Resource Summary Report

Generated by NIF on Apr 20, 2025

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 mimimum 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

Funding:

Resource Name: MDBLOCKS

Resource ID: SCR_009280

Alternate IDs: nlx_154461

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250420T015756+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 NIF.

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