

Resource Summary Report

Generated by [RRID](#) on Aug 5, 2026

Metab

RRID:SCR_016877

Type: Tool

Proper Citation

Metab (RRID:SCR_016877)

Resource Information

URL: <https://github.com/reinkk/Metab>

Proper Citation: Metab (RRID:SCR_016877)

Description: Software package as a metabolomic data processing pipeline in R codes.

Abbreviations: Metab

Synonyms: Metabolome, Metab 1.0

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

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

Keywords: metabolomic, data, processing, pipeline, analysis, datasets

Funding: Health Research Council of New Zealand

Availability: Free, Available for download, Freely available

Resource Name: Metab

Resource ID: SCR_016877

Alternate URLs: <https://bioconductor.org/packages/release/bioc/html/Metab.html>

License: GPL

Record Creation Time: 20220129T080332+0000

Record Last Update: 20260805T044400+0000

Ratings and Alerts

No rating or validation information has been found for Metab.

No alerts have been found for Metab.

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](#) .

Sulek K, et al. (2014) Hair metabolomics: identification of fetal compromise provides proof of concept for biomarker discovery. Theranostics, 4(9), 953.