

Resource Summary Report

Generated by [nidm-terms](#) on Sep 21, 2026

HAPLOREC

RRID:SCR_009055

Type: Tool

Proper Citation

HAPLOREC (RRID:SCR_009055)

Resource Information

URL: <http://www.cs.helsinki.fi/group/genetics/haplotyping.html>

Proper Citation: HAPLOREC (RRID:SCR_009055)

Description: Software application for population-based haplotyping (entry from Genetic Analysis Software)

Abbreviations: HAPLOREC

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Resource Name: HAPLOREC

Resource ID: SCR_009055

Alternate IDs: nlx_154036

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260919T195757+0000

Ratings and Alerts

No rating or validation information has been found for HAPLOREC.

No alerts have been found for HAPLOREC.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Zhang LJ, et al. (2017) Phylogeographic patterns of *Lygus pratensis* (Hemiptera: Miridae): Evidence for weak genetic structure and recent expansion in northwest China. PloS one, 12(4), e0174712.

Eronen L, et al. (2006) HaploRec: efficient and accurate large-scale reconstruction of haplotypes. BMC bioinformatics, 7, 542.

Salem RM, et al. (2005) A comprehensive literature review of haplotyping software and methods for use with unrelated individuals. Human genomics, 2(1), 39.

Weale ME, et al. (2004) A survey of current software for haplotype phase inference. Human genomics, 1(2), 141.