

Resource Summary Report

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MULTIPOPOTAGSELECT

RRID:SCR_009307

Type: Tool

Proper Citation

MULTIPOPOTAGSELECT (RRID:SCR_009307)

Resource Information

URL: <http://droog.gs.washington.edu/multiPopTagSelect.html>

Proper Citation: MULTIPOPOTAGSELECT (RRID:SCR_009307)

Description: Software program that selects a near-minimal set of tagging single-nucleotide polymorphisms (tagSNPs) that account for all observed patterns of linkage disequilibrium (LD) in multiple populations. (entry from Genetic Analysis Software)

Abbreviations: MULTIPOPOTAGSELECT

Resource Type: software application, software resource

Keywords: gene, genetic, genomic, perl

Resource Name: MULTIPOPOTAGSELECT

Resource ID: SCR_009307

Alternate IDs: nlx_154497

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260912T200244+0000

Ratings and Alerts

No rating or validation information has been found for MULTIPOPOTAGSELECT.

No alerts have been found for MULTIPOPOTAGSELECT.

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 [kravitz2](#) .

Abbott D, et al. (2012) An international collaborative family-based whole genome quantitative trait linkage scan for myopic refractive error. Molecular vision, 18, 720.