

# Resource Summary Report

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

## HAPLO 1

RRID:SCR\_009058

Type: Tool

### Proper Citation

HAPLO 1 (RRID:SCR\_009058)

### Resource Information

**URL:** [http://watson.hgen.pitt.edu/register/soft\\_doc.html](http://watson.hgen.pitt.edu/register/soft_doc.html),

**Proper Citation:** HAPLO 1 (RRID:SCR\_009058)

**Description:** THIS RESOURCE IS NO LONGER IN SERVICE. Documented on May 16,2023. Software application that assigns each person the haplotype with the highest conditional probability using Elston-Stewart algorithm (entry from Genetic Analysis Software)

**Abbreviations:** HAPLO 1

**Synonyms:** Haplotyping with computation of conditional probabilities

**Resource Type:** software application, software resource

**Keywords:** gene, genetic, genomic, fortran 77, (ansi standard), unix, sunos, solaris, hp

**Availability:** THIS RESOURCE IS NO LONGER IN SERVICE

**Resource Name:** HAPLO 1

**Resource ID:** SCR\_009058

**Alternate IDs:** nlx\_154039

**Record Creation Time:** 20220129T080250+0000

**Record Last Update:** 20260727T040449+0000

### Ratings and Alerts

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

No alerts have been found for HAPLO 1.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.