

Resource Summary Report

Generated by [RRID](#) on Aug 10, 2026

Spacemake

RRID:SCR_022207

Type: Tool

Proper Citation

Spacemake (RRID:SCR_022207)

Resource Information

URL: <https://github.com/rajewsky-lab/spacemake>

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Description: Software pipeline for processing and analysis of large scale spatial transcriptomics data. Enables reproducible data processing from raw sequencing data to automatically generated downstream analysis reports.

Synonyms: spacemake

Resource Type: data analysis software, software resource, software application, data processing software

Defining Citation: [DOI:10.1101/2021.11.07.467598](https://doi.org/10.1101/2021.11.07.467598)

Keywords: spatial dataset, gene counts, large scale spatial transcriptomics data, transcriptomics data, spatial transcriptomics data, raw sequencing data, analysis reports

Availability: Free, Available for download, Freely available

Resource Name: Spacemake

Resource ID: SCR_022207

License: GNU GPL v2

Record Creation Time: 20220427T191217+0000

Record Last Update: 20260810T040750+0000

Ratings and Alerts

No rating or validation information has been found for Spacemake.

No alerts have been found for Spacemake.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Schott M, et al. (2024) Protocol for high-resolution 3D spatial transcriptomics using Open-ST. STAR protocols, 6(1), 103521.

Legnini I, et al. (2023) Spatiotemporal, optogenetic control of gene expression in organoids. Nature methods, 20(10), 1544.