

Resource Summary Report

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Scfind

RRID:SCR_017339

Type: Tool

Proper Citation

Scfind (RRID:SCR_017339)

Resource Information

URL: <https://scfind.sanger.ac.uk/>

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Description: Software R package as search tool for single cell RNA-seq data by gene lists. Builds index from scRNA-seq datasets which organizes information in suitable and compact manner so that datasets can be very efficiently searched for either cells or cell types in which given list of genes is expressed.

Resource Type: software application, data or information resource, software resource, data processing software

Keywords: Single, cell, RNA-seq, data, gene, list, build, index, organize

Availability: Free, Available for download, Freely available

Resource Name: Scfind

Resource ID: SCR_017339

Alternate URLs: <https://genat.uk/post/scfind/>, <http://bioconductor.org/packages/scfind/>, <https://github.com/hemberg-lab/scfind>

License: GNU GPLv3

Record Creation Time: 20220129T080334+0000

Record Last Update: 20260729T044429+0000

Ratings and Alerts

No rating or validation information has been found for Scfind.

No alerts have been found for Scfind.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Townes FW, et al. (2023) Nonnegative spatial factorization applied to spatial genomics. Nature methods, 20(2), 229.