

Resource Summary Report

Generated by [nidm-terms](#) on Sep 23, 2026

HAPLOCLUSTERS

RRID:SCR_007439

Type: Tool

Proper Citation

HAPLOCLUSTERS (RRID:SCR_007439)

Resource Information

URL: <http://bioinf.wehi.edu.au/folders/melanie/haploclusters.html>

Proper Citation: HAPLOCLUSTERS (RRID:SCR_007439)

Description: Software program designed to detect excess haplotypes sharing in datasets consisting of case and control haplotypes. Excess haplotype sharing can be seen around disease loci in case samples since LD persists longer here than in the controls where LD is persisting only according to the relatedness of the individuals in the population, i.e. the age of the population. (entry from Genetic Analysis Software)

Abbreviations: HAPLOCLUSTERS

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, bio.tools

Related Condition: Aging

Resource Name: HAPLOCLUSTERS

Resource ID: SCR_007439

Alternate IDs: nlx_154014, biotools:haploclusters

Alternate URLs: <https://bio.tools/haploclusters>

Record Creation Time: 20220129T080241+0000

Record Last Update: 20260919T195754+0000

Ratings and Alerts

No rating or validation information has been found for HAPLOCLUSTERS.

No alerts have been found for HAPLOCLUSTERS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.