

Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 18, 2026

SeQuiLa

RRID:SCR_017220

Type: Tool

Proper Citation

SeQuiLa (RRID:SCR_017220)

Resource Information

URL: <https://github.com/ZSI-Bio/bdg-sequila>

Proper Citation: SeQuiLa (RRID:SCR_017220)

Description: Software tool for genomic intervals querying and processing built on top of Apache Spark. Elastic, fast and scalable SQL oriented solution for processing and querying genomic intervals.

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

Defining Citation: [PMID:30428005](#)

Keywords: genomic, intervals, querying, processing, Apache, Spark, SQL, analysis, bio.tools

Funding: National Science Center ;
Polish budget funds for science

Availability: Free, Available for download, Freely available

Resource Name: SeQuiLa

Resource ID: SCR_017220

Alternate IDs: biotools:SeQuiLa-cov

Alternate URLs: <http://biodatageeks.org/sequila/>, <https://bio.tools/SeQuiLa-cov>

License: Apache License 2.0

Record Creation Time: 20220129T080334+0000

Record Last Update: 20260818T043607+0000

Ratings and Alerts

No rating or validation information has been found for SeQuiLa.

No alerts have been found for SeQuiLa.

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 [SPARC SAWG](#) .

Wiewi??rka M, et al. (2019) SeQuiLa-cov: A fast and scalable library for depth of coverage calculations. GigaScience, 8(8).