**PHYLIP**

RRID:SCR_006244  
Type: Tool

**Proper Citation**

PHYLIP (RRID:SCR_006244)

**Resource Information**


**Description:** A free package of software programs for inferring phylogenies (evolutionary trees). The source code is distributed (in C), and executables are also distributed. In particular, already-compiled executables are available for Windows (95/98/NT/2000/me/xp/Vista), Mac OS X, and Linux systems. Older executables are also available for Mac OS 8 or 9 systems.

**Resource Name:** PHYLIP

**Proper Citation:** PHYLIP (RRID:SCR_006244)

**Resource Type:** Resource, software resource, software application, data processing software, source code

**Keywords:** phylogeny prediction, evolutionary tree

**Resource ID:** SCR_006244

**Parent Organization:** University of Washington; Seattle; USA

**Funding Agency:** DOE, NIGMS, NSF

**Availability:** Free

**Website Status:** Last checked up

**Alternate IDs:** nif-0000-06708
Abbreviations: PHYLIP

Mentions Count: 2938

Ratings and Alerts

No rating or validation information has been found for PHYLIP.

No alerts have been found for PHYLIP.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 2938 mentions in open access literature.

Listed below are recent publications. The full list is available at FDI Lab - SciCrunch Infrastructure.


Csáky V, et al. (2020) Genetic insights into the social organisation of the Avar period elite in the 7th century AD Carpathian Basin. Scientific reports, 10(1), 948.


Deng L, et al. (2020) Occurrence and genetic characteristics of Cryptosporidium spp. and Enterocytozoon bieneusi in pet red squirrels (Sciurus vulgaris) in China. Scientific reports, 10(1), 1026.