Resource Summary Report

Generated by NIF on Apr 25, 2025

SeqLib

RRID:SCR_024091

Type: Tool

Proper Citation

SeqLib (RRID:SCR_024091)

Resource Information

URL: https://github.com/walaj/SeqLib

Proper Citation: SeqLib (RRID:SCR_024091)

Description: Software C++ htslib/bwa-mem/fermi interface for interrogating sequence data

Synonyms: libseqlib

Resource Type: software resource, software application

Defining Citation: DOI:10.1093/bioinformatics/btw741

Keywords: interrogating sequence data,

Funding:

Availability: Free, Available for download, Freely available,

Resource Name: SeqLib

Resource ID: SCR_024091

Alternate IDs: OMICS_14682

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

License: Apache License, Version 2.0

Record Creation Time: 20230824T050211+0000

Record Last Update: 20250421T054511+0000

Ratings and Alerts

No rating or validation information has been found for SeqLib.

No alerts have been found for SeqLib.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 1 mentions in open access literature.

Listed below are recent publications. The full list is available at NIF.

De Mita S, et al. (2011) Molecular adaptation in flowering and symbiotic recognition pathways: insights from patterns of polymorphism in the legume Medicago truncatula. BMC evolutionary biology, 11, 229.