

Resource Summary Report

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

dadasnake

RRID:SCR_019149

Type: Tool

Proper Citation

dadasnake (RRID:SCR_019149)

Resource Information

URL: <https://github.com/a-h-b/dadasnake>

Proper Citation: dadasnake (RRID:SCR_019149)

Description: Software tool as snakemake implementation of DADA2 to process amplicon sequencing data for microbial ecology.

Resource Type: software application, software resource, data processing software

Defining Citation: [DOI:10.1101/2020.05.17.095679](https://doi.org/10.1101/2020.05.17.095679)

Keywords: Snakemake, DADA2, amplicon sequencing data, microbial ecology

Availability: Free, Available for download, Freely available

Resource Name: dadasnake

Resource ID: SCR_019149

License: GNU GPLv3.0

Record Creation Time: 20220129T080343+0000

Record Last Update: 20260727T040730+0000

Ratings and Alerts

No rating or validation information has been found for dadasnake.

No alerts have been found for dadasnake.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Gebauer L, et al. (2021) Soil Texture, Sampling Depth and Root Hairs Shape the Structure of ACC Deaminase Bacterial Community Composition in Maize Rhizosphere. *Frontiers in microbiology*, 12, 616828.

Wei??becker C, et al. (2020) Dadasnake, a Snakemake implementation of DADA2 to process amplicon sequencing data for microbial ecology. *GigaScience*, 9(12).