Resource Summary Report

Generated by NIF on Apr 29, 2025

JAligner

RRID:SCR_024050

Type: Tool

Proper Citation

JAligner (RRID:SCR_024050)

Resource Information

URL: https://github.com/ahmedmoustafa/JAligner

Proper Citation: JAligner (RRID:SCR_024050)

Description: Open source software Java implementation of the Needleman? Wunsch and Smith-Waterman algorithms for biological pairwise sequence alignment with the affine gap penalty model.

Synonyms: jaligner

Resource Type: data processing software, image analysis software, alignment software, software resource, software application

Keywords: Java, pairwise sequence alignment, affine gap penalty model,

Funding:

Availability: Free, Available for download, Freely available,

Resource Name: JAligner

Resource ID: SCR_024050

Alternate IDs: OMICS_20021

Alternate URLs: https://sources.debian.org/src/jaligner/

License: GPL-2.0 license

Record Creation Time: 20230824T050211+0000

Record Last Update: 20250429T060309+0000

Ratings and Alerts

No rating or validation information has been found for JAligner.

No alerts have been found for JAligner.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.