# **Resource Summary Report**

Generated by NIF on Apr 26, 2025

# **PhySamp**

RRID:SCR\_024159

Type: Tool

### **Proper Citation**

PhySamp (RRID:SCR\_024159)

#### **Resource Information**

**URL:** https://jydu.github.io/physamp/

**Proper Citation:** PhySamp (RRID:SCR\_024159)

Description: Software package dedicated to phylogenetic sampling. Used to sample

sequence alignment according to its corresponding phylogenetic tree.

Synonyms: physamp

Resource Type: software toolkit, software library, software resource

**Defining Citation:** PMID:26055961

**Keywords:** phylogenetic sampling, sample sequence alignment, corresponding phylogenetic

tree,

Funding:

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

Resource Name: PhySamp

Resource ID: SCR\_024159

Alternate IDs: OMICS\_21686

Alternate URLs: https://sources.debian.org/src/physamp/

**Record Creation Time: 20230824T050212+0000** 

**Record Last Update:** 20250426T061003+0000

## **Ratings and Alerts**

No rating or validation information has been found for PhySamp.

No alerts have been found for PhySamp.

### **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 <u>NIF</u>.

Meteyer CU, et al. (2022) Plant pathogens provide clues to the potential origin of bat white-nose syndrome Pseudogymnoascus destructans. Virulence, 13(1), 1020.