Resource Summary Report

Generated by RRID on Apr 11, 2025

TAGSTER

RRID:SCR_009413
Type: Tool

Proper Citation

TAGSTER (RRID:SCR_009413)

Resource Information

URL: http://www.niehs.nih.gov/research/resources/software/epidemiology/tagster/

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Description: Software tool to select, evaluate and visualize LD tag SNPs for single or multiple populations. The input files can be a set of dumped genotype files from International HapMap Project (http://www.hapmap.org/) (Hapmap format) or Seattle SNPs (http://pga.gs.washington.edu/) (Prettybase format). The ouput is a set of LD tag SNPs for single or multiple populations. (entry from Genetic Analysis Software)

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, perl, r, linux, macos, ms-windows

Funding:

Resource Name: TAGSTER

Resource ID: SCR_009413

Alternate IDs: nlx_154673

Record Creation Time: 20220129T080252+0000

Record Last Update: 20250411T055326+0000

Ratings and Alerts

No rating or validation information has been found for TAGSTER.

No alerts have been found for TAGSTER.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Hebbar P, et al. (2023) Linkage analysis using whole exome sequencing data implicates SLC17A1, SLC17A3, TATDN2 and TMEM131L in type 1 diabetes in Kuwaiti families. Scientific reports, 13(1), 14978.

Velez Edwards DR, et al. (2017) Nonsteroidal Anti-inflammatory Drug Interaction with Prostacyclin Synthase Protects from Miscarriage. Scientific reports, 7(1), 9874.