Resource Summary Report

Generated by <u>NIF</u> on Apr 23, 2025

MLBGH

RRID:SCR_009297 Type: Tool

Proper Citation

MLBGH (RRID:SCR_009297)

Resource Information

URL: http://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 resource, software application

Keywords: gene, genetic, genomic, c, unix

Funding:

Resource Name: MLBGH

Resource ID: SCR_009297

Alternate IDs: nlx_154488

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250421T053725+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: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at <u>NIF</u>.

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