## **Resource Summary Report**

Generated by RRID on Apr 29, 2025

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

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: 20250429T054642+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: <u>SciCrunch Registry</u>

### **Usage and Citation Metrics**

We have not found any literature mentions for this resource.