

Resource Summary Report

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

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.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.