Resource Summary Report

Generated by FDI Lab - SciCrunch.org on Apr 13, 2025

Bio-SamTools

RRID:SCR_024066
Type: Tool

Proper Citation

Bio-SamTools (RRID:SCR_024066)

Resource Information

URL: https://metacpan.org/dist/Bio-SamTools

Proper Citation: Bio-SamTools (RRID:SCR_024066)

Description: Software Perl interface to SamTools library for DNA sequencing.

Synonyms: libbio-samtools-perl

Resource Type: software application, software resource

Keywords: Perl interface, SamTools library, DNA sequencing,

Funding:

Availability: Free, Available for download, Freely available,

Resource Name: Bio-SamTools

Resource ID: SCR_024066

Alternate URLs: https://sources.debian.org/src/libbio-samtools-perl/

License: perl_5

Record Creation Time: 20230824T050211+0000

Record Last Update: 20250412T060622+0000

Ratings and Alerts

No rating or validation information has been found for Bio-SamTools.

No alerts have been found for Bio-SamTools.

Data and Source Information

Source: SciCrunch Registry

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

Sandhu N, et al. (2024) Development of Novel KASP Markers for Improved Germination in Deep-Sown Direct Seeded Rice. Rice (New York, N.Y.), 17(1), 33.

Shanmugam V, et al. (2014) Whole genome sequencing reveals potential targets for therapy in patients with refractory KRAS mutated metastatic colorectal cancer. BMC medical genomics, 7, 36.

Collins JE, et al. (2012) Incorporating RNA-seq data into the zebrafish Ensembl genebuild. Genome research, 22(10), 2067.

Weiss GJ, et al. (2012) Paired tumor and normal whole genome sequencing of metastatic olfactory neuroblastoma. PloS one, 7(5), e37029.

Demeure MJ, et al. (2012) Cancer of the ampulla of Vater: analysis of the whole genome sequence exposes a potential therapeutic vulnerability. Genome medicine, 4(7), 56.