

# Resource Summary Report

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

## MetaPhyl

RRID:SCR\_001169

Type: Tool

### Proper Citation

MetaPhyl (RRID:SCR\_001169)

### Resource Information

**URL:** <http://www.cs.ucr.edu/~tanaseio/metaphyl.htm>

**Proper Citation:** MetaPhyl (RRID:SCR\_001169)

**Description:** Software implementing a supervised classification method for metagenomic samples that takes advantage of the natural structure of microbial community data encoded by phylogenetic trees.

**Abbreviations:** MetaPhyl

**Resource Type:** software resource

**Defining Citation:** [PMID:24369151](#)

**Keywords:** c++, linux, mac osx, phylogenetic tree, command line, classification, metagenomic

**Availability:** Free, Available for download, Freely available

**Resource Name:** MetaPhyl

**Resource ID:** SCR\_001169

**Alternate IDs:** OMICS\_02189

**Record Creation Time:** 20220129T080206+0000

**Record Last Update:** 20260801T190124+0000

### Ratings and Alerts

No rating or validation information has been found for MetaPhyl.

No alerts have been found for MetaPhyl.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.