

# Resource Summary Report

Generated by [RRID](#) on Aug 2, 2026

## [phylobase](#)

RRID:SCR\_024292

Type: Tool

---

### Proper Citation

phylobase (RRID:SCR\_024292)

---

### Resource Information

**URL:** <https://cran.r-project.org/package=phylobase>

**Proper Citation:** phylobase (RRID:SCR\_024292)

**Description:** Software R package provides base S4 class for comparative methods, incorporating one or more trees and trait data. Used for phylogenetic analysis.

**Resource Type:** software resource, software toolkit

**Keywords:** associate, manipulate, explore, plot, phylogenetic trees and data, phylogenetic analysis,

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

**Resource Name:** phylobase

**Resource ID:** SCR\_024292

**Alternate URLs:** <https://sources.debian.org/src/r-cran-phylobase/>

**License:** GPL-2 | GPL-3 [expanded from: GPL (? 2)]

**Record Creation Time:** 20230830T050217+0000

**Record Last Update:** 20260801T190850+0000

---

### Ratings and Alerts

No rating or validation information has been found for phylobase.

No alerts have been found for phylobase.

---

### 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](#) .

Chaguza C, et al. (2023) The population-level impact of Enterococcus faecalis genetics on intestinal colonization and extraintestinal infection. Microbiology spectrum, 11(6), e0020123.