Resource Summary Report

Generated by Kravitz on Sep 10, 2026

LIPED

RRID:SCR_009255
Type: Tool

Proper Citation

LIPED (RRID:SCR_009255)

Resource Information

URL: https://github.com/gaow/genetic-analysis-software/blob/master/pages/LIPED.md

Proper Citation: LIPED (RRID:SCR_009255)

Description: Software application (entry from Genetic Analysis Software)

Abbreviations: LIPED

Synonyms: Likelihoods in Pedigrees

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, fortran, ms-windows

Resource Name: LIPED

Resource ID: SCR_009255

Alternate IDs: nlx_154432

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260905T133248+0000

Ratings and Alerts

No rating or validation information has been found for LIPED.

No alerts have been found for LIPED.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5 mentions in open access literature.

Listed below are recent publications. The full list is available at Kravitz .

Eiberg H, et al. (2019) A splice-site variant in the lncRNA gene RP1-140A9.1 cosegregates in the large Volkmann cataract family. Molecular vision, 25, 1.

Minocherhomji S, et al. (2014) Epigenetic remodelling and dysregulation of DLGAP4 is linked with early-onset cerebellar ataxia. Human molecular genetics, 23(23), 6163.

Anjum I, et al. (2010) A mutation in the FOXE3 gene causes congenital primary aphakia in an autosomal recessive consanguineous Pakistani family. Molecular vision, 16, 549.

Govil M, et al. (2008) MLIP: using multiple processors to compute the posterior probability of linkage. BMC bioinformatics, 9 Suppl 6(Suppl 6), S2.

Bartlett CW, et al. (2005) Two novel quantitative trait linkage analysis statistics based on the posterior probability of linkage: application to the COGA families. BMC genetics, 6 Suppl 1(Suppl 1), S121.