

Resource Summary Report

Generated by [nidm-terms](#) on Sep 23, 2026

MView

RRID:SCR_024129

Type: Tool

Proper Citation

MView (RRID:SCR_024129)

Resource Information

URL: <https://desmid.github.io/mview/>

Proper Citation: MView (RRID:SCR_024129)

Description: Web-compatible database search or multiple alignment viewer, Software tool as command line utility that extracts and reformats the results of a sequence database search or a multiple alignment, optionally adding HTML markup for web page layout. It can also be used as a filter to extract and convert searches or alignments to common formats.

Synonyms: mview

Resource Type: software application, software resource

Defining Citation: [PMID:9632837](#)

Keywords: database search, multiple alignment viewer,

Availability: Free, Available for download, Freely available,

Resource Name: MView

Resource ID: SCR_024129

Alternate IDs: OMICS_15907

Old URLs: <https://sources.debian.org/src/mview/>

License: GNU General Public License v2 or any later version

Record Creation Time: 20230824T050212+0000

Record Last Update: 20260919T195828+0000

Ratings and Alerts

No rating or validation information has been found for MView.

No alerts have been found for MView.

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 [nidm-terms](#) .

Sun W, et al. (2026) Antimicrobial resistance, virulence defects, and evolutionary dynamics of multidrug-resistant *Klebsiella pneumoniae* from human and animal hosts in Central China. *Microbiology spectrum*, 14(2), e0226925.

Orkwis JA, et al. (2026) Helminth proteins recapitulate key structural motifs of MIF to facilitate immunomodulation of host receptors. *iScience*, 29(7), 116513.

Nishio S, et al. (2024) ZP2 cleavage blocks polyspermy by modulating the architecture of the egg coat. *Cell*, 187(6), 1440.

Anders J, et al. (2023) RNAcode_Web - Convenient identification of evolutionary conserved protein coding regions. *Journal of integrative bioinformatics*, 20(3).

Hermida Lorenzo RJ, et al. (2021) Metagenomic Snapshots of Viral Components in Guinean Bats. *Microorganisms*, 9(3).