

# Resource Summary Report

Generated by [NIF](#) on Sep 11, 2026

## Atropos

RRID:SCR\_023962

Type: Tool

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### Proper Citation

Atropos (RRID:SCR\_023962)

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### Resource Information

**URL:** <https://github.com/jdidion/atropos>

**Proper Citation:** Atropos (RRID:SCR\_023962)

**Description:** Software tool for specific, sensitive, and speedy trimming of NGS reads.

**Synonyms:** atropos

**Resource Type:** software application, software resource, source code

**Defining Citation:** [PMID:28875074](#)

**Keywords:** speedy trimming of NGS reads, NGS reads, Next Generation Sequencing,

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

**Resource Name:** Atropos

**Resource ID:** SCR\_023962

**Alternate IDs:** OMICS\_20869

**Alternate URLs:** <https://sources.debian.org/src/atropos/>

**License URLs:** <https://github.com/jdidion/atropos/blob/1.1/LICENSE>

**Record Creation Time:** 20230824T050210+0000

**Record Last Update:** 20260905T133102+0000

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### Ratings and Alerts

No rating or validation information has been found for Atropos.

No alerts have been found for Atropos.

## Data and Source Information

**Source:** [SciCrunch Registry](#)

## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Michel M, et al. (2025) Noninvasive Multicancer Detection Using DNA Hypomethylation of LINE-1 Retrotransposons. Clinical cancer research : an official journal of the American Association for Cancer Research, 31(7), 1275.

Griffiths JA, et al. (2024) Peripheral neuronal activation shapes the microbiome and alters gut physiology. Cell reports, 43(4), 113953.

Henon C, et al. (2024) Single-cell multiomics profiling reveals heterogeneous transcriptional programs and microenvironment in DSRCTs. Cell reports. Medicine, 5(6), 101582.