

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

SEQUENCE LD/SEQUENCE LDHOT

RRID:SCR_009379

Type: Tool

Proper Citation

SEQUENCE LD/SEQUENCE LDHOT (RRID:SCR_009379)

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 application, software resource

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: 20260905T133251+0000

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.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.