

# Resource Summary Report

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## Bio-Tools-Phylo-PAML

RRID:SCR\_024069

Type: Tool

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### Proper Citation

Bio-Tools-Phylo-PAML (RRID:SCR\_024069)

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### Resource Information

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

**Proper Citation:** Bio-Tools-Phylo-PAML (RRID:SCR\_024069)

**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:** 20260821T194248+0000

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### 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.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We have not found any literature mentions for this resource.