

Resource Summary Report

Generated by [Kravitz](#) on Sep 11, 2026

single cell RNA-seq Data Visualization and Analysis

RRID:SCR_023598

Type: Tool

Proper Citation

single cell RNA-seq Data Visualization and Analysis (RRID:SCR_023598)

Resource Information

URL: http://cancer-pku.cn:3838/CRC_Leukocyte/

Proper Citation: single cell RNA-seq Data Visualization and Analysis (RRID:SCR_023598)

Description: Interactive web server to explore and analyze single cell RNA-seq data. R shiny based tool used for single cell RNA-seq data visualization and analysis.

Abbreviations: scDVA

Resource Type: data access protocol, software resource, web service

Defining Citation: [PMID:32302573](#)

Keywords: single cell RNA-seq data, data visualization and analysis service, data visualization and analysis,

Availability: Free, Freely available

Resource Name: single cell RNA-seq Data Visualization and Analysis

Resource ID: SCR_023598

Alternate URLs: <https://github.com/liziyie/scDVA>

Record Creation Time: 20230523T050222+0000

Record Last Update: 20260905T133024+0000

Ratings and Alerts

No rating or validation information has been found for single cell RNA-seq Data Visualization and Analysis.

No alerts have been found for single cell RNA-seq Data Visualization and Analysis.

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 [Kravitz](#) .

Yan C, et al. (2026) ALKBH5/IGF2BP1-mediated m6A demethylation of TRIM37 promotes pancreatic cancer tumorigenesis and glycolysis by mediating RBMX degradation. Cell biology and toxicology, 42(1).

Cui J, et al. (2025) Mechanism of dexmedetomidine in inhibiting proliferation, migration, and invasion of esophageal squamous cell carcinoma cell through coagulation factor II receptor. Discover oncology, 16(1), 1112.

Zhu T, et al. (2025) PD-L1/ITGB4 Axis Modulates Sensitivity of Hepatocellular Carcinoma to Sorafenib via FAK/AKT/mTOR Signaling Pathway. ImmunoTargets and therapy, 14, 815.

Chen X, et al. (2022) RUNX3/H3K27me3 Co-Expression Defines a Better Prognosis in Surgically Resected Stage I and Postoperative Chemotherapy-Naive Non-Small-Cell Lung Cancer. Journal of oncology, 2022, 5752263.

Zhu H, et al. (2022) Comprehensive bioinformatics analysis for MEF2 family genes in gastric cancer. Translational cancer research, 11(11), 4057.