

# Resource Summary Report

Generated by [kravitz2](#) on Jul 31, 2026

## SEQUENCE LD/SEQUENCE LDHOT

RRID:SCR\_009379

Type: Tool

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### Proper Citation

SEQUENCE LD/SEQUENCE LDHOT (RRID:SCR\_009379)

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### Resource Information

**URL:** <http://www.maths.lancs.ac.uk/~fearnhea/Hotspot/>

**Proper Citation:** SEQUENCE LD/SEQUENCE LDHOT (RRID:SCR\_009379)

**Description:** Software program that analyzes sequence data. It obtains an approximation to the likelihood of a summary of the data (as such it can be thought of as a marginal likelihood approach). It does not use all the information in the data, but computationally it can be substantially more efficient than the full-likelihood methods (and hence able to analyze larger data sets). (entry from Genetic Analysis Software)

**Resource Type:** software resource, software application

**Keywords:** gene, genetic, genomic

**Resource Name:** SEQUENCE LD/SEQUENCE LDHOT

**Resource ID:** SCR\_009379

**Alternate IDs:** nlx\_154614

**Old URLs:** <http://www.maths.lancs.ac.uk/~fearnhea/Software.html>

**Record Creation Time:** 20220129T080252+0000

**Record Last Update:** 20260731T042809+0000

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### Ratings and Alerts

No rating or validation information has been found for SEQUENCE LD/SEQUENCE LDHOT.

No alerts have been found for SEQUENCE LD/SEQUENCE LDHOT.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.