

# Resource Summary Report

Generated by [NIF](#) on Aug 19, 2026

## SeqPig

RRID:SCR\_008548

Type: Tool

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### Proper Citation

SeqPig (RRID:SCR\_008548)

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### Resource Information

**URL:** <http://seqpig.sourceforge.net/>

**Proper Citation:** SeqPig (RRID:SCR\_008548)

**Description:** A software library for Apache Pig for the distributed analysis of large sequencing datasets on Hadoop clusters.

**Abbreviations:** SeqPig

**Resource Type:** software resource

**Defining Citation:** [PMID:24149054](#)

**Keywords:** mapreduce/hadoop

**Resource Name:** SeqPig

**Resource ID:** SCR\_008548

**Alternate IDs:** OMICS\_01226

**Record Creation Time:** 20220129T080248+0000

**Record Last Update:** 20260815T182346+0000

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### Ratings and Alerts

No rating or validation information has been found for SeqPig.

No alerts have been found for SeqPig.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Maarala AI, et al. (2021) Distributed hybrid-indexing of compressed pan-genomes for scalable and fast sequence alignment. PloS one, 16(8), e0255260.

Schumacher A, et al. (2014) SeqPig: simple and scalable scripting for large sequencing data sets in Hadoop. Bioinformatics (Oxford, England), 30(1), 119.