

# Resource Summary Report

Generated by T1D on Aug 2, 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 resource, software application

**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:** 20260801T191039+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.