

Resource Summary Report

Generated by [NIF](#) on Aug 19, 2026

SynBrowse

RRID:SCR_011807

Type: Tool

Proper Citation

SynBrowse (RRID:SCR_011807)

Resource Information

URL: <http://www.synbrowse.org/>

Proper Citation: SynBrowse (RRID:SCR_011807)

Description: A generic sequence comparison tool for visualizing genome alignments both within and between species.

Abbreviations: SynBrowse

Synonyms: Synteny Browser

Resource Type: software resource

Resource Name: SynBrowse

Resource ID: SCR_011807

Alternate IDs: OMICS_00946

Record Creation Time: 20220129T080306+0000

Record Last Update: 20260815T182434+0000

Ratings and Alerts

No rating or validation information has been found for SynBrowse.

No alerts have been found for SynBrowse.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Van K, et al. (2008) Sequence level analysis of recently duplicated regions in soybean [Glycine max (L.) Merr.] genome. DNA research : an international journal for rapid publication of reports on genes and genomes, 15(2), 93.

Shin JH, et al. (2008) The lipoxygenase gene family: a genomic fossil of shared polyploidy between Glycine max and Medicago truncatula. BMC plant biology, 8, 133.