

Resource Summary Report

Generated by [RRID](#) on Sep 10, 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 [RRID](#) .

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.