

Resource Summary Report

Generated by [FDI Lab - SciCrunch.org](#) on Apr 16, 2025

POP BAM

RRID:SCR_000464

Type: Tool

Proper Citation

POP BAM (RRID:SCR_000464)

Resource Information

URL: <http://popbam.sourceforge.net/>

Proper Citation: POP BAM (RRID:SCR_000464)

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

Defining Citation: [PMID:24027417](#)

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

Funding:

Availability: BSD License, MIT License

Resource Name: POP BAM

Resource ID: SCR_000464

Alternate IDs: biotools:popbam, OMICS_01559

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

Record Creation Time: 20220129T080201+0000

Record Last Update: 20250416T063220+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.