

Resource Summary Report

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GseaVis

RRID:SCR_028574

Type: Tool

Proper Citation

GseaVis (RRID:SCR_028574)

Resource Information

URL: <https://github.com/junjunlab/GseaVis>

Proper Citation: GseaVis (RRID:SCR_028574)

Description: Software R package for enhanced visualization of gene set enrichment analysis in biomedicine. Used to visualize Gene Set Enrichment Analysis (GSEA) results.

Resource Type: software resource, software toolkit, source code

Defining Citation: [DOI:10.1002/mdr2.70000](https://doi.org/10.1002/mdr2.70000)

Keywords: Gene Set Enrichment Analysis, visualization of gene set enrichment analysis,

Funding: China Pharmaceutical University ;
National Key Research and Development Program of China ;
National Natural Science Foundation of China

Availability: Free, Available for download, Freely available

Resource Name: GseaVis

Resource ID: SCR_028574

License: MIT license

Record Creation Time: 20260619T043006+0000

Record Last Update: 20260919T200134+0000

Ratings and Alerts

No rating or validation information has been found for GseaVis.

No alerts have been found for GseaVis.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 34 mentions in open access literature.

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

Niu S, et al. (2026) TBC1 domain family member 23 is essential for STING-mediated anti-melanoma effect. *Molecular biomedicine*, 7(1).

Luo M, et al. (2026) H2AC14 is a novel and reliable prognostic marker for CRC. *Medicine*, 105(2), e47026.

Lu X, et al. (2026) Molecular taxonomy of pancreatic neuroendocrine tumors reveals BEND2-fusions-driven transcriptional plasticity and therapeutic vulnerabilities. *Cell reports. Medicine*, 7(3), 102642.

Rong Y, et al. (2026) Oxygen-sensing histone demethylase KDM6A modulates chondrocyte-to-osteoblast transdifferentiation by activating the Wnt/catenin pathway. *International journal of molecular medicine*, 58(2).

Wang B, et al. (2026) Integrated single-cell RNA sequencing and Bulk-RNA technologies reveal the immunological characteristics of lactylation related-genes in glioblastoma. *PloS one*, 21(6), e0351849.

Sun S, et al. (2026) The comprehensive landscape of TTMV::RARA fusion-driven acute myeloid leukemia: from viral integration mechanisms to clinical outcomes. *Haematologica*, 111(4), 1254.

Kopylova IV, et al. (2026) Convergent transcriptomic signature in iPSC-dopaminergic neurons of hereditary Parkinson's disease. *Life science alliance*, 9(7).

Yi X, et al. (2026) Dibutyl phthalate induces sarcopenia via TNF/TNFR1-mediated proteolytic and pyroptotic axes: evidence from NHANES and experimental models. *Frontiers in immunology*, 17, 1853039.

Cheng X, et al. (2026) Reversing adenosine-mediated immunosuppression in triple-negative breast cancer by synergistic chemo-immunotherapy via stimuli-responsive nanomedicines. *EBioMedicine*, 123, 106059.

Zheng X, et al. (2026) Integrative bioinformatics and experiments identify RIBC2 as a key regulator in the esophageal cancer. *PloS one*, 21(2), e0340850.

Xie X, et al. (2026) Targeted inhibition of PARP-1 in pulmonary epithelial cells and macrophages via SPA-functionalized microparticles attenuates sepsis-induced lung injury. *Materials today. Bio*, 37, 102955.

Zhang W, et al. (2026) SULT1E1 suppresses hepatocellular carcinoma progression via activation of the TGF- β /SMAD signaling pathway and cell cycle arrest. *BMC gastroenterology*, 26(1).

Li C, et al. (2026) Serotonergic System-Targeted Nucleic Acid Hydrogel Coordinates Excitability Restoration and Circuit Reconstruction for Spinal Cord Injury Therapy. *Advanced materials* (Deerfield Beach, Fla.), 38(11), e21427.

Chen S, et al. (2026) Single-Cell and Machine Learning Analysis Reveal Novel Inflammatory Macrophage Subtypes and Biomarkers in Periodontitis. *International dental journal*, 76(1), 103983.

Hu X, et al. (2026) B7-H3 interaction with cMet enhances lipolysis to drive malignant progression of glioma via pSTAT3-ATGL axis. *Journal of translational medicine*, 24(1).

Wu L, et al. (2025) KCTD9 expression predicts immunotherapy response and enhances anti-PD-1 efficacy in colon adenocarcinoma. *International journal of surgery* (London, England), 111(10), 6755.

Zhang Z, et al. (2025) Preliminary Analysis of Placental DNA Methylation Profiles in Piglets with Extreme Birth Weight Variations. *Animals : an open access journal from MDPI*, 15(15).

Sun P, et al. (2025) Disruption of tryptophan metabolism by high-fat diet-triggered maternal immune activation promotes social behavioral deficits in male mice. *Nature communications*, 16(1), 2105.

Zhou Z, et al. (2025) Identification and external validation of a prognostic signature based on hypoxia-glycolysis-related genes for kidney renal clear cell carcinoma. *Open medicine* (Warsaw, Poland), 20(1), 20251305.

Yue L, et al. (2025) Human pluripotent stem cell-derived skin organoids enabled pathophysiological model of *Mycobacterium tuberculosis* infection. *Nature communications*, 16(1), 10831.