

# Resource Summary Report

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

## SynTView

RRID:SCR\_011939

Type: Tool

### Proper Citation

SynTView (RRID:SCR\_011939)

### Resource Information

**URL:** <http://genopole.pasteur.fr/SynTView/>

**Proper Citation:** SynTView (RRID:SCR\_011939)

**Description:** An interactive multi-view genome browser for next-generation comparative microorganism genomics.

**Abbreviations:** SynTView

**Resource Type:** software resource

**Keywords:** bio.tools

**Resource Name:** SynTView

**Resource ID:** SCR\_011939

**Alternate IDs:** OMICS\_01500, biotools:syntview

**Alternate URLs:** <https://bio.tools/syntview>

**Record Creation Time:** 20220129T080307+0000

**Record Last Update:** 20260905T132716+0000

### Ratings and Alerts

No rating or validation information has been found for SynTView.

No alerts have been found for SynTView.

### Data and Source Information

## Usage and Citation Metrics

We found 7 mentions in open access literature.

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

L??-Bury P, et al. (2023) Yersiniomics, a Multi-Omics Interactive Database for Yersinia Species. Microbiology spectrum, 11(2), e0382622.

Bussotti G, et al. (2018) Leishmania Genome Dynamics during Environmental Adaptation Reveal Strain-Specific Differences in Gene Copy Number Variation, Karyotype Instability, and Telomeric Amplification. mBio, 9(6).

Gonzalez-Hilarion S, et al. (2016) Intron retention-dependent gene regulation in Cryptococcus neoformans. Scientific reports, 6, 32252.

Drini S, et al. (2016) Species- and Strain-Specific Adaptation of the HSP70 Super Family in Pathogenic Trypanosomatids. Genome biology and evolution, 8(6), 1980.

Glaser P, et al. (2016) Demography and Intercontinental Spread of the USA300 Community-Acquired Methicillin-Resistant Staphylococcus aureus Lineage. mBio, 7(1), e02183.

Bouchez V, et al. (2015) New Data on Vaccine Antigen Deficient Bordetella pertussis Isolates. Vaccines, 3(3), 751.

Lechat P, et al. (2013) SynTView - an interactive multi-view genome browser for next-generation comparative microorganism genomics. BMC bioinformatics, 14, 277.