

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

HAPLOTYPE ESTIMATION

RRID:SCR_009231

Type: Tool

Proper Citation

HAPLOTYPE ESTIMATION (RRID:SCR_009231)

Resource Information

URL: <http://www.bioinf.mdc-berlin.de/projects/hap/>

Proper Citation: HAPLOTYPE ESTIMATION (RRID:SCR_009231)

Description: Haplotype estimation service available in two variants: one for genetic data from unrelated probands (e.g., case/control studies) and one for core family data (entry from Genetic Analysis Software)

Abbreviations: HAPLOTYPE ESTIMATION

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Resource Name: HAPLOTYPE ESTIMATION

Resource ID: SCR_009231

Alternate IDs: nlx_154388

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260727T040455+0000

Ratings and Alerts

No rating or validation information has been found for HAPLOTYPE ESTIMATION.

No alerts have been found for HAPLOTYPE ESTIMATION.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.