SAMtools/BCFtools

RRID:SCR_005227
Type: Tool

Proper Citation

SAMtools/BCFtools (RRID:SCR_005227)

Resource Information

**URL:** [http://samtools.sourceforge.net/mpileup.shtml](http://samtools.sourceforge.net/mpileup.shtml)

**Description:** Provide various utilities for manipulating alignments in the SAM format, including sorting, merging, indexing and generating alignments in a per-position format.

**Resource Name:** SAMtools/BCFtools

**Proper Citation:** SAMtools/BCFtools (RRID:SCR_005227)

**Resource Type:** Resource, software resource

**Keywords:** snp, indel

**Resource ID:** SCR_005227

**Parent Organization:** SAMTOOLS

**Website Status:** Last checked up

**Alternate IDs:** OMICS_00090

**Abbreviations:** BCFtools

**Mentions Count:** 134

Ratings and Alerts

No rating or validation information has been found for SAMtools/BCFtools.

No alerts have been found for SAMtools/BCFtools.
Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 134 mentions in open access literature.

**Listed below are recent publications.** The full list is available at FDI Lab - SciCrunch Infrastructure.


Perdigão J, et al. (2020) Using genomics to understand the origin and dispersion of multidrug and extensively drug resistant tuberculosis in Portugal. Scientific reports, 10(1), 2600.


