

Resource Summary Report

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[eProbalign](#)

RRID:SCR_013247

Type: Tool

Proper Citation

eProbalign (RRID:SCR_013247)

Resource Information

URL: <http://probalign.njit.edu/probalign/login>

Proper Citation: eProbalign (RRID:SCR_013247)

Description: Data analysis service that computes maximal expected accuracy multiple sequence alignments from partition function posterior probabilities.

Synonyms: eProbalign web server, EProbalign

Resource Type: analysis service resource, production service resource, data analysis service, service resource

Defining Citation: [PMID:17485479](#)

Keywords: multiple sequence alignments, partition function posterior probabilities, bio.tools

Funding: NIGMS R01 GM073082

Resource Name: eProbalign

Resource ID: SCR_013247

Alternate IDs: OMICS_00975, biotools:eprobalign

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

Record Creation Time: 20220129T080315+0000

Record Last Update: 20260819T044347+0000

Ratings and Alerts

No rating or validation information has been found for eProbalign.

No alerts have been found for eProbalign.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.