Resource Summary Report

Generated by NIF on Apr 19, 2025

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

Funding:

Resource Name: GLFSINGLE/GLFTRIO/GLFMULTIPLES

Resource ID: SCR_013128

Alternate IDs: nlx_154358

Record Creation Time: 20220129T080314+0000

Record Last Update: 20250419T055344+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 NIF.

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