

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

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 library, software resource, software toolkit

Defining Citation: [PMID:26055961](#)

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

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: 20260905T133104+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 [Kravitz](#) .

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