

Resource Summary Report

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16SMicrobiomeMLFS

RRID:SCR_027170

Type: Tool

Proper Citation

16SMicrobiomeMLFS (RRID:SCR_027170)

Resource Information

URL: <https://github.com/nach00gar/16SMicrobiomeMLFS>

Proper Citation: 16SMicrobiomeMLFS (RRID:SCR_027170)

Description: Code and supporting data for the article: "Exploring the role of normalization and feature selection in microbiome disease classification pipelines."

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

Keywords: Microbiome, Amplicon, Machine Learning, Feature Selection

Related Condition: Arthritis, Clostridium difficile Infection, Colorectal Cancer, Human Immunodeficiency Virus, Crohn's Disease, Inflammatory Bowel Disease, Cirrhosis, Minimal Hepatic Encephalopathy, Obesity, Parkinson's Disease

Availability: Free, Available for download, Freely available

Resource Name: 16SMicrobiomeMLFS

Resource ID: SCR_027170

License: GNU GPL v3

Record Creation Time: 20250709T061412+0000

Record Last Update: 20260801T042951+0000

Ratings and Alerts

No rating or validation information has been found for 16SMicrobiomeMLFS.

No alerts have been found for 16SMicrobiomeMLFS.

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 [kravitz2](#) .

Garach Vélez I, et al. (2025) Exploring the role of normalization and feature selection in microbiome disease classification pipelines. GigaScience, 14.