

Resource Summary Report

Generated by [PRECISE-TBI](#) on Aug 10, 2026

HAPBLOCK

RRID:SCR_009220

Type: Tool

Proper Citation

HAPBLOCK (RRID:SCR_009220)

Resource Information

URL: <http://www-hto.usc.edu/msms/HapBlock/>

Proper Citation: HAPBLOCK (RRID:SCR_009220)

Description: Software application (entry from Genetic Analysis Software)

Abbreviations: HAPBLOCK

Synonyms: a dynamical programming algorithm for haplotype block partitioning

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, c, ms-windows, linux, unix, solaris

Resource Name: HAPBLOCK

Resource ID: SCR_009220

Alternate IDs: nlx_154375

Record Creation Time: 20220129T080251+0000

Record Last Update: 20260810T040507+0000

Ratings and Alerts

No rating or validation information has been found for HAPBLOCK.

No alerts have been found for HAPBLOCK.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

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

Orfanou E, et al. (2026) Reconstruction of the lifeways of Central European Late Bronze Age communities using ancient DNA, isotope and osteoarchaeological analyses. *Nature communications*, 17(1).

Moon H, et al. (2026) Ancient genomes reveal an extensive kinship network and endogamy in a Three-Kingdoms period society in Korea. *Science advances*, 12(15), eady8614.

Gretzinger J, et al. (2025) Ancient DNA connects large-scale migration with the spread of Slavs. *Nature*, 646(8084), 384.

Ghalichi A, et al. (2024) The rise and transformation of Bronze Age pastoralists in the Caucasus. *Nature*, 635(8040), 917.

Gnecchi-Ruscone GA, et al. (2024) Network of large pedigrees reveals social practices of Avar communities. *Nature*, 629(8011), 376.

Penske S, et al. (2023) Early contact between late farming and pastoralist societies in southeastern Europe. *Nature*, 620(7973), 358.

Nascimento AVD, et al. (2018) Genome-wide association study using haplotype alleles for the evaluation of reproductive traits in Nelore cattle. *PloS one*, 13(8), e0201876.

Netzer C, et al. (2008) Replication study of the insulin receptor gene in migraine with aura. *Genomics*, 91(6), 503.