Resource Summary Report

Generated by <u>NIF</u> on May 19, 2025

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 resource, software application, data processing software

Defining Citation: DOI:10.1101/2020.05.17.095679

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

Funding:

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: 20250519T204541+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 <u>NIF</u>.

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.

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