

# Resource Summary Report

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## RELATIVEFINDER

RRID:SCR\_009356

Type: Tool

### Proper Citation

RELATIVEFINDER (RRID:SCR\_009356)

### Resource Information

**URL:** <http://genome.sph.umich.edu/wiki/RelativeFinder>

**Proper Citation:** RELATIVEFINDER (RRID:SCR\_009356)

**Description:** Software program for checking relationships between pairs of individuals. There are many excellent programs that carry out similar tasks. Some of the unique features in relativeFinder are the batch mode options, that allow large jobs to be divided into many smaller jobs (suitable for deployment on a compute cluster environment), and the flexibility of the underlying Merlin engine, which allows relative finder to handle large pedigrees and consider a variety of alternate relationships -- including potential relationships specified by the user on the fly. (entry from Genetic Analysis Software)

**Abbreviations:** RELATIVEFINDER

**Resource Type:** software resource, software application

**Keywords:** gene, genetic, genomic

**Resource Name:** RELATIVEFINDER

**Resource ID:** SCR\_009356

**Alternate IDs:** nlx\_154571

**Record Creation Time:** 20220129T080252+0000

**Record Last Update:** 20260801T191100+0000

### Ratings and Alerts

No rating or validation information has been found for RELATIVEFINDER.

No alerts have been found for RELATIVEFINDER.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [ASWG](#) .

Stuart PE, et al. (2015) Genome-wide Association Analysis of Psoriatic Arthritis and Cutaneous Psoriasis Reveals Differences in Their Genetic Architecture. American journal of human genetics, 97(6), 816.

Lange LA, et al. (2014) Whole-exome sequencing identifies rare and low-frequency coding variants associated with LDL cholesterol. American journal of human genetics, 94(2), 233.