

# Resource Summary Report

Generated by [dkNET](#) on Apr 23, 2025

## EHP

RRID:SCR\_009170

Type: Tool

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### Proper Citation

EHP (RRID:SCR\_009170)

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### Resource Information

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

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**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 resource, software application

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

**Funding:**

**Resource Name:** EHP

**Resource ID:** SCR\_009170

**Alternate IDs:** SCR\_009101, nlx\_154293, nlx\_154194

**Record Creation Time:** 20220129T080251+0000

**Record Last Update:** 20250422T055523+0000

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## Ratings and Alerts

No rating or validation information has been found for EHP.

No alerts have been found for EHP.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.