

# Resource Summary Report

Generated by [dkNET](#) on Aug 16, 2026

## GLFSINGLE/GLFTRIO/GLFMULTIPLES

RRID:SCR\_013128

Type: Tool

---

### Proper Citation

GLFSINGLE/GLFTRIO/GLFMULTIPLES (RRID:SCR\_013128)

---

### Resource Information

**URL:** <http://genome.sph.umich.edu/wiki/GlfSingle>

**Proper Citation:** GLFSINGLE/GLFTRIO/GLFMULTIPLES (RRID:SCR\_013128)

**Description:** Software application that is a GLF-based variant caller for next-generation sequencing data. It takes one/three/multiple GLF format genotype likelihood files as input and generates a VCF-format set of variant calls as output. (entry from Genetic Analysis Software)

**Resource Type:** software application, software resource

**Keywords:** gene, genetic, genomic

**Resource Name:** GLFSINGLE/GLFTRIO/GLFMULTIPLES

**Resource ID:** SCR\_013128

**Alternate IDs:** nlx\_154358

**Record Creation Time:** 20220129T080314+0000

**Record Last Update:** 20260815T182949+0000

---

### Ratings and Alerts

No rating or validation information has been found for GLFSINGLE/GLFTRIO/GLFMULTIPLES.

No alerts have been found for GLFSINGLE/GLFTRIO/GLFMULTIPLES.

---

### Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [dkNET](#) .

Lo Y, et al. (2015) Comparing variant calling algorithms for target-exon sequencing in a large sample. BMC bioinformatics, 16, 75.