

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

HAPLOBLOCK

RRID:SCR_009223

Type: Tool

Proper Citation

HAPLOBLOCK (RRID:SCR_009223)

Resource Information

URL: <http://bioinfo.cs.technion.ac.il/haploblock/>

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Description: Software package which provides an integrated approach to haplotype block identification, haplotype resolution and linkage disequilibrium mapping, suitable for high-density phased or unphased SNP data. (entry from Genetic Analysis Software)

Abbreviations: HAPLOBLOCK

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, unix, solaris, linux, macos x

Resource Name: HAPLOBLOCK

Resource ID: SCR_009223

Alternate IDs: nlx_154379

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260905T133247+0000

Ratings and Alerts

No rating or validation information has been found for HAPLOBLOCK.

No alerts have been found for HAPLOBLOCK.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Kwakye A, et al. (2026) Rare jackpot individuals drive rapid adaptation in Threespine Stickleback. *Nature communications*, 17(1).

Temel ??G, et al. (2019) The association between the chromosome 9p21 CDKN2B-AS1 gene variants and the lipid metabolism: A pre-diagnostic biomarker for coronary artery disease. *Anatolian journal of cardiology*, 21(1), 31.

Jia HM, et al. (2019) The red bayberry genome and genetic basis of sex determination. *Plant biotechnology journal*, 17(2), 397.

Scheet P, et al. (2006) A fast and flexible statistical model for large-scale population genotype data: applications to inferring missing genotypes and haplotypic phase. *American journal of human genetics*, 78(4), 629.