

Resource Summary Report

Generated by [NIF](#) on Jul 31, 2026

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++ htlib/bwa-mem/fermi interface for interrogating sequence data

Synonyms: libseqlib

Resource Type: software resource, software application

Defining Citation: [DOI:10.1093/bioinformatics/btw741](https://doi.org/10.1093/bioinformatics/btw741)

Keywords: interrogating sequence data,

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: 20260731T043154+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.