

# Resource Summary Report

Generated by T1D on Aug 2, 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 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:** 20260801T191058+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.