

Resource Summary Report

Generated by [NIF](#) on Sep 4, 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: 20260904T000429+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 [NIF](#) .

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.