

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

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/>

Proper Citation: TAGSTER (RRID:SCR_009413)

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 output is a set of LD tag SNPs for single or multiple populations. (entry from Genetic Analysis Software)

Resource Type: software application, software resource

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

Resource Name: TAGSTER

Resource ID: SCR_009413

Alternate IDs: nlx_154673

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260805T044222+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.