

Resource Summary Report

Generated by T1D on Aug 5, 2026

POP BAM

RRID:SCR_000464

Type: Tool

Proper Citation

POP BAM (RRID:SCR_000464)

Resource Information

URL: <https://sourceforge.net/projects/popbam/>

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Description: A tool to perform evolutionary or population-based analyses of next-generation sequencing data. POP BAM takes a BAM file as its input and can compute many widely used evolutionary genetics measures in sliding windows across a genome.

Abbreviations: POP BAM

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

Defining Citation: [PMID:24027417](https://pubmed.ncbi.nlm.nih.gov/24027417/)

Keywords: next-generation sequencing, evolution, population, bam, genome, evolutionary genetics, c++, short read, sequence alignment, sliding window, command-line, bio.tools

Availability: Free, Available for download, Freely available

Resource Name: POP BAM

Resource ID: SCR_000464

Alternate IDs: biotools:popbam, OMICS_01559

Alternate URLs: <https://bio.tools/popbam>

Old URLs: <http://popbam.sourceforge.net/>

License: BSD License, MIT License

Record Creation Time: 20220129T080201+0000

Record Last Update: 20260805T044008+0000

Ratings and Alerts

No rating or validation information has been found for POPBAM.

No alerts have been found for POPBAM.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.