

Resource Summary Report

Generated by PRECISE-TBI on Sep 8, 2026

MINERVA

RRID:SCR_026679

Type: Tool

Proper Citation

MINERVA (RRID:SCR_026679)

Resource Information

URL: <https://minerva.uni.lu>

Proper Citation: MINERVA (RRID:SCR_026679)

Description: Standalone web server for visualization, exploration and management of molecular networks encoded in SBGN-compliant format.

Abbreviations: MINERVA

Synonyms: Molecular Interaction NEtwoRk VisuAlization

Resource Type: software resource, web application

Keywords: standalone webserver for visualization, exploration and management, molecular networks, SBGN-compliant format,

Availability: Free, Freely available

Resource Name: MINERVA

Resource ID: SCR_026679

License: Affero GPL

License URLs: <https://minerva.pages.uni.lu/doc/tos/>

Record Creation Time: 20250401T060330+0000

Record Last Update: 20260905T133528+0000

Ratings and Alerts

No rating or validation information has been found for MINERVA.

No alerts have been found for MINERVA.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 4 mentions in open access literature.

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

Balaur I, et al. (2025) FAIR assessment of Disease Maps fosters open science and scientific crowdsourcing in systems biomedicine. *Scientific data*, 12(1), 851.

Hemedan AA, et al. (2025) Protocol to identify regulatory modules in Parkinson's disease progression using miRNA data and Boolean modeling. *STAR protocols*, 6(2), 103769.

Hemedan AA, et al. (2024) Cohort-specific boolean models highlight different regulatory modules during Parkinson's disease progression. *iScience*, 27(10), 110956.

Gawron P, et al. (2023) Visualization of automatically combined disease maps and pathway diagrams for rare diseases. *Frontiers in bioinformatics*, 3, 1101505.