

Resource Summary Report

Generated by [nidm-terms](#) on Jul 29, 2026

MDBLOCKS

RRID:SCR_009280

Type: Tool

Proper Citation

MDBLOCKS (RRID:SCR_009280)

Resource Information

URL: http://ib.berkeley.edu/labs/slatkin/eriq/software/mdb_web/

Proper Citation: MDBLOCKS (RRID:SCR_009280)

Description: Software application using the minimum description length model to delineate haplotype blocks (entry from Genetic Analysis Software)

Abbreviations: MDBLOCKS

Synonyms: Minimum Description length method for haplotype BLOCKS

Resource Type: software application, software resource

Keywords: gene, genetic, genomic

Resource Name: MDBLOCKS

Resource ID: SCR_009280

Alternate IDs: nlx_154461

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260727T040457+0000

Ratings and Alerts

No rating or validation information has been found for MDBLOCKS.

No alerts have been found for MDBLOCKS.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Indap AR, et al. (2005) Analysis of concordance of different haplotype block partitioning algorithms. BMC bioinformatics, 6, 303.