Resource Summary Report

Generated by dkNET on Apr 30, 2025

glfMultiples

RRID:SCR_010763

Type: Tool

Proper Citation

glfMultiples (RRID:SCR_010763)

Resource Information

URL: http://genome.sph.umich.edu/wiki/GlfMultiples

Proper Citation: glfMultiples (RRID:SCR_010763)

Description: A GLF-based variant caller for next-generation sequencing data.

Abbreviations: glfMultiples

Resource Type: software resource

Funding:

Resource Name: glfMultiples

Resource ID: SCR_010763

Alternate IDs: OMICS_00061

Record Creation Time: 20220129T080300+0000

Record Last Update: 20250420T014508+0000

Ratings and Alerts

No rating or validation information has been found for glfMultiples.

No alerts have been found for glfMultiples.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 3 mentions in open access literature.

Listed below are recent publications. The full list is available at dkNET.

Onda Y, et al. (2018) Multiplex PCR Targeted Amplicon Sequencing (MTA-Seq): Simple, Flexible, and Versatile SNP Genotyping by Highly Multiplexed PCR Amplicon Sequencing. Frontiers in plant science, 9, 201.

Jun G, et al. (2015) An efficient and scalable analysis framework for variant extraction and refinement from population-scale DNA sequence data. Genome research, 25(6), 918.

Emond MJ, et al. (2015) Exome Sequencing of Phenotypic Extremes Identifies CAV2 and TMC6 as Interacting Modifiers of Chronic Pseudomonas aeruginosa Infection in Cystic Fibrosis. PLoS genetics, 11(6), e1005273.