

Resource Summary Report

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

Bio-Tools-Phylo-PAML

RRID:SCR_024069

Type: Tool

Proper Citation

Bio-Tools-Phylo-PAML (RRID:SCR_024069)

Resource Information

URL: <https://metacpan.org/dist/Bio-Tools-Phylo-PAML>

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Description: Software package used to parse output from the PAML programs codeml, baseml, basemlg, codemlsites and yn00. You can use the Bio-Tools-Run-Phylo-PAML modules to actually run some of the PAML programs, but this module is only useful to parse the output.

Synonyms: libbio-tools-phylo-paml-perl, Bio-Tools-Phylo-PAML - Parses output from the PAML programs codeml, baseml, codemlsites and yn00, basemlg

Resource Type: software resource, software toolkit

Keywords: parse output, PAML programs, codeml, baseml, basemlg, codemlsites, yn00.

Availability: Free, Available for download, Freely available,

Resource Name: Bio-Tools-Phylo-PAML

Resource ID: SCR_024069

Alternate URLs: <https://sources.debian.org/src/libbio-tools-phylo-paml-perl/>

License: perl_5

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260905T133103+0000

Ratings and Alerts

No rating or validation information has been found for Bio-Tools-Phylo-PAML.

No alerts have been found for Bio-Tools-Phylo-PAML.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.