

# Resource Summary Report

Generated by T1D on Aug 10, 2026

## SeQuiLa

RRID:SCR\_017220

Type: Tool

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### Proper Citation

SeQuiLa (RRID:SCR\_017220)

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### 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 analysis software, software resource, software application, data processing software

**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:** 20260810T040715+0000

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## Ratings and Alerts

No rating or validation information has been found for SeQuiLa.

No alerts have been found for SeQuiLa.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [T1D](#) .

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