

Resource Summary Report

Generated by T1D on Aug 2, 2026

METAREP

RRID:SCR_011926

Type: Tool

Proper Citation

METAREP (RRID:SCR_011926)

Resource Information

URL: <http://www.jcvi.org/metarep/>

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Description: A tool for high-performance comparative metagenomics that allows users to view, query, browse, and compare metagenomics annotation profiles from short reads or assemblies. Users can use statistical tests, hierarchical clustering, multidimensional scaling, and heat maps to compare multiple datasets at various functional and taxonomic levels.

Abbreviations: METAREP

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

Defining Citation: [DOI:10.1093/bioinformatics/btq455](https://doi.org/10.1093/bioinformatics/btq455)

Keywords: microbiome, comparison, comparative metagenomics, annotation, short read, short assembly, bio.tools

Availability: Open source

Resource Name: METAREP

Resource ID: SCR_011926

Alternate IDs: biotools:metarep, OMICS_01480

Alternate URLs: <https://bio.tools/metarep>

License: Open unspecified license

Record Creation Time: 20220129T080307+0000

Record Last Update: 20260803T040604+0000

Ratings and Alerts

No rating or validation information has been found for METAREP.

No alerts have been found for METAREP.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 8 mentions in open access literature.

Listed below are recent publications. The full list is available at [T1D](#) .

Aalismail NA, et al. (2019) Functional metagenomic analysis of dust-associated microbiomes above the Red Sea. Scientific reports, 9(1), 13741.

Bagarolli RA, et al. (2017) Probiotics modulate gut microbiota and improve insulin sensitivity in DIO mice. The Journal of nutritional biochemistry, 50, 16.

Díez B, et al. (2016) Metagenomic Analysis of the Indian Ocean Picocyanobacterial Community: Structure, Potential Function and Evolution. PloS one, 11(5), e0155757.

Garg N, et al. (2016) Spatial Molecular Architecture of the Microbial Community of a Peltigera Lichen. mSystems, 1(6).

Dudhagara P, et al. (2015) Web Resources for Metagenomics Studies. Genomics, proteomics & bioinformatics, 13(5), 296.

Llorens-Marès T, et al. (2015) Connecting biodiversity and potential functional role in modern euxinic environments by microbial metagenomics. The ISME journal, 9(7), 1648.

Mandal RS, et al. (2015) Metagenomic surveys of gut microbiota. Genomics, proteomics & bioinformatics, 13(3), 148.

Prakash T, et al. (2012) Functional assignment of metagenomic data: challenges and applications. Briefings in bioinformatics, 13(6), 711.