

Resource Summary Report

Generated by [ASWG](#) on Aug 3, 2026

HTSJDK

RRID:SCR_024036

Type: Tool

Proper Citation

HTSJDK (RRID:SCR_024036)

Resource Information

URL: <http://samtools.github.io/htsjdk/>

Proper Citation: HTSJDK (RRID:SCR_024036)

Description: Software implementation of unified Java library for accessing common file formats, such as SAM and VCF, used for high-throughput sequencing data. There are also an number of useful utilities for manipulating HTS data.

Synonyms: htsjdk

Resource Type: software resource, software application

Keywords: accessing common file formats, SAM, VCF, manipulating HTS data,

Availability: Free, Available for download, Freely available,

Resource Name: HTSJDK

Resource ID: SCR_024036

Alternate URLs: <https://sources.debian.org/src/htsjdk/>, <https://github.com/samtools/htsjdk>

License: MIT license/ Apache License v2.0/ LGPL

Record Creation Time: 20230824T050211+0000

Record Last Update: 20260801T191117+0000

Ratings and Alerts

No rating or validation information has been found for HTSJDK.

No alerts have been found for HTSJDK.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Wen X, et al. (2025) Deciphering splicing heterogeneity at single-cell resolution by SCSES. Nature communications, 16(1), 9459.

Simmons SK, et al. (2024) Experimental and Computational Methods for Allelic Imbalance Analysis from Single-Nucleus RNA-seq Data. bioRxiv : the preprint server for biology.