

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

Auspice

RRID:SCR_023966

Type: Tool

Proper Citation

Auspice (RRID:SCR_023966)

Resource Information

URL: <https://github.com/nextstrain/auspice>

Proper Citation: Auspice (RRID:SCR_023966)

Description: Web application for visualizing pathogen evolution. Interactive web app for visualizing phylogenomic data.

Synonyms: auspice

Resource Type: software resource, source code

Keywords: visualizing pathogen evolution, visualizing phylogenomic data,

Availability: Free, Available for download, Freely available

Resource Name: Auspice

Resource ID: SCR_023966

Alternate URLs: <https://docs.nextstrain.org/projects/auspice/en/stable/>

Old URLs: <https://sources.debian.org/src/auspice/>

License: GNU Affero General Public License

Record Creation Time: 20230824T050210+0000

Record Last Update: 20260905T133437+0000

Ratings and Alerts

No rating or validation information has been found for Auspice.

No alerts have been found for Auspice.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

Listed below are recent publications. The full list is available at [Kravitz](#) .

Li SH, et al. (2026) Childhood immunological imprinting of cross-subtype antibodies targeting the hemagglutinin head domain of influenza viruses. *Cell host & microbe*, 34(5), 873.

Mbaye A, et al. (2025) Genomic and epidemiological analysis of SARS-CoV-2 variants isolated in Guinea: a routine sequencing implementation. *BMC infectious diseases*, 25(1), 3.

Pitts J, et al. (2025) Oral dosing of the nucleoside analog obeldesivir is efficacious against RSV infection in African green monkeys. *Nature communications*, 16(1), 6437.

Sitharam N, et al. (2024) SARS-CoV-2 Genomic Epidemiology Dashboards: A Review of Functionality and Technological Frameworks for the Public Health Response. *Genes*, 15(7).

Alathari S, et al. (2024) In field use of water samples for genomic surveillance of infectious spleen and kidney necrosis virus (ISKNV) infecting tilapia fish in Lake Volta, Ghana. *PeerJ*, 12, e17605.

Choi CH, et al. (2024) Genomic Analysis of Monkeypox Virus During the 2023 Epidemic in Korea. *Journal of Korean medical science*, 39(18), e165.

Agius JE, et al. (2022) SARS-CoV-2 Within-Host and in vitro Genomic Variability and Sub-Genomic RNA Levels Indicate Differences in Viral Expression Between Clinical Cohorts and in vitro Culture. *Frontiers in microbiology*, 13, 824217.

Padane A, et al. (2022) Rapidly rising cases with Omicron in Senegal. *New microbes and new infections*, 45, 100959.