

Resource Summary Report

Generated by SPARC SAWG on Aug 9, 2026

GenoViewer

RRID:SCR_001203

Type: Tool

Proper Citation

GenoViewer (RRID:SCR_001203)

Resource Information

URL: <http://www.genoviewer.com/>

Proper Citation: GenoViewer (RRID:SCR_001203)

Description: THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 23,2022. Open source viewer / browser software for the SAM / BAM format commonly used in the assembly tasks of Next Generation Sequencing data.

Abbreviations: GenoViewer

Resource Type: software resource

Defining Citation: [PMID:22359445](https://pubmed.ncbi.nlm.nih.gov/22359445/)

Keywords: next-generation sequencing, sequence, mutation, windows, linux, mac os x, genome, browser, sam, bam, fasta, gff, read error, snp, mnp, insertion, deletion

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: GenoViewer

Resource ID: SCR_001203

Alternate IDs: OMICS_02146

Alternate URLs: <https://github.com/astrid/GenoViewer>

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260808T185726+0000

Ratings and Alerts

No rating or validation information has been found for GenoViewer.

No alerts have been found for GenoViewer.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.