Kalign
RRID:SCR_011810
Type: Tool

**Proper Citation**

Kalign (RRID:SCR_011810)

**Resource Information**

**URL:** [http://www.ebi.ac.uk/Tools/msa/kalign/](http://www.ebi.ac.uk/Tools/msa/kalign/)

**Description:** A fast and accurate multiple sequence alignment algorithm.

**Resource Name:** Kalign

**Proper Citation:** Kalign (RRID:SCR_011810)

**Resource Type:** Resource, software resource

**Resource ID:** SCR_011810

**Parent Organization:** European Bioinformatics Institute

**References:** [PMID: 16343337]

**Availability:** Free

**Website Status:** Last checked up

**Alternate IDs:** OMICS_00978

**Abbreviations:** Kalign

**Mentions Count:** 64

**Ratings and Alerts**

No rating or validation information has been found for Kalign.
No alerts have been found for Kalign.

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Data and Source Information

Source: SciCrunch Registry

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Usage and Citation Metrics

We found 64 mentions in open access literature.

Listed below are recent publications. The full list is available at scicrunch.


Julca I, et al. (2018) Phylogenomics of the olive tree (Olea europaea) reveals the relative
contribution of ancient allo- and autopolyploidization events. BMC biology, 16(1), 15.


