

Resource Summary Report

Generated by PRECISE-TBI on Aug 1, 2026

SIMULA

RRID:SCR_009390

Type: Tool

Proper Citation

SIMULA (RRID:SCR_009390)

Resource Information

URL: <http://www.chg.duke.edu/research/simla30.html>

Proper Citation: SIMULA (RRID:SCR_009390)

Description: THIS RESOURCE IS NO LONGER IN SERVICE, documented September 29, 2016. Simulation program that generates data sets of families for use in linkage and association studies. SIMLA_3.2 is a major upgrade to versions 2.3 and 3.0 that provides the ability to simulate two disease loci and two environmental covariates. Gene-gene and gene-environment interactions may also be simulated which jointly determine the disease risk of all pedigree members.

Abbreviations: SIMULA

Resource Type: software resource, software application

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

Availability: THIS RESOURCE IS NO LONGER IN SERVICE

Resource Name: SIMULA

Resource ID: SCR_009390

Alternate IDs: nlx_154629

Old URLs: <http://www.chg.duke.edu/research/simla30.html>

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260731T042817+0000

Ratings and Alerts

No rating or validation information has been found for SIMULA.

No alerts have been found for SIMULA.

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 [PRECISE-TBI](#) .

Skovlund E, et al. (1999) Repeated significance tests on accumulating survival data. Journal of clinical epidemiology, 52(11), 1083.