

Resource Summary Report

Generated by [NIF](#) on Aug 29, 2026

[sim4cc](#)

RRID:SCR_001204

Type: Tool

Proper Citation

sim4cc (RRID:SCR_001204)

Resource Information

URL: <http://ccb.jhu.edu/software/sim4cc/>

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Description: Software tool as cross species spliced alignment program. Heuristic sequence alignment tool for comparing cDNA sequence with genomic sequence containing homolog of gene in another species.

Resource Type: alignment software, data processing software, image analysis software, software application, software resource

Defining Citation: [PMID:19429899](#)

Keywords: Cross species spliced alignment, unix, sequence alignment, cDNA sequence, genomic sequence, homolog, gene, splice, bio.tools

Funding: NLM R01 LM006845;
NSF CLS20163A;
Sloan Research Fellowship

Availability: Free, Available for download, Freely available

Resource Name: sim4cc

Resource ID: SCR_001204

Alternate IDs: biotools:sim4cc, OMICS_02145

Alternate URLs: <https://bio.tools/sim4cc>

License: GNU General Public License, v3

Record Creation Time: 20220129T080206+0000

Record Last Update: 20260829T182037+0000

Ratings and Alerts

No rating or validation information has been found for sim4cc.

No alerts have been found for sim4cc.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.