Mutans-Group (Fig. 5), while streptococcal species not commonly associated with the oral microbiome, were not affected (Fig. 7).

Cases of invasive disease caused by streptococci belonging to the Anginosus-group (*S. anginosus*, *S. constellatus*, *S. intermedius*) also showed no reduction (Fig. 8).

### CONCLUSIONS

- The SARS-CoV-2 pandemic had a strong reducing effect on invasive streptococcal disease caused by *S. pneumoniae*, *S. pyogenes* and oral streptococci of the Mitis-, Salivarius- and Mutans-Group
- No comparable effect was observed for invasive GBS and SD disease or for disease caused by Anginosus-group streptococci.
- These reductions in case numbers are obviously limited to respiratory transmitted streptococci, and therefore seem to be related to non-pharmaceutical interventions (face masks, social distancing, working from home, school closures) during the SARS-CoV-2 pandemic.
- Our results might also be explained by differences in dependence on preceding viral infections of the different streptococcal species.

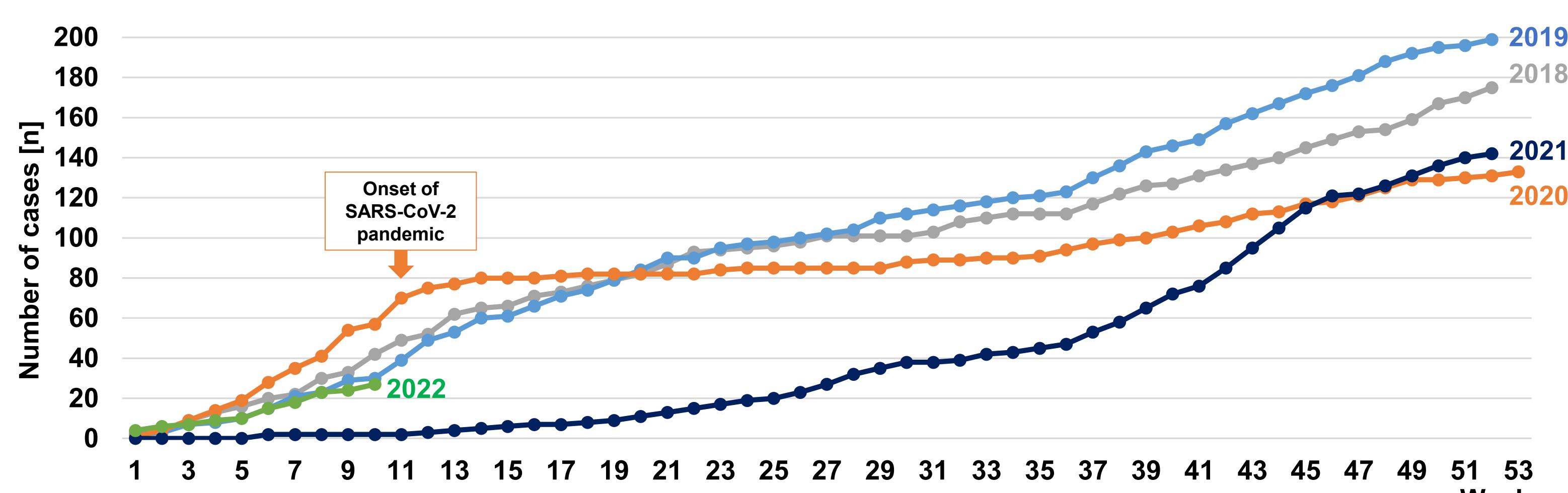


Fig. 1: Cumulative number of reported IPD cases from children <18 years of age in Germany

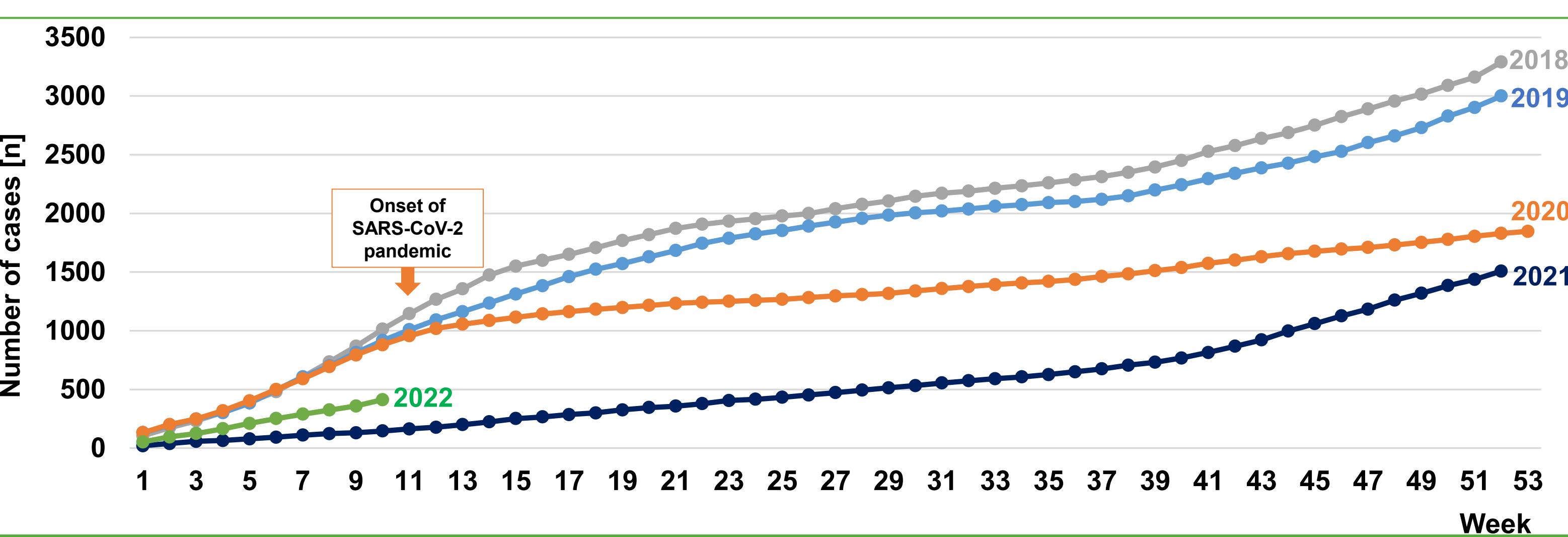


Fig. 2: Cumulative number of reported IPD cases from adults ≥18 years of age in Germany

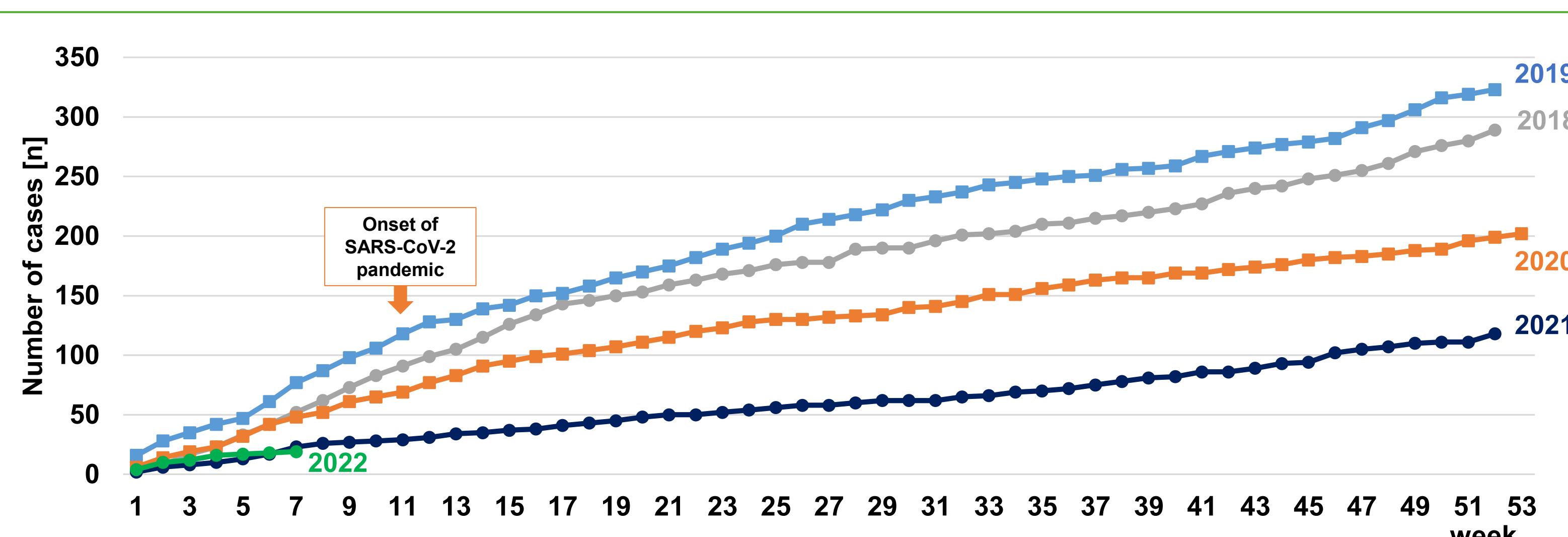


Fig. 3: Cumulative number of reported invasive Group-A streptococcal disease cases among all ages in Germany.

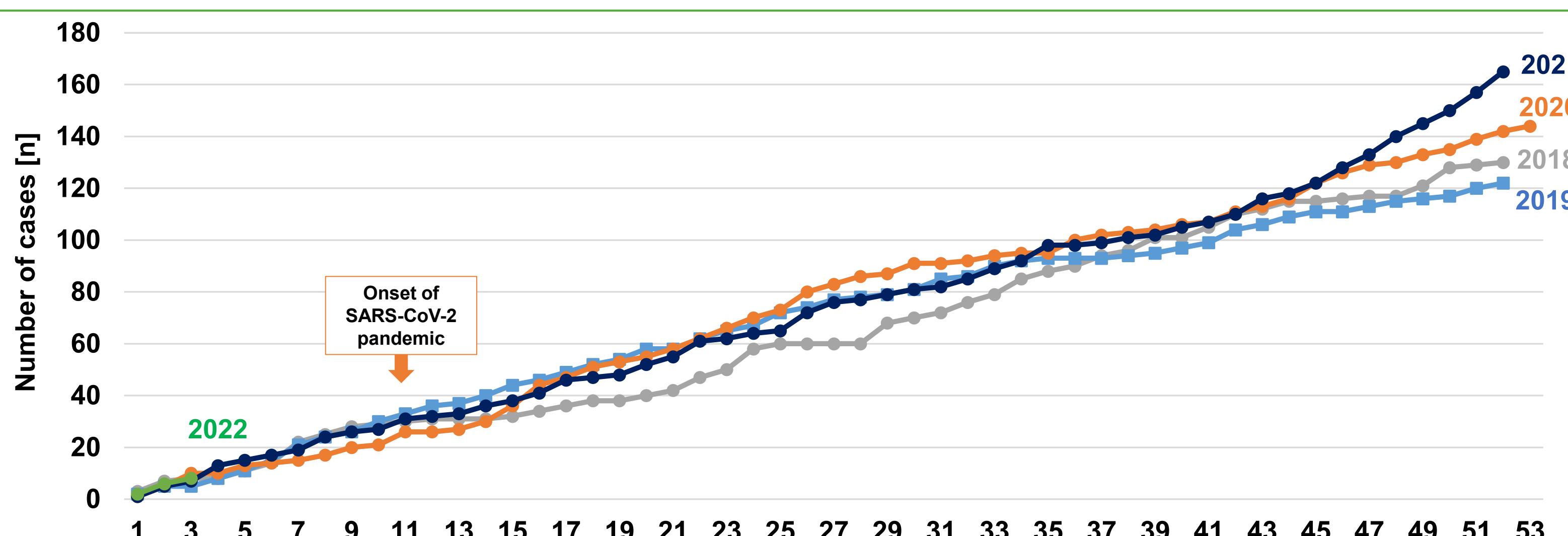


Fig. 4: Cumulative number of reported invasive Group-B streptococcal disease cases among all ages in Germany.

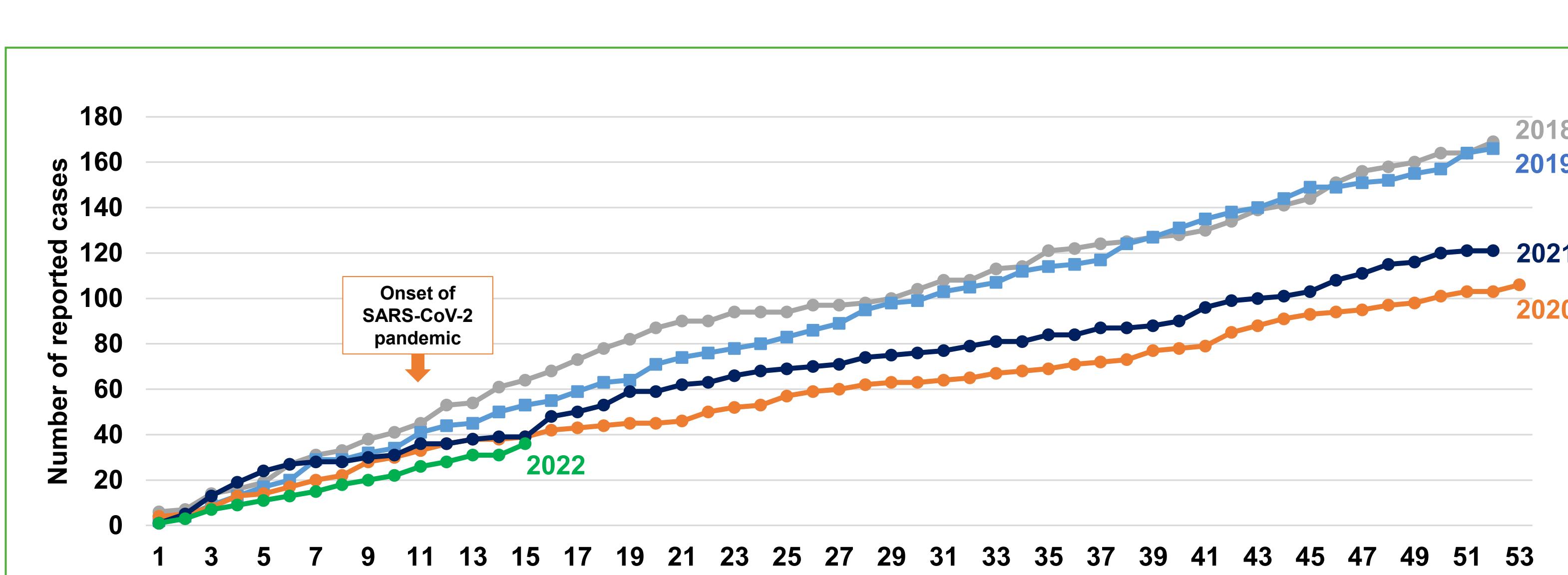


Fig. 5: Cumulative number of reported invasive infections with streptococci of the Mitis-, Salivarius- and Mutans-group among all ages in Germany

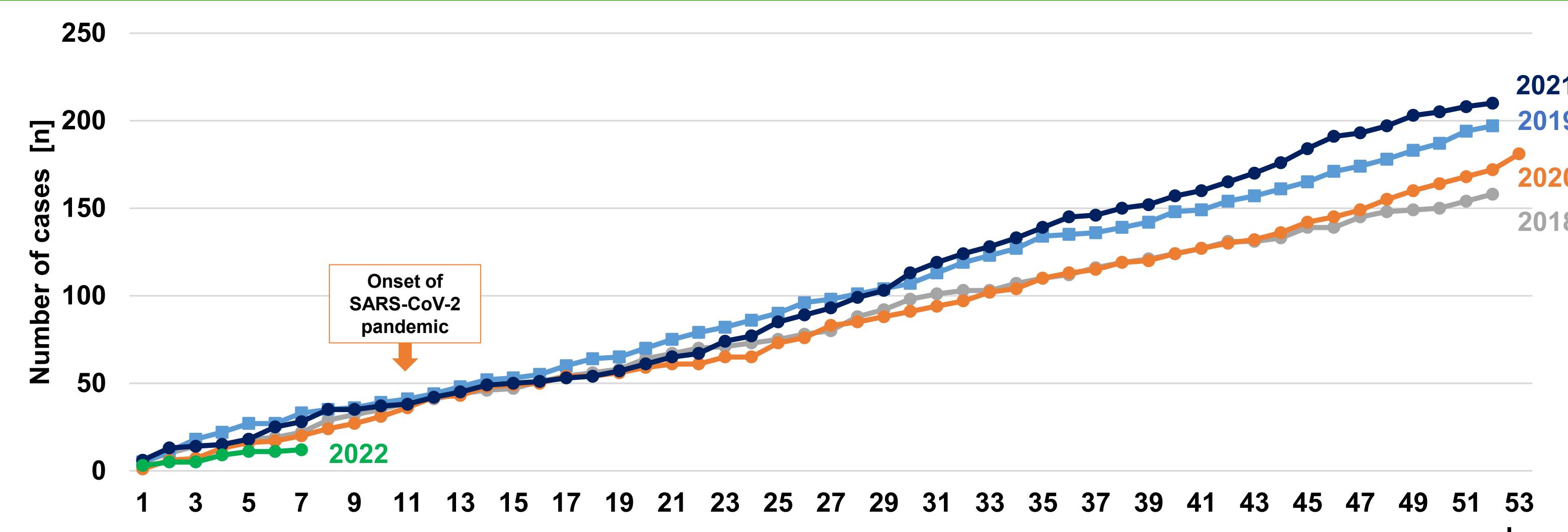


Fig. 6: Cumulative number of reported invasive *Streptococcus dysgalactiae* disease cases among all ages in Germany

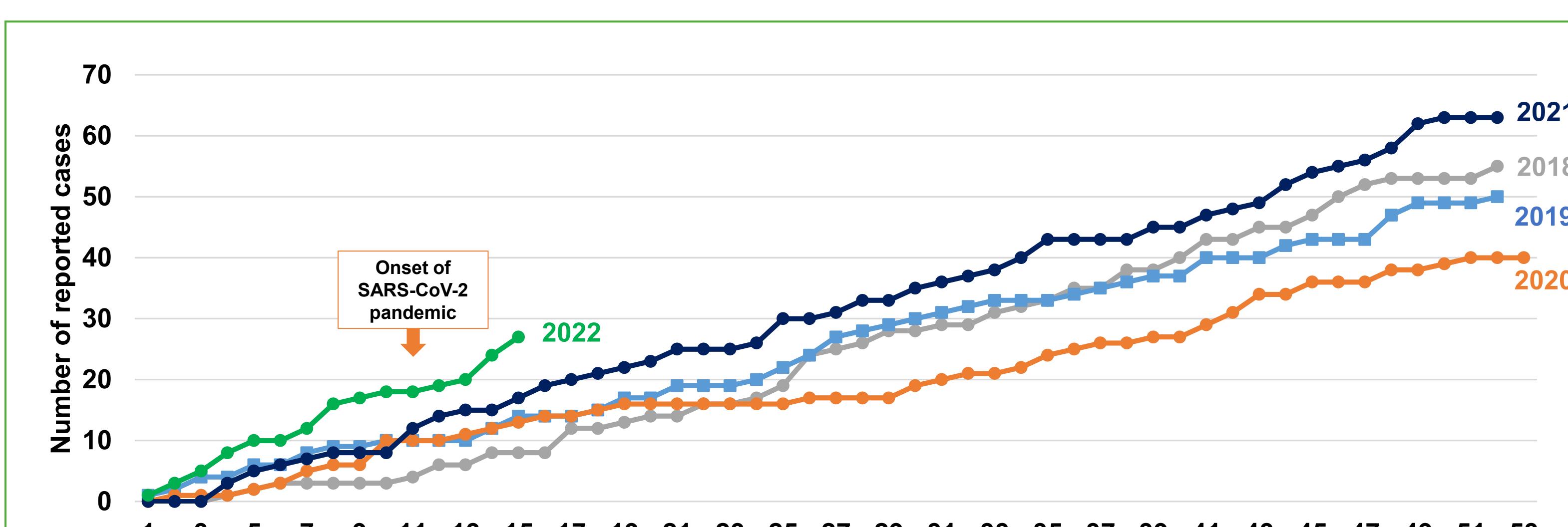


Fig. 7: Cumulative number of reported invasive infections with viridans streptococci NOT belonging to the Mitis-, Salivarius- and Mutans-group among all ages in Germany

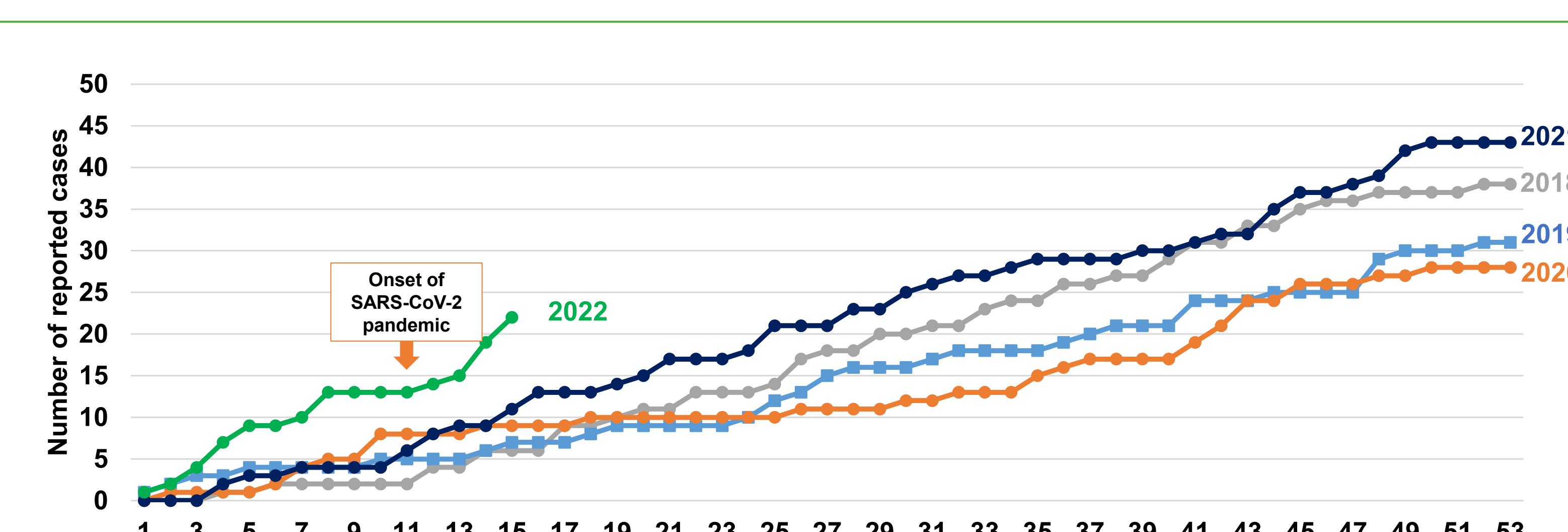


Fig. 8: Cumulative number of reported invasive infections with streptococci of the Anginosus-group among all ages in Germany