Resource Summary Report

Generated by NIF on Aug 29, 2026

AgileVariantMapper

RRID:SCR_010770
Type: Tool

Proper Citation

AgileVariantMapper (RRID:SCR_010770)

Resource Information

URL: http://www.insilicase.co.uk/agile/AgileVariantMapper.aspx

Proper Citation: AgileVariantMapper (RRID:SCR_010770)

Description: Software that visualises sequence variant data from whole exome data, so that it is possible to identify autozygous regions in consanguineous individuals.

Abbreviations: AgileVariantMapper

Synonyms: AgileVariantMapper - Autozygosity mapping using Next generation sequence data

Resource Type: software resource

Defining Citation: PMID:23090942

Resource Name: AgileVariantMapper

Resource ID: SCR_010770

Alternate IDs: OMICS_00122

Record Creation Time: 20220129T080300+0000

Record Last Update: 20260815T182408+0000

Ratings and Alerts

No rating or validation information has been found for AgileVariantMapper.

No alerts have been found for AgileVariantMapper.

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 .

Boone PM, et al. (2016) Hutterite-type cataract maps to chromosome 6p21.32-p21.31, cosegregates with a homozygous mutation in LEMD2, and is associated with sudden cardiac death. Molecular genetics & genomic medicine, 4(1), 77.