

Resource Summary Report

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

[GBrowse syn](#)

RRID:SCR_012971

Type: Tool

Proper Citation

GBrowse syn (RRID:SCR_012971)

Resource Information

URL: http://gmod.org/wiki/GBrowse_syn

Proper Citation: GBrowse syn (RRID:SCR_012971)

Description: A GBrowse-based synteny browser designed to display multiple genomes, with a central reference species compared to two or more additional species.

Abbreviations: GBrowse_syn

Synonyms: Generic Synteny Browser

Resource Type: software resource

Resource Name: GBrowse syn

Resource ID: SCR_012971

Alternate IDs: OMICS_00935

Record Creation Time: 20220129T080313+0000

Record Last Update: 20260905T132730+0000

Ratings and Alerts

No rating or validation information has been found for GBrowse syn.

No alerts have been found for GBrowse syn.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 9 mentions in open access literature.

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

Wang M, et al. (2017) Origin and evolution of MIR1444 genes in Salicaceae. Scientific reports, 7, 39740.

Bu L, et al. (2015) Early evolutionary history and genomic features of gene duplicates in the human genome. BMC genomics, 16(1), 621.

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

Chan ET, et al. (2012) Considerations for creating and annotating the budding yeast Genome Map at SGD: a progress report. Database : the journal of biological databases and curation, 2012, bar057.

Visendi P, et al. (2011) TparvaDB: a database to support Theileria parva vaccine development. Database : the journal of biological databases and curation, 2011, bar015.

Arnaiz O, et al. (2011) ParameciumDB in 2011: new tools and new data for functional and comparative genomics of the model ciliate Paramecium tetraurelia. Nucleic acids research, 39(Database issue), D632.

Harris TW, et al. (2010) WormBase: a comprehensive resource for nematode research. Nucleic acids research, 38(Database issue), D463.

Sen TZ, et al. (2009) MaizeGDB becomes 'sequence-centric'. Database : the journal of biological databases and curation, 2009, bap020.

Winsor GL, et al. (2009) Pseudomonas Genome Database: facilitating user-friendly, comprehensive comparisons of microbial genomes. Nucleic acids research, 37(Database issue), D483.