

# Resource Summary Report

Generated by [SPARC SAWG](#) on Aug 9, 2026

## KAnalyze

RRID:SCR\_001323

Type: Tool

---

### Proper Citation

KAnalyze (RRID:SCR\_001323)

---

### Resource Information

**URL:** <http://sourceforge.net/projects/kanalyze/>

**Proper Citation:** KAnalyze (RRID:SCR\_001323)

**Description:** A Java toolkit designed to convert DNA and RNA sequences into k-mers.

**Resource Type:** software resource

**Defining Citation:** [PMID:24642064](#)

**Keywords:** standalone software, java, bio.tools

**Availability:** Free, Available for download, Freely available

**Resource Name:** KAnalyze

**Resource ID:** SCR\_001323

**Alternate IDs:** biotools:kanalyze, OMICS\_03565

**Alternate URLs:** <https://bio.tools/kanalyze>

**License:** GNU Lesser General Public License, v3

**Record Creation Time:** 20220129T080206+0000

**Record Last Update:** 20260808T185726+0000

---

### Ratings and Alerts

No rating or validation information has been found for KAnalyze.

No alerts have been found for KAnalyze.

## 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 [SPARC SAWG](#) .

Manekar SC, et al. (2018) A benchmark study of k-mer counting methods for high-throughput sequencing. GigaScience, 7(12).

Flygare S, et al. (2016) Taxonomer: an interactive metagenomics analysis portal for universal pathogen detection and host mRNA expression profiling. Genome biology, 17(1), 111.