

Resource Summary Report

Generated by SPARC SAWG on Jul 28, 2026

GenomeSmasher

RRID:SCR_002406

Type: Tool

Proper Citation

GenomeSmasher (RRID:SCR_002406)

Resource Information

URL: <https://code.google.com/p/genome-smasher/>

Proper Citation: GenomeSmasher (RRID:SCR_002406)

Description: Software repository for tools used to create diploid FASTA files with containing snps, indels, duplications, deletions, and translocations. They can be used to create artificial genomes for next-gen sequencing simulations.

Resource Type: software application, software resource

Keywords: fasta file creation, file creation, artificial genome creator, diploid fasta files, sequencing simulations

Availability: Open source

Resource Name: GenomeSmasher

Resource ID: SCR_002406

Alternate IDs: OMICS_00251

License: GNU GPL v3

Record Creation Time: 20220129T080213+0000

Record Last Update: 20260727T040314+0000

Ratings and Alerts

No rating or validation information has been found for GenomeSmasher.

No alerts have been found for GenomeSmasher.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.