

Resource Summary Report

Generated by [kravitz2](#) on Jul 30, 2026

NW-align

RRID:SCR_024138

Type: Tool

Proper Citation

NW-align (RRID:SCR_024138)

Resource Information

URL: <https://zhanggroup.org/NW-align/>

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Description: Software tool as alignment program for protein sequence-to-sequence alignments based on the standard Needleman-Wunsch dynamic programming algorithm.

Synonyms: nw-align

Resource Type: software application, data access protocol, software resource, data processing software, alignment software, web service, image analysis software

Keywords: protein sequence-to-sequence alignments, standard Needleman-Wunsch dynamic programming algorithm,

Availability: Free, Freely available,

Resource Name: NW-align

Resource ID: SCR_024138

Old URLs: <https://sources.debian.org/src/nw-align/>

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260729T044602+0000

Ratings and Alerts

No rating or validation information has been found for NW-align.

No alerts have been found for NW-align.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.