Resource Summary Report

Generated by <u>NIF</u> on Apr 28, 2025

NW-align

RRID:SCR_024138 Type: Tool

Proper Citation

NW-align (RRID:SCR_024138)

Resource Information

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

Proper Citation: NW-align (RRID:SCR_024138)

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

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

Funding:

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: 20250428T054408+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.