

Resource Summary Report

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HAPLOT

RRID:SCR_009228

Type: Tool

Proper Citation

HAPLOT (RRID:SCR_009228)

Resource Information

URL: <http://info.med.yale.edu/genetics/kkidd/programs.html>

Proper Citation: HAPLOT (RRID:SCR_009228)

Description: A simple software program for graphical presentation of haplotype block structures, tagSNP selection and SNP variation. (entry from Genetic Analysis Software)

Abbreviations: HAPLOT

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, fortran, vms

Resource Name: HAPLOT

Resource ID: SCR_009228

Alternate IDs: nlx_154385

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260919T195801+0000

Ratings and Alerts

No rating or validation information has been found for HAPLOT.

No alerts have been found for HAPLOT.

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 [nidm-terms](#) .

Zhang L, et al. (2024) Integrating GWAS, RNA-Seq and functional analysis revealed that BnaA02.SE mediates silique elongation by affecting cell proliferation and expansion in *Brassica napus*. *Plant biotechnology journal*, 22(10), 2907.

Gu S, et al. (2018) Recent Selection on a Class I ADH Locus Distinguishes Southwest Asian Populations Including Ashkenazi Jews. *Genes*, 9(9).

Zanetti D, et al. (2015) Potential Signals of Natural Selection in the Top Risk Loci for Coronary Artery Disease: 9p21 and 10q11. *PloS one*, 10(8), e0134840.

Zabetian CP, et al. (2003) The structure of linkage disequilibrium at the DBH locus strongly influences the magnitude of association between diallelic markers and plasma dopamine beta-hydroxylase activity. *American journal of human genetics*, 72(6), 1389.

Zabetian CP, et al. (2001) A quantitative-trait analysis of human plasma-dopamine beta-hydroxylase activity: evidence for a major functional polymorphism at the DBH locus. *American journal of human genetics*, 68(2), 515.