

Resource Summary Report

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

Phyx

RRID:SCR_024164

Type: Tool

Proper Citation

Phyx (RRID:SCR_024164)

Resource Information

URL: <https://github.com/FePhyFoFum/phyx>

Proper Citation: Phyx (RRID:SCR_024164)

Description: Software phylogenetic tools for unix. Used to perform phylogenetics analyses on trees and sequences. Collection of programs written in C ++ to explore, manipulate, analyze and simulate phylogenetic objects.

Resource Type: software resource, software library, software toolkit

Defining Citation: [PMID:28174903](#)

Keywords: phylogenetic tools for unix, perform phylogenetics analyses on trees and sequences, C ++, explore phylogenetic objects, manipulate phylogenetic objects, analyze and simulate phylogenetic objects,

Availability: Free, Available for download, Freely available,

Resource Name: Phyx

Resource ID: SCR_024164

Alternate URLs: <https://sources.debian.org/src/phyx/>

License: GPL-3.0 license

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260801T190853+0000

Ratings and Alerts

No rating or validation information has been found for Phyx.

No alerts have been found for Phyx.

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](#) .

Chia KS, et al. (2024) The N-terminal domains of NLR immune receptors exhibit structural and functional similarities across divergent plant lineages. The Plant cell, 36(7), 2491.