

Resource Summary Report

Generated by SPARC SAWG on Sep 15, 2026

EHP

RRID:SCR_009170

Type: Tool

Proper Citation

EHP (RRID:SCR_009170)

Resource Information

URL: <http://gaow.github.io/genetic-analysis-software/e-1.html#ehp>

Proper Citation: EHP (RRID:SCR_009170)

Description: Software application that provides variance estimates for haplotype frequency estimates, it allows several kinds of missing information in the genotype data, it also allows for combined genotype data of different pool sizes. This program can be used for testing haplotype-disease associations in case control studies by calculating the likelihood ratio test: $2 \log(\text{likelihood for cases}) + 2 \log(\text{likelihood for controls}) - 2 \log(\text{likelihood for case+controls})$. (entry from Genetic Analysis Software)

Synonyms: Estimation of Haplotype frequencies from Pooled DNA samples EH, R/EHP, EHP.R

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, r/s-plus

Resource Name: EHP

Resource ID: SCR_009170

Alternate IDs: SCR_009101, nlx_154194, nlx_154293

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260912T200241+0000

Ratings and Alerts

No rating or validation information has been found for EHP.

No alerts have been found for EHP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.