

Resource Summary Report

Generated by [dkNET](#) on Sep 2, 2026

MUTAGENESYS

RRID:SCR_009309

Type: Tool

Proper Citation

MUTAGENESYS (RRID:SCR_009309)

Resource Information

URL: <http://magnet.c2b2.columbia.edu/mutagenesys/>

Proper Citation: MUTAGENESYS (RRID:SCR_009309)

Description: Software application that uses genome-wide genotype data to estimate disease susceptibility. Our system integrates three data sources: HapMap, whole-genome marker correlation data, and OMIM database. It accepts SNP data of individuals as query input and delivers disease susceptibility hypotheses even if the original set of typed SNPs is incomplete. (entry from Genetic Analysis Software)

Abbreviations: MUTAGENESYS

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Resource Name: MUTAGENESYS

Resource ID: SCR_009309

Alternate IDs: nlx_154499

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260829T183107+0000

Ratings and Alerts

No rating or validation information has been found for MUTAGENESYS.

No alerts have been found for MUTAGENESYS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.