## **Resource Summary Report**

Generated by FDI Lab - SciCrunch.org on Apr 18, 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:** 20250416T063535+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 FDI Lab - SciCrunch.org.

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