Kalign

RRID:SCR_011810
Type: Tool

Proper Citation

Kalign (RRID:SCR_011810)

Resource Information

URL: http://www.ebi.ac.uk/Tools/msa/kalign/
Proper Citation: Kalign (RRID:SCR_011810)
Description: A fast and accurate multiple sequence alignment algorithm.
Abbreviations: Kalign
Resource Type: software resource
Defining Citation: PMID:16343337, DOI:10.1093/bioinformatics/btz795
Keywords: bio.tools
Availability: Free
Resource Name: Kalign
Resource ID: SCR_011810
Alternate IDs: biotools:kalign, OMICS_00978
Alternate URLs: https://bio.tools/kalign, https://sources.debian.org/src/kalign/

Ratings and Alerts

No rating or validation information has been found for Kalign.

No alerts have been found for Kalign.
Usage and Citation Metrics

We found 103 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [RRID](#).


Mewa DB, et al. (2023) ANTHRACNOSE RESISTANCE GENE2 confers fungal resistance in sorghum. The Plant journal: for cell and molecular biology, 113(2), 308.


Liedtke HC, et al. (2022) Chromosome-level assembly, annotation and phylome of Pelobates cultripes, the western spadefoot toad. DNA research: an international journal for rapid publication of reports on genes and genomes, 29(3).


Tan Y, et al. (2021) Unification and extensive diversification of M/Orf3-related ion channel proteins in coronaviruses and other nidoviruses. Virus evolution, 7(1), veab014.