

Resource Summary Report

Generated by [nidm-terms](#) on Jul 28, 2026

RiboSnake

RRID:SCR_025608

Type: Tool

Proper Citation

RiboSnake (RRID:SCR_025608)

Resource Information

URL: <https://github.com/IKIM-Essen/RiboSnake>

Proper Citation: RiboSnake (RRID:SCR_025608)

Description: Software 16S rRNA analysis workflow with QIIME2 and Snakemake.

Resource Type: software resource, source code, software application, data processing software, workflow software

Defining Citation: [DOI:10.1101/2024.08.06.606757](https://doi.org/10.1101/2024.08.06.606757)

Keywords: 16S rRNA analysis workflow, 16S rRNA, analysis workflow,

Funding: Medical Informatics Initiative Germany

Availability: Free, Available for download, Freely available,

Resource Name: RiboSnake

Resource ID: SCR_025608

License: MIT license

Record Creation Time: 20240813T053254+0000

Record Last Update: 20260728T043727+0000

Ratings and Alerts

No rating or validation information has been found for RiboSnake.

No alerts have been found for RiboSnake.

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 [nidm-terms](#) .

D??rr AK, et al. (2024) RiboSnake - a user-friendly, robust, reproducible, multipurpose and documentation-extensive pipeline for 16S rRNA gene microbiome analysis. GigaByte (Hong Kong, China), 2024, gigabyte132.