

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

SynView

RRID:SCR_001106

Type: Tool

Proper Citation

SynView (RRID:SCR_001106)

Resource Information

URL: <http://gmod.org/wiki/SynView>

Proper Citation: SynView (RRID:SCR_001106)

Description: A software tool for comparative visualization of genomes. It is based in the GBrowse frameworks and integrates with the annotation features.

Resource Type: sequence analysis software, software resource, software application, data analysis software, data processing software

Defining Citation: [PMID:16844709](#), [DOI:10.1093/bioinformatics/btl389](#)

Keywords: visualization, genome, comparison, gbrowse, sequence analysis software

Resource Name: SynView

Resource ID: SCR_001106

Alternate IDs: OMICS_00947

Record Creation Time: 20220129T080205+0000

Record Last Update: 20260808T185725+0000

Ratings and Alerts

No rating or validation information has been found for SynView.

No alerts have been found for SynView.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Pavlopoulos GA, et al. (2015) Visualizing genome and systems biology: technologies, tools, implementation techniques and trends, past, present and future. GigaScience, 4, 38.