

# Resource Summary Report

Generated by [dkNET](#) on Jul 29, 2026

## R/METASIM

RRID:SCR\_009370

Type: Tool

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### Proper Citation

R/METASIM (RRID:SCR\_009370)

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### 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 application, software resource

**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:** 20260727T040500+0000

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### 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](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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