

Resource Summary Report

Generated by [dkNET](#) on Apr 18, 2025

UMAKE

RRID:SCR_009093

Type: Tool

Proper Citation

UMAKE (RRID:SCR_009093)

Resource Information

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

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Description: Software pipeline to detect SNPs and call their genotypes from a list of BAM files. UMAKE pipeline has been successfully applied in detecting SNPs from many large-scale next-generation sequencing studies. (entry from Genetic Analysis Software)

Abbreviations: UMAKE

Resource Type: software resource, software application

Keywords: gene, genetic, genomic

Funding:

Resource Name: UMAKE

Resource ID: SCR_009093

Alternate IDs: nlx_154110

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250416T063536+0000

Ratings and Alerts

No rating or validation information has been found for UMAKE.

No alerts have been found for UMAKE.

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](#).

Hu Y, et al. (2013) Accurate local-ancestry inference in exome-sequenced admixed individuals via off-target sequence reads. *American journal of human genetics*, 93(5), 891.

Orrù V, et al. (2013) Genetic variants regulating immune cell levels in health and disease. *Cell*, 155(1), 242.

Zhan X, et al. (2013) Identification of a rare coding variant in complement 3 associated with age-related macular degeneration. *Nature genetics*, 45(11), 1375.