

Resource Summary Report

Generated by [SPARC SAWG](#) on Sep 17, 2026

GSV

RRID:SCR_011803

Type: Tool

Proper Citation

GSV (RRID:SCR_011803)

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

Ratings and Alerts

No rating or validation information has been found for GSV.

No alerts have been found for GSV.

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 [SPARC SAWG](#) .

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.