

# Resource Summary Report

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## 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:** data processing software, software application, software resource, source code, 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:** 20260912T200448+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](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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.