

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

crb-blast

RRID:SCR_024325

Type: Tool

Proper Citation

crb-blast (RRID:SCR_024325)

Resource Information

URL: <https://github.com/cboursnell/crb-blast>

Proper Citation: crb-blast (RRID:SCR_024325)

Description: Software tool for finding orthologs between one set of sequences and another. This is particularly useful in genome and transcriptome annotation.

Synonyms: Conditional Reciprocal Best Blast

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

Keywords: finding orthologs between one set of sequences and another, genome and transcriptome annotation,

Availability: Free, Available for download, Freely available,

Resource Name: crb-blast

Resource ID: SCR_024325

Alternate URLs: <https://sources.debian.org/src/ruby-crb-blast/>

License: MIT license

Record Creation Time: 20230830T050217+0000

Record Last Update: 20260905T133030+0000

Ratings and Alerts

No rating or validation information has been found for crb-blast.

No alerts have been found for crb-blast.

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](#) .

Feigin CY, et al. (2023) Convergent deployment of ancestral functions during the evolution of mammalian flight membranes. Science advances, 9(12), eade7511.