

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 19, 2026

## GSV

RRID:SCR\_011803

Type: Tool

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### Proper Citation

GSV (RRID:SCR\_011803)

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### Resource Information

**URL:** <http://cas-bioinfo.cas.unt.edu/gsv/homepage.php>

**Proper Citation:** GSV (RRID:SCR\_011803)

**Description:** Software that allows users to upload files which contain synteny regions between two or more genomes and interactively visualize the synteny between them.

**Abbreviations:** GSV

**Synonyms:** Genome Synteny Viewer

**Resource Type:** software resource

**Resource Name:** GSV

**Resource ID:** SCR\_011803

**Alternate IDs:** OMICS\_00939

**Record Creation Time:** 20220129T080306+0000

**Record Last Update:** 20260912T195740+0000

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### Ratings and Alerts

No rating or validation information has been found for GSV.

No alerts have been found for GSV.

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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 [PRECISE-TBI](#) .

Ács-Szabó L, et al. (2018) Assembly of Schizosaccharomyces cryophilus chromosomes and their comparative genomic analyses revealed principles of genome evolution of the haploid fission yeasts. Scientific reports, 8(1), 14629.

Zuo LH, et al. (2017) The first complete chloroplast genome sequences of Ulmus species by de novo sequencing: Genome comparative and taxonomic position analysis. PloS one, 12(2), e0171264.