

Resource Summary Report

Generated by RRID on Aug 11, 2026

Ruby on Rails

RRID:SCR_022129

Type: Tool

Proper Citation

Ruby on Rails (RRID:SCR_022129)

Resource Information

URL: <https://rubyonrails.org/>

Proper Citation: Ruby on Rails (RRID:SCR_022129)

Description: Open source web application development framework. Model View Controller framework, providing default structures for database, web service, and web pages.

Abbreviations: RoR

Resource Type: web application, software library, software toolkit, software resource

Defining Citation: [DOI:10.1109/MS.2007.176](https://doi.org/10.1109/MS.2007.176)

Keywords: MVC, model–view–controller, web application development framework, Rails

Availability: Free, Available for download, Freely available

Resource Name: Ruby on Rails

Resource ID: SCR_022129

Alternate IDs: <https://www.wikidata.org/wiki/Q190478>

Alternate URLs: <https://github.com/rails/rails>

License: <https://github.com/rails/rails/blob/main/MIT-LICENSE>

Record Creation Time: 20220421T050138+0000

Record Last Update: 20260811T043700+0000

Ratings and Alerts

No rating or validation information has been found for Ruby on Rails.

No alerts have been found for Ruby on Rails.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 10 mentions in open access literature.

Listed below are recent publications. The full list is available at [RRID](#) .

Kruger R, et al. (2026) MiMeDB 2.0: the Human Microbial Metabolome Database for 2026. Nucleic acids research, 54(D1), D623.

Jackson H, et al. (2025) MarkerDB 2.0: a comprehensive molecular biomarker database for 2025. Nucleic acids research, 53(D1), D1415.

Wishart DS, et al. (2025) The Natural Products Magnetic Resonance Database (NP-MRD) for 2025. Nucleic acids research, 53(D1), D700.

Diedrich K, et al. (2024) User-centric design of a 3D search interface for protein-ligand complexes. Journal of computer-aided molecular design, 38(1), 23.

Knox C, et al. (2024) DrugBank 6.0: the DrugBank Knowledgebase for 2024. Nucleic acids research, 52(D1), D1265.

Wishart DS, et al. (2024) PathBank 2.0-the pathway database for model organism metabolomics. Nucleic acids research, 52(D1), D654.

Gupta C, et al. (2024) Enhancing bug allocation in software development: a multi-criteria approach using fuzzy logic and evolutionary algorithms. PeerJ. Computer science, 10, e2111.

Neveu V, et al. (2023) The human microbial exposome: expanding the Exposome-Explorer database with gut microbial metabolites. Scientific reports, 13(1), 1946.

Diedrich K, et al. (2023) PoseEdit: enhanced ligand binding mode communication by interactive 2D diagrams. Journal of computer-aided molecular design, 37(10), 491.

Mitsuhashi N, et al. (2022) TogoVar: A comprehensive Japanese genetic variation database. Human genome variation, 9(1), 44.