

Resource Summary Report

Generated by [kravitz2](#) on Sep 17, 2026

TRIMHAP

RRID:SCR_013512

Type: Tool

Proper Citation

TRIMHAP (RRID:SCR_013512)

Resource Information

URL: <http://vipbg.vcu.edu/vipbg/trimhap/>

Proper Citation: TRIMHAP (RRID:SCR_013512)

Description: Software application for linkage disequilibrium mapping based on ancestral founder haplotypes. Method uses haplotype data from general pedigrees. (entry from Genetic Analysis Software)

Abbreviations: TRIMHAP

Synonyms: TRIMmed-HAPlotype (previously named HAL: Haplotype ALgorithm)

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, fortran, unix

Resource Name: TRIMHAP

Resource ID: SCR_013512

Alternate IDs: nlx_154683

Record Creation Time: 20220129T080316+0000

Record Last Update: 20260912T200249+0000

Ratings and Alerts

No rating or validation information has been found for TRIMHAP.

No alerts have been found for TRIMHAP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.