

# Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 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:** 20260905T133249+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](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [PRECISE-TBI](#) .

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