

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

PSAR-Align

RRID:SCR_011814

Type: Tool

Proper Citation

PSAR-Align (RRID:SCR_011814)

Resource Information

URL: <http://bioen-compbio.bioen.illinois.edu/PSAR-Align/>

Proper Citation: PSAR-Align (RRID:SCR_011814)

Description: Software for improving multiple sequence alignment using probabilistic sampling.

Abbreviations: PSAR-Align

Synonyms: PSAR-Align: improving multiple sequence alignment using probabilistic sampling

Resource Type: software resource

Defining Citation: [PMID:24222208](#)

Keywords: c++, bio.tools

Availability: Free

Resource Name: PSAR-Align

Resource ID: SCR_011814

Alternate IDs: OMICS_00987, biotools:psar

Alternate URLs: <https://bio.tools/psar>

Record Creation Time: 20220129T080306+0000

Record Last Update: 20260905T132712+0000

Ratings and Alerts

No rating or validation information has been found for PSAR-Align.

No alerts have been found for PSAR-Align.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Mochales-Riaño G, et al. (2025) Chromosome-level reference genome for the medically important Arabian horned viper (*Cerastes gasperettii*). GigaScience, 14.