

Resource Summary Report

Generated by [RRID](#) on Aug 15, 2026

[gNOMO2](#)

RRID:SCR_025293

Type: Tool

Proper Citation

gNOMO2 (RRID:SCR_025293)

Resource Information

URL: <https://github.com/muzafferarikan/gNOMO2>

Proper Citation: gNOMO2 (RRID:SCR_025293)

Description: Software as comprehensive and modular pipeline for integrated multi-omics analyses of microbiomes. Enables integration of taxonomic and functional analyses in microbiome multi-omics data.

Resource Type: software toolkit, source code, software resource

Defining Citation: [DOI:10.1101/2023.12.03.569767](https://doi.org/10.1101/2023.12.03.569767)

Keywords: microbiome, multi-omics, data integration, amplicon sequencing, metagenomics, metatranscriptomics, metaproteomics

Funding: TUBITAK

Availability: Free, Available for download, Freely available

Resource Name: gNOMO2

Resource ID: SCR_025293

License: MIT license

Record Creation Time: 20240430T053222+0000

Record Last Update: 20260815T042934+0000

Ratings and Alerts

No rating or validation information has been found for gNOMO2.

No alerts have been found for gNOMO2.

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 [RRID](#) .

Arikan M, et al. (2024) gNOMO2: a comprehensive and modular pipeline for integrated multi-omics analyses of microbiomes. GigaScience, 13.