Resource Summary Report

Generated by dkNET on May 18, 2025

sim4cc

RRID:SCR_001204

Type: Tool

Proper Citation

sim4cc (RRID:SCR_001204)

Resource Information

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

Proper Citation: sim4cc (RRID:SCR_001204)

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

Defining Citation: PMID:19429899

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

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

Availability: Free, 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: 20250517T055458+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.