

Resource Summary Report

Generated by SPARC SAWG on Aug 1, 2026

POA

RRID:SCR_024172

Type: Tool

Proper Citation

POA (RRID:SCR_024172)

Resource Information

URL: <https://sourceforge.net/projects/poamsa/>

Proper Citation: POA (RRID:SCR_024172)

Description: Software application for multiple sequence alignment in bioinformatics. Has superior ability to handle branching / indels in the alignment.

Synonyms: poa, Partial Order Alignment

Resource Type: software resource, software library, software toolkit

Defining Citation: [DOI:10.1093/bioinformatics/bth126](https://doi.org/10.1093/bioinformatics/bth126)

Keywords: sequence alignment, multiple sequence alignment, handle branching in alignment, handle indels in alignment,

Availability: Free, Available for download, Freely available,

Resource Name: POA

Resource ID: SCR_024172

Alternate IDs: OMICS_14254

Alternate URLs: <https://sources.debian.org/src/poa/>

License: GNU General Public License version 2.0 (GPLv2)

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260801T190853+0000

Ratings and Alerts

No rating or validation information has been found for POA.

No alerts have been found for POA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.