Resource Summary Report

Generated by dkNET on Apr 18, 2025

muliAlignFree

RRID:SCR_013188

Type: Tool

Proper Citation

muliAlignFree (RRID:SCR_013188)

Resource Information

URL: http://www-rcf.usc.edu/~fsun/Programs/multiAlignFree/multiAlignFreemain.html

Proper Citation: muliAlignFree (RRID:SCR_013188)

Description: R package intended to implement a program for multiple alignment-free

sequence comparison based on long genome sequence or NGS data.

Abbreviations: muliAlignFree

Synonyms: muliAlignFree: Multiple Alignment-free Sequence Comparison

Resource Type: software resource

Defining Citation: PMID:23990418

Funding:

Availability: Free

Resource Name: muliAlignFree

Resource ID: SCR_013188

Alternate IDs: OMICS_00981

Record Creation Time: 20220129T080314+0000

Record Last Update: 20250410T070341+0000

Ratings and Alerts

No rating or validation information has been found for muliAlignFree.

No alerts have been found for muliAlignFree.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We have not found any literature mentions for this resource.