

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

HAP 1

RRID:SCR_000837

Type: Tool

Proper Citation

HAP 1 (RRID:SCR_000837)

Resource Information

URL: <http://research.calit2.net/hap/>

Proper Citation: HAP 1 (RRID:SCR_000837)

Description: Software application (entry from Genetic Analysis Software)

Abbreviations: HAP 1

Synonyms: haplotype resolution using imperfect phylogeny

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: HAP 1

Resource ID: SCR_000837

Alternate IDs: nlx_154371

Record Creation Time: 20220129T080203+0000

Record Last Update: 20260727T040251+0000

Ratings and Alerts

No rating or validation information has been found for HAP 1.

No alerts have been found for HAP 1.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Kalis M, et al. (2007) Variants in the FFAR1 gene are associated with beta cell function. PloS one, 2(11), e1090.

Marchini J, et al. (2006) A comparison of phasing algorithms for trios and unrelated individuals. American journal of human genetics, 78(3), 437.

Scheet P, et al. (2006) A fast and flexible statistical model for large-scale population genotype data: applications to inferring missing genotypes and haplotypic phase. American journal of human genetics, 78(4), 629.