

# Resource Summary Report

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

## SynBrowse

RRID:SCR\_011807

Type: Tool

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### Proper Citation

SynBrowse (RRID:SCR\_011807)

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### 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

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### Ratings and Alerts

No rating or validation information has been found for SynBrowse.

No alerts have been found for SynBrowse.

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### 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 [dkNET](#) .

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.

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.