

Resource Summary Report

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scDbfFinder

RRID:SCR_022700

Type: Tool

Proper Citation

scDbfFinder (RRID:SCR_022700)

Resource Information

URL: <https://github.com/plger/scDbfFinder>

Proper Citation: scDbfFinder (RRID:SCR_022700)

Description: Software package for detection and handling of doublets/multiplets in single cell sequencing data.

Resource Type: software toolkit, software resource

Defining Citation: [PMID:35814628](#)

Keywords: single cell sequencing data, doublets detection, multiplets detection, single cell, sequencing data

Availability: Free, Available for download, Freely available

Resource Name: scDbfFinder

Resource ID: SCR_022700

Alternate URLs: <https://bioconductor.org/packages/scDbfFinder/>

License: GNU General Public License v3.0

Record Creation Time: 20220825T050144+0000

Record Last Update: 20260810T040822+0000

Ratings and Alerts

No rating or validation information has been found for scDbfFinder.

No alerts have been found for scDbfFinder.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 174 mentions in open access literature.

Listed below are recent publications. The full list is available at [kravitz2](#) .

Gao Y, et al. (2026) Altered Inflammatory Signature in a C9ORF72 -ALS iPSC-Derived Motor Neuron and Microglia Coculture Model. *Glia*, 74(1), e70084.

Vandenbempt V, et al. (2026) PTPN2 deficiency amplifies inflammatory signalling and impairs functional maturation of human stem cell-derived islets. *Stem cell research & therapy*, 17(1).

Zhang Y, et al. (2026) Co-occurrence of transcriptionally distinct persister cell states underpins neoadjuvant therapy resistance in triple-negative breast cancer. *Genome medicine*, 18(1).

Simmons SK, et al. (2026) Experimental and computational methods for allelic imbalance analysis from single-nucleus RNA-seq data. *Genome biology*, 27(1).

Wang H, et al. (2026) Peripheral monocyte ratio as a predictive biomarker for metastasis-directed therapy in prostate cancer: insights from an exploratory study. *BMC cancer*, 26(1).

Sethna CR, et al. (2026) Genome-wide cell type-specific and sex-specific transcriptional dysregulation in the islet of Langerhans underlies islet dysfunction in Down syndrome-related diabetes. *bioRxiv : the preprint server for biology*.

Zamboni M, et al. (2026) The regulatory code of injury-responsive enhancers enables precision cell-state targeting in the CNS. *Nature neuroscience*, 29(2), 337.

Wu CC, et al. (2026) Serial Spatial Transcriptomes Reveal Regulatory Transitions in Maize Leaf Development. *Plant biotechnology journal*, 24(5), 2787.

Fritz L, et al. (2026) Differentiation-induced reduction in functional diversity restricts the ability of cytomegalovirus-specific CD8 T cells to eliminate virus-infected cells. *EBioMedicine*, 123, 106107.

Lee H, et al. (2026) A multi-omic single-cell landscape of perinatal mouse skin maps lineage specification and reveals shared dynamics in human fetal skin. *Experimental & molecular medicine*, 58(4), 1269.

Xiong Y, et al. (2026) Molecular and Chromatin Accessibility Programs Underlying Epithelial Injury and Impaired Regeneration in Neonatal Necrotizing Enterocolitis. *Cellular and molecular gastroenterology and hepatology*, 20(6), 101730.

Anilkumar Sithara A, et al. (2026) Composition and Functional State of T and NK Cells in the Extramedullary Myeloma Tumor Microenvironment. *Blood cancer discovery*, 7(2), 250.

Yashar WM, et al. (2026) PU.1 inhibition sensitizes stem-monocytic AML to BCL2 blockade. *bioRxiv : the preprint server for biology*.

Johnson LA, et al. (2026) Clonal expansion of cytotoxic CD8⁺ T cells in lecanemab-associated ARIA. *Nature communications*, 17(1).

Sun H, et al. (2026) Identification and characterization of MARCO-expressing tumor-associated macrophages in pancreatic ductal adenocarcinoma with pan-cancer relevance. *NPJ precision oncology*, 10(1).

Kroehling L, et al. (2026) A highly resolved integrated single-cell atlas of HPV-negative head and neck cancer. *Communications medicine*, 6(1).

Aalling NN, et al. (2026) Sustained diabetes remission induced by FGF1 involves a shift in transcriptionally distinct AgRP neuron subpopulations. *Molecular metabolism*, 103, 102300.

Valdivieso K, et al. (2026) Transcription-independent induction of rapid-onset senescence is integral to healing. *Nature cell biology*, 28(6), 1281.

Singh S, et al. (2026) Temporal single-cell transcriptional dynamics of murine pancreatic islet remodeling during hyperglycaemia progression. *Molecular metabolism*, 110, 102387.

Weinberg MS, et al. (2026) Bacillus Calmette-Guérin (BCG) immunotherapy reprograms CNS immunity and alters Alzheimer's biomarkers: results from two open-label clinical trials. *Communications medicine*, 6(1).