Resource Summary Report

Generated by NIF on May 8, 2025

SNVMix

RRID:SCR_013050
Type: Tool

Proper Citation

SNVMix (RRID:SCR_013050)

Resource Information

URL: http://compbio.bccrc.ca/software/snvmix/

Proper Citation: SNVMix (RRID:SCR_013050)

Description: Software designed to detect single nucleotide variants from next generation sequencing data.

Abbreviations: SNVMix

Resource Type: software resource

Funding:

Resource Name: SNVMix

Resource ID: SCR_013050

Alternate IDs: OMICS_00077

Record Creation Time: 20220129T080314+0000

Record Last Update: 20250420T014631+0000

Ratings and Alerts

No rating or validation information has been found for SNVMix.

No alerts have been found for SNVMix.

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.

Ha G, et al. (2012) Integrative analysis of genome-wide loss of heterozygosity and monoallelic expression at nucleotide resolution reveals disrupted pathways in triple-negative breast cancer. Genome research, 22(10), 1995.