RAxML
RRID:SCR_006086
Type: Tool

Proper Citation
RAxML (RRID:SCR_006086)

Resource Information

URL: https://github.com/stamatak/standard-RAxML

Description: Software program for phylogenetic analyses of large datasets under maximum likelihood.

Resource Name: RAxML
Proper Citation: RAxML (RRID:SCR_006086)

Resource Type: Resource, software resource, software application, data analysis software, data processing software

Keywords: phylogeny

Resource ID: SCR_006086

References: PMID: 24451623, PMID: 16928733, PMID: 15608047

Availability: GNU General Public License

Website Status: Last checked up

Alternate IDs: OMICS_02242

Abbreviations: RAxML

Mentions Count: 5576

Ratings and Alerts
No rating or validation information has been found for RAxML.

No alerts have been found for RAxML.

Data and Source Information
Source: SciCrunch Registry

Usage and Citation Metrics

We found 5576 mentions in open access literature.

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


Ding X, et al. (2020) Genome sequence of the agarwood tree Aquilaria sinensis (Lour.) Spreng: the first chromosome-level draft genome in the Thymelaeaceae family. GigaScience, 9(3).


Adrián-Kalchauser I, et al. (2020) The round goby genome provides insights into mechanisms that may facilitate biological invasions. BMC biology, 18(1), 11.


