Resource Summary Report

Generated by <u>NIF</u> on Apr 21, 2025

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 resource, software application

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

Funding:

Resource Name: HAPLO 2

Resource ID: SCR_009059

Alternate IDs: nlx_154040

Record Creation Time: 20220129T080250+0000

Record Last Update: 20250421T053713+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: <u>SciCrunch Registry</u>

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

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