

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 7, 2026

PCMA

RRID:SCR_024156

Type: Tool

Proper Citation

PCMA (RRID:SCR_024156)

Resource Information

URL: <http://prodata.swmed.edu/pcma/pcma.php>

Proper Citation: PCMA (RRID:SCR_024156)

Description: Software tool for multiple sequence alignment based on profile consistency. Used to construct multiple sequence alignment given set of protein sequences.

Synonyms: pcma, Profile Consistency Multiple Sequence Alignment

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

Defining Citation: [PMID:12584134](#)

Keywords: multiple sequence alignment, profile consistency, construct multiple sequence alignment, protein sequences,

Availability: Free, Available for download, Freely available,

Resource Name: PCMA

Resource ID: SCR_024156

Alternate IDs: OMICS_14997

Old URLs: <https://sources.debian.org/src/pcma/>

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260905T133029+0000

Ratings and Alerts

No rating or validation information has been found for PCMA.

No alerts have been found for PCMA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.