

Resource Summary Report

Generated by [NIF](#) on Jul 30, 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 toolkit, software resource, software library

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: 20260731T043144+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](#)

Usage and Citation Metrics

We found 5 mentions in open access literature.

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

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 N₂O 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.