

Resource Summary Report

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PRESTO: Genetic Association Analysis Software

RRID:SCR_013285

Type: Tool

Proper Citation

PRESTO: Genetic Association Analysis Software (RRID:SCR_013285)

Resource Information

URL: <http://faculty.washington.edu/browning/presto/presto.html>

Proper Citation: PRESTO: Genetic Association Analysis Software (RRID:SCR_013285)

Description: Software application that performs permutation testing and computes empirical distributions of order statistics for one and two stage association studies with stratified or unstratified data.

Resource Type: software application, software resource

Defining Citation: [DOI:10.1093/bioinformatics/btu138](https://doi.org/10.1093/bioinformatics/btu138)

Keywords: gene, genetic, genomic, java, ms-windows, unix, solaris, linux, macos

Resource Name: PRESTO: Genetic Association Analysis Software

Resource ID: SCR_013285

Alternate IDs: nlx_154549

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260727T040551+0000

Ratings and Alerts

No rating or validation information has been found for PRESTO: Genetic Association Analysis Software.

No alerts have been found for PRESTO: Genetic Association Analysis Software.

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 [nidm-terms](#) .

Okoye GD, et al. (2025) Single-cell map of innate-like lymphocyte response to *Francisella tularensis* infection reveals interleukin-17-dependent protection by MAIT cells. *iScience*, 28(3), 111810.

Rubelt F, et al. (2016) Individual heritable differences result in unique cell lymphocyte receptor repertoires of na⁺ve and antigen-experienced cells. *Nature communications*, 7, 11112.