

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

Multipoint Identical-by-descent Method

RRID:SCR_004676

Type: Tool

Proper Citation

Multipoint Identical-by-descent Method (RRID:SCR_004676)

Resource Information

URL: <http://ftp://morgan.med.utah.edu/pub/Mim>

Proper Citation: Multipoint Identical-by-descent Method (RRID:SCR_004676)

Description: Software application using multipoint IBD method for partitioning genetic variance of quantitative traits to specific chromosome regions using data on nuclear families. (entry from Genetic Analysis Software)

Abbreviations: MIM

Resource Type: software application, software resource

Defining Citation: [PMID:9433587](#)

Keywords: gene, genetic, genomic, c, unix

Resource Name: Multipoint Identical-by-descent Method

Resource ID: SCR_004676

Alternate IDs: nlx_154482

Record Creation Time: 20220129T080225+0000

Record Last Update: 20260805T044111+0000

Ratings and Alerts

No rating or validation information has been found for Multipoint Identical-by-descent Method.

No alerts have been found for Multipoint Identical-by-descent Method.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.