

# Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

## LIPED

RRID:SCR\_009255

Type: Tool

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### Proper Citation

LIPED (RRID:SCR\_009255)

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### 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:** 20260805T044220+0000

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### Ratings and Alerts

No rating or validation information has been found for LIPED.

No alerts have been found for LIPED.

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### 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 [RRID](#) .

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.