

Resource Summary Report

Generated by [dkNET](#) on Aug 27, 2026

[pysam](#)

RRID:SCR_021017

Type: Tool

Proper Citation

pysam (RRID:SCR_021017)

Resource Information

URL: <https://pysam.readthedocs.io/en/latest/api.html>

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Description: Software tool as interface for reading and writing SAM files. Python module to read and manipulate mapped short read sequence data stored in SAM/BAM files. Lightweight wrapper of htlib C-API.

Resource Type: software resource, software toolkit

Keywords: Interface, reading SAM files, writing SAM files, mapped short read sequence data, htlib C-API wrapper, SAM/BAM files

Availability: Free, Available for download, Freely available

Resource Name: pysam

Resource ID: SCR_021017

Alternate URLs: <https://github.com/pysam-developers/pysam>

License: MIT License

Record Creation Time: 20220129T080353+0000

Record Last Update: 20260821T194148+0000

Ratings and Alerts

No rating or validation information has been found for pysam.

No alerts have been found for pysam.

Data and Source Information

Usage and Citation Metrics

We found 164 mentions in open access literature.

Listed below are recent publications. The full list is available at [dkNET](#) .

Wäneskog M, et al. (2026) High-throughput methods enabling random duplications, deletions, or nucleotide-constrained mutagenesis of entire DNA motifs. *Nucleic acids research*, 54(6).

Chen L, et al. (2026) Animal-associated jumbo phages as widespread and active modulators of gut microbiome ecology and metabolism. *Science advances*, 12(6), eaeb6265.

Pastore S, et al. (2026) Mapping human pre-rRNA processing and modification at single nucleotide resolution using long read nanopore sequencing. *Nature communications*, 17(1).

Simmons SK, et al. (2026) Experimental and computational methods for allelic imbalance analysis from single-nucleus RNA-seq data. *Genome biology*, 27(1).

Hallast P, et al. (2026) Population-scale Y chromosome assemblies reveal recurrent remodeling within constrained architectures. *bioRxiv : the preprint server for biology*.

Rishik S, et al. (2026) SmallBARN 2026: a kingdom-wide bacterial sRNA resource. *Nucleic acids research*, 54(D1), D342.

Jeong R, et al. (2026) Functional analysis of NPR2 variants supports the therapeutic rationale for CNP in short stature. *American journal of human genetics*, 113(1), 117.

Engin B, et al. (2026) Atlas of HIV cis-regulatory elements reveals extensive transcriptional variation across clades, isolates, and within individuals. *bioRxiv : the preprint server for biology*.

Bucher P, et al. (2026) CAR-adapted PIK3CD base editing enhances T cell anti-tumor potency. *Nature cancer*, 7(2), 368.

Ben Chehida Y, et al. (2026) Genetic parallelism underpins convergent mimicry coloration in Lepidoptera across 120 million years of evolution. *PLoS biology*, 24(4), e3003742.

Kolling Iv FW, et al. (2026) Multisite Assessment of Methods for Cell Preservation Upstream of Single-Cell RNA Sequencing. *Journal of biomolecular techniques : JBT*, 37(2), 9.

Hunt M, et al. (2026) Addressing pandemic-wide systematic errors in the SARS-CoV-2 phylogeny. *Nature methods*, 23(3), 653.

Choi JH, et al. (2026) Universal conditional knockout approach to multiple species, from fish to human induced pluripotent stem cells. *Nucleic acids research*, 54(9).

Ji HJ, et al. (2026) Accurate serotype identification of *Streptococcus pneumoniae* using nanopore Cas9-targeted serotype identification (nCATSerotyping). *Journal of clinical microbiology*, 64(2), e0098425.

Jasnauskaitė M, et al. (2026) Structure and mechanism of antiphage retron Eco2. *Nature structural & molecular biology*, 33(2), 330.

Dziagwa C, et al. (2026) Disruption of tRNA threonylation triggers RIG-I mediated anti-tumour immune response. *Nature communications*, 17(1).

Kim JW, et al. (2025) Pairwise analysis of plasma cell-free DNA before and after palliative second-line paclitaxel plus ramucirumab treatment in patients with metastatic gastric cancer. *Gastric cancer : official journal of the International Gastric Cancer Association and the Japanese Gastric Cancer Association*, 28(4), 620.

Momen-Roknabadi A, et al. (2025) Detection of Early-Stage Colorectal Cancer Using Cell-Free oncRNA Biomarkers and Artificial Intelligence. *Clinical cancer research : an official journal of the American Association for Cancer Research*, 31(15), 3229.

Seluzicki CM, et al. (2025) Regulation of translation elongation and integrated stress response in heat-shocked neurons. *Cell reports*, 44(5), 115639.

Calderón-Osorno M, et al. (2025) The influence of depth on the global deep-sea plasmidome. *Scientific reports*, 15(1), 2959.