

# Resource Summary Report

Generated by RRID on Aug 5, 2026

## L-POP

RRID:SCR\_009262

Type: Tool

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### Proper Citation

L-POP (RRID:SCR\_009262)

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### Resource Information

**URL:** <http://statgen.iop.kcl.ac.uk/lpop/>

**Proper Citation:** L-POP (RRID:SCR\_009262)

**Description:** Software application that detects population stratification in samples of unrelated individuals for whom a number of unlinked genotypes have been measured. (entry from Genetic Analysis Software), THIS RESOURCE IS NO LONGER IN SERVICE. Documented on September 16,2025.

**Abbreviations:** L-POP

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

**Keywords:** gene, genetic, genomic, ms-dos

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** L-POP

**Resource ID:** SCR\_009262

**Alternate IDs:** nlx\_154441

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

**Record Last Update:** 20260805T044220+0000

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### Ratings and Alerts

No rating or validation information has been found for L-POP.

No alerts have been found for L-POP.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](#) .

Koenen KC, et al. (2009) Protocol for investigating genetic determinants of posttraumatic stress disorder in women from the Nurses' Health Study II. BMC psychiatry, 9, 29.

Wu B, et al. (2006) PSMIX: an R package for population structure inference via maximum likelihood method. BMC bioinformatics, 7, 317.