

Resource Summary Report

Generated by PRECISE-TBI on Aug 1, 2026

R/METASIM

RRID:SCR_009370

Type: Tool

Proper Citation

R/METASIM (RRID:SCR_009370)

Resource Information

URL: <https://cran.r-project.org/web/packages/rmetasim/index.html>

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Description: An R package that uses an individual-based approach to simulate distributions of genotypes that result from arbitrary within and among population demographies (including extinction/recolonization). These distributions can be used to test new or existing population-genetics summary statistics or develop null distributions under various demographies. (entry from Genetic Analysis Software)

Synonyms: METApopulation SIMulation

Resource Type: software resource, software application

Keywords: gene, genetic, genomic, r, c++, unix, ms-windows, macos

Resource Name: R/METASIM

Resource ID: SCR_009370

Alternate IDs: nlx_154592

Old URLs: <http://linum.cofc.edu/software.html>

Record Creation Time: 20220129T080252+0000

Record Last Update: 20260731T042808+0000

Ratings and Alerts

No rating or validation information has been found for R/METASIM.

No alerts have been found for R/METASIM.

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](#) .

Carvajal-Rodr  guez A, et al. (2010) Simulation of genes and genomes forward in time. Current genomics, 11(1), 58.