

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

HAPLO 2

RRID:SCR_009059

Type: Tool

Proper Citation

HAPLO 2 (RRID:SCR_009059)

Resource Information

URL: <http://krunch.med.yale.edu/haplo/>

Proper Citation: HAPLO 2 (RRID:SCR_009059)

Description: Software application that estimates frequencies of multi-site haplotypes using the EM algorithm (entry from Genetic Analysis Software)

Abbreviations: HAPLO 2

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, fortran, dec vax

Resource Name: HAPLO 2

Resource ID: SCR_009059

Alternate IDs: nlx_154040

Record Creation Time: 20220129T080250+0000

Record Last Update: 20260805T044217+0000

Ratings and Alerts

No rating or validation information has been found for HAPLO 2.

No alerts have been found for HAPLO 2.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Weale ME, et al. (2004) A survey of current software for haplotype phase inference. Human genomics, 1(2), 141.