

Resource Summary Report

Generated by SPARC SAWG on Aug 10, 2026

MLBGH

RRID:SCR_009297

Type: Tool

Proper Citation

MLBGH (RRID:SCR_009297)

Resource Information

URL: <http://ftp://ftp.biomath.jussieu.fr/pub/mlbgh> (not available)

Proper Citation: MLBGH (RRID:SCR_009297)

Description: Software application that is an extension of the GENEHUNTER program to perform sib-pair and sib-ship linkage analysis using the Maximum Likelihood Binomial (MLB) method. (entry from Genetic Analysis Software)

Abbreviations: MLBGH

Synonyms: Maximum-Likelihood-Binomial analysis of affected sib-pair and sibship data extended from the GENEHUNTER program

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, unix

Resource Name: MLBGH

Resource ID: SCR_009297

Alternate IDs: nlx_154488

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260810T040509+0000

Ratings and Alerts

No rating or validation information has been found for MLBGH.

No alerts have been found for MLBGH.

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 [SPARC SAWG](#) .

Alale TY, et al. (2024) Genomic signatures of hybridization between *Ixodes ricinus* and *Ixodes persulcatus* in natural populations. *Ecology and evolution*, 14(5), e11415.