

Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

[ruby-bio](#)

RRID:SCR_024322

Type: Tool

Proper Citation

ruby-bio (RRID:SCR_024322)

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: 20260801T190850+0000

Ratings and Alerts

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

No alerts have been found for ruby-bio.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.