



Pathogenwatch

for *Streptococcus pneumoniae*

One-stop platform for pneumococcal genomics: from analyses to visualisation

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⁹<https://www.pneumogen.net/gps/>

We have tailored the web application Pathogenwatch to offer *in silico* detection and characterisation of whole-genome sequence (WGS) data from *Streptococcus pneumoniae*.

(A) Drag and drop assemblies/raw sequence data to obtain the following output and manage them through your personal account. Uploaded data are kept private by default and metadata can be integrated without upload.

Pathogenwatch's core features:

- Assembly quality metric
- Species prediction
- MLST
- Antimicrobial resistance determinant detection

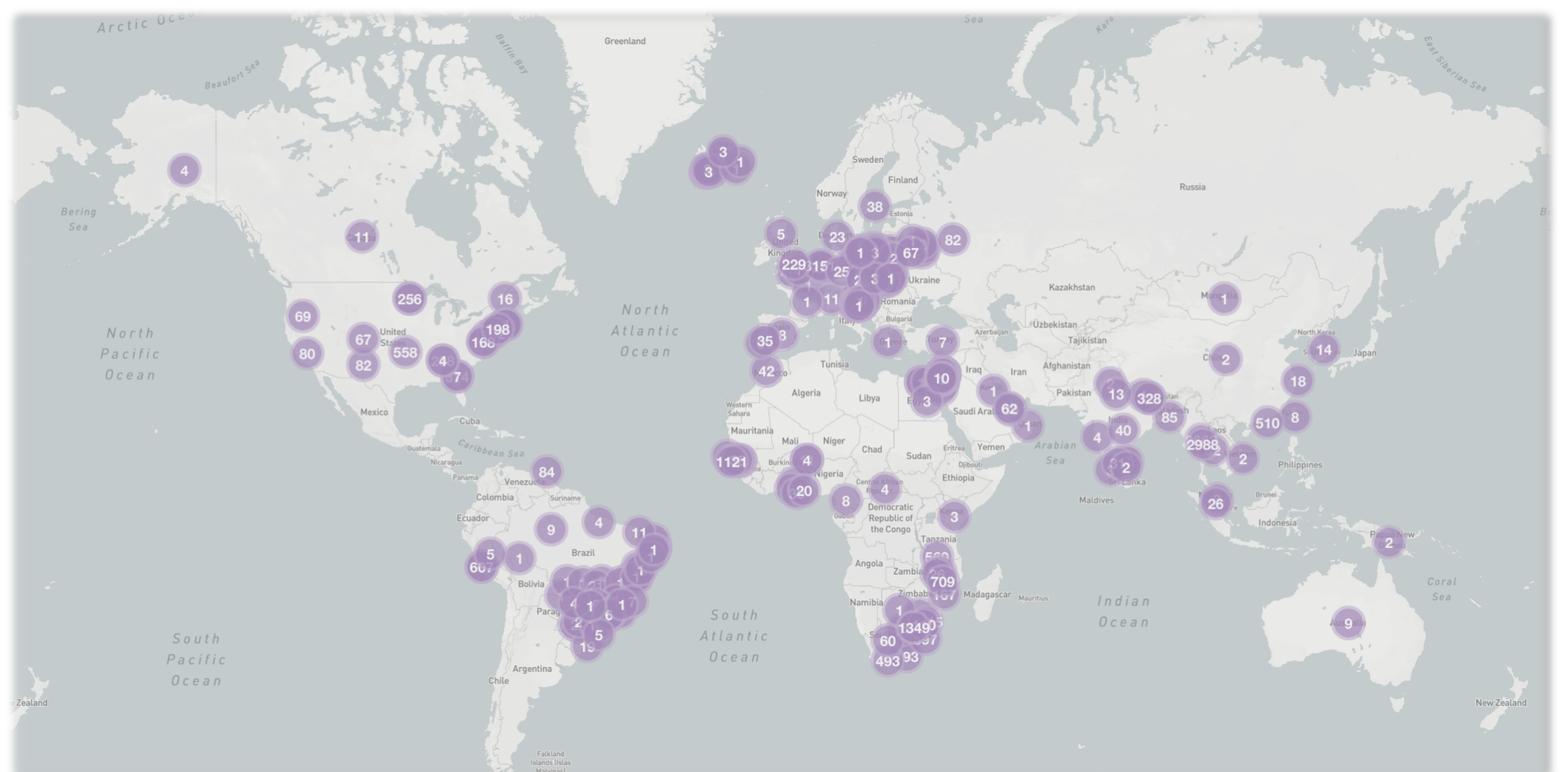
Pneumococcus-specific features:

- Global pneumococcal Sequencing Cluster (GPSC) using PopPUNK
- *In silico* serotype using SeroBA

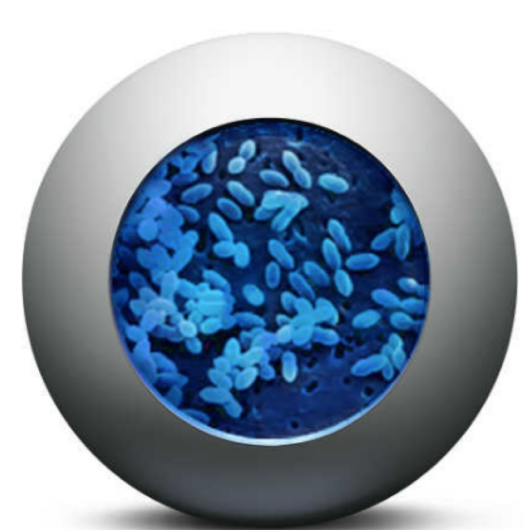
(B) Access and download publicly available >20,000 pneumococcal genomes, including ~13,000 generated from the Global Pneumococcal Sequencing (GPS) project.



A step-by-step tutorial video on the GPS website



Pneumo Pathogenwatch is brought to you by



The Global Pneumococcal Sequencing Project

A worldwide genomic survey of the impact of vaccination on the pathogen population



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