

Resource Summary Report

Generated by PRECISE-TBI on Sep 18, 2026

BAMseek

RRID:SCR_000672

Type: Tool

Proper Citation

BAMseek (RRID:SCR_000672)

Resource Information

URL: <https://code.google.com/p/bamseek/>

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Description: A Large File Viewer for BAM and SAM alignment files.

Abbreviations: BAMseek

Synonyms: bamseek - Browse large BAM and SAM alignment files.

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

Keywords: matlab, sam, bam, viewer, alignment, format, large file, vcf, sff, fastq, sequencing

Availability: Free, Available for download, Freely available

Resource Name: BAMseek

Resource ID: SCR_000672

Alternate IDs: OMICS_00877

License: GNU General Public License, v3

Record Creation Time: 20220129T080202+0000

Record Last Update: 20260912T195513+0000

Ratings and Alerts

No rating or validation information has been found for BAMseek.

No alerts have been found for BAMseek.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.