

# Resource Summary Report

Generated by T1D on Aug 1, 2026

## NanoLyse

RRID:SCR\_024125

Type: Tool

### Proper Citation

NanoLyse (RRID:SCR\_024125)

### Resource Information

**URL:** <https://github.com/wdecoster/nanolyse>

**Proper Citation:** NanoLyse (RRID:SCR\_024125)

**Description:** Software package to remove reads mapping to the lambda phage genome from a fastq file.

**Synonyms:** nanolyse

**Resource Type:** software resource, software library, software toolkit

**Keywords:** remove reads mapping, lambda phage genome, fastq file,

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

**Resource Name:** NanoLyse

**Resource ID:** SCR\_024125

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

**License:** GPL-3.0 license

**Record Creation Time:** 20230824T050212+0000

**Record Last Update:** 20260801T042916+0000

### Ratings and Alerts

No rating or validation information has been found for NanoLyse.

No alerts have been found for NanoLyse.

### Data and Source Information

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

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## Usage and Citation Metrics

We found 5 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [T1D](#) .

Ritter EJ, et al. (2025) The assembly and annotation of two teinturier grapevine varieties, Dakapo and Rubired. GigaByte (Hong Kong, China), 2025, gigabyte149.

Webster VL, et al. (2024) Revealing the genome of the microsporidian Vairimorpha bombi, a potential driver of bumble bee declines in North America. G3 (Bethesda, Md.), 14(4).

Sugimura Y, et al. (2024) Impact of rice GENERAL REGULATORY FACTOR14h (GF14h) on low-temperature seed germination and its application to breeding. PLoS genetics, 20(8), e1011369.

Roothans N, et al. (2024) Aerobic denitrification as an N<sub>2</sub>O source from microbial communities. The ISME journal, 18(1).

Hemmings SJ, et al. (2023) Long-read Sequencing and de novo Genome Assembly of Three Aspergillus fumigatus Genomes. Mycopathologia, 188(4), 409.