Resource Summary Report

Generated by RRID on Aug 5, 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: 20260805T044219+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 6 mentions in open access literature.

Listed below are recent publications. The full list is available at RRID .

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.