

Resource Summary Report

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scRNAtoolVis

RRID:SCR_023916

Type: Tool

Proper Citation

scRNAtoolVis (RRID:SCR_023916)

Resource Information

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

Proper Citation: scRNAtoolVis (RRID:SCR_023916)

Description: Software package to make scRNA-seq plot.

Resource Type: software resource, software application

Keywords: scRNA-seq plot, scRNA-seq, scRNA-seq data visualization, plot, graph

Funding:

Availability: Free, Available for download, Freely available

Resource Name: scRNAtoolVis

Resource ID: SCR_023916

Record Creation Time: 20230809T050222+0000

Record Last Update: 20250513T062427+0000

Ratings and Alerts

No rating or validation information has been found for scRNAtoolVis.

No alerts have been found for scRNAtoolVis.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 6 mentions in open access literature.

Listed below are recent publications. The full list is available at [FDI Lab - SciCrunch.org](#).

Yu Q, et al. (2025) Causal genes identification of giant cell arteritis in CD4+?Memory t cells: an integration of multi-omics and expression quantitative trait locus analysis. Inflammation research : official journal of the European Histamine Research Society ... [et al.], 74(1), 3.

Chai D, et al. (2025) Lipid nanoparticles deliver DNA-encoded biologics and induce potent protective immunity. Molecular cancer, 24(1), 12.

Danev N, et al. (2024) Comparative transcriptomic analysis of bovine mesenchymal stromal cells reveals tissue-source and species-specific differences. iScience, 27(2), 108886.

Lu R, et al. (2024) 3D spheroid culture synchronizes heterogeneous MSCs into an immunomodulatory phenotype with enhanced anti-inflammatory effects. iScience, 27(9), 110811.

Chen G, et al. (2024) Bulk and single-cell alternative splicing analyses reveal roles of TRA2B in myogenic differentiation. Cell proliferation, 57(2), e13545.

Chen YZ, et al. (2024) HMGB2 drives tumor progression and shapes the immunosuppressive microenvironment in hepatocellular carcinoma: insights from multi-omics analysis. Frontiers in immunology, 15, 1415435.