

Resource Summary Report

Generated by [NIF](#) on Sep 3, 2026

EHAP

RRID:SCR_009169

Type: Tool

Proper Citation

EHAP (RRID:SCR_009169)

Resource Information

URL: http://wpicr.wpic.pitt.edu/WPICCompGen/ehap_v1.htm

Proper Citation: EHAP (RRID:SCR_009169)

Description: Software application detecting association between haplotypes and phenotypes (entry from Genetic Analysis Software)

Abbreviations: EHAP

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Resource Name: EHAP

Resource ID: SCR_009169

Alternate IDs: nlx_154292

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260829T183107+0000

Ratings and Alerts

No rating or validation information has been found for EHAP.

No alerts have been found for EHAP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Wu Y, et al. (2024) Assessing the effectiveness of the expanded hepatitis A vaccination program in China: an interrupted time series design. *BMJ global health*, 9(2).

Ang CS, et al. (2022) Getting more out of FLAG-Tag co-immunoprecipitation mass spectrometry experiments using FAIMS. *Journal of proteomics*, 254, 104473.

Hewitt G, et al. (2021) Defective ALC1 nucleosome remodeling confers PARPi sensitization and synthetic lethality with HRD. *Molecular cell*, 81(4), 767.

Halder I, et al. (2010) Polymorphisms in the CRP gene moderate an association between depressive symptoms and circulating levels of C-reactive protein. *Brain, behavior, and immunity*, 24(1), 160.

Yu CE, et al. (2007) Comprehensive analysis of APOE and selected proximate markers for late-onset Alzheimer's disease: patterns of linkage disequilibrium and disease/marker association. *Genomics*, 89(6), 655.