

Resource Summary Report

Generated by [RRID](#) on Aug 4, 2026

[ROSIE](#)

RRID:SCR_018764

Type: Tool

Proper Citation

ROSIE (RRID:SCR_018764)

Resource Information

URL: <https://rosie.graylab.jhu.edu/docking2>

Proper Citation: ROSIE (RRID:SCR_018764)

Description: Unified web framework for Rosetta applications. Web interface for selected Rosetta protocols. Web front end for Rosetta software suite. Provides common user interface for Rosetta protocols, stable application programming interface for developers to add additional protocols, flexible back-end to allow leveraging of computer cluster resources shared by Rosetta Commons member institutions, and centralized administration by Rosetta Commons to ensure continuous maintenance. Offers general and speedy paradigm for serverification of Rosetta applications. Lowers barriers to Rosetta use for broader biological community.

Abbreviations: ROSIE

Synonyms: Rosetta Online Server that Includes Everyone

Resource Type: data access protocol, software resource, application programming interface, web application

Defining Citation: [PMID:23717507](#)

Keywords: Web interface for Rosetta, Rosetta online server, Rosetta application serverification, Rosetta user interface

Funding: NIGMS R01 GM073151;
NIGMS R01 GM07822;
NIGMS R21 GM102716;
NCRR R00 RR024107;
NCI U54 CA143907;
NEI PN2 EY016586;
NIGMS T32 GM 88118;
Taiwan Governmental Scholarship for Study Abroad ;

Howard Hughes Medical and Institute International Student Research Fellowship ; NSF

Availability: Restricted

Resource Name: ROSIE

Resource ID: SCR_018764

Alternate URLs: <https://rosie.rosettacommons.org/>

Record Creation Time: 20220129T080341+0000

Record Last Update: 20260804T043703+0000

Ratings and Alerts

No rating or validation information has been found for ROSIE.

No alerts have been found for ROSIE.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 7 mentions in open access literature.

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

Wang Y, et al. (2025) A unique mechanism of snRNP core assembly. Nature communications, 16(1), 3166.

Krapež G, et al. (2025) In Vitro Functional Validation of an Anti-FREM2 Nanobody for Glioblastoma Cell Targeting. Antibodies (Basel, Switzerland), 14(1).

Xi C, et al. (2024) Manipulating the molecular specificity of transcriptional biosensors for tryptophan metabolites and analogs. Cell reports. Physical science, 5(10).

Das NK, et al. (2024) Structural basis for a highly conserved RNA-mediated enteroviral genome replication. Nucleic acids research, 52(18), 11218.

Nieto-Fabregat F, et al. (2024) Computational toolbox for the analysis of protein-glycan interactions. Beilstein journal of organic chemistry, 20, 2084.

Yunaini L, et al. (2023) In silico docking analysis of beta-defensin 20 against cation channel sperm-associated protein 1-4 to predict its role in the sperm maturation. Asian journal of andrology, 25(4), 528.

Solorza J, et al. (2021) Effects of Interleukin-1? in Glycinergic Transmission at the Central Amygdala. Frontiers in pharmacology, 12, 613105.