

# Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

## [ruby-bio](#)

RRID:SCR\_024322

Type: Tool

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### Proper Citation

ruby-bio (RRID:SCR\_024322)

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### Resource Information

**URL:** <https://bioruby.org/>

**Proper Citation:** ruby-bio (RRID:SCR\_024322)

**Description:** Software tools and libraries for bioinformatics and molecular biology, for the Ruby programming language. BioRuby has components for sequence analysis, pathway analysis, protein modelling and phylogenetic analysis; it supports many widely used data formats and provides easy access to databases, external programs and public web services, including BLAST, KEGG, GenBank, MEDLINE and GO.

**Resource Type:** software resource, software toolkit

**Keywords:** Ruby programming language, sequence analysis, pathway analysis, protein modelling, phylogenetic analysis,

**Availability:** Free, Available for download, Freely available,

**Resource Name:** ruby-bio

**Resource ID:** SCR\_024322

**Alternate URLs:** <https://sources.debian.org/src/ruby-bio/>

**Record Creation Time:** 20230830T050217+0000

**Record Last Update:** 20260905T133108+0000

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### Ratings and Alerts

No rating or validation information has been found for ruby-bio.

No alerts have been found for ruby-bio.

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### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.