Resource Summary Report

Generated by NIF on Apr 23, 2025

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 resource, software application

Keywords: gene, genetic, genomic

Funding:

Resource Name: EHAP

Resource ID: SCR_009169

Alternate IDs: nlx_154292

Record Creation Time: 20220129T080251+0000

Record Last Update: 20250421T053722+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.