

# Resource Summary Report

Generated by [ASWG](#) on Sep 5, 2026

## Slide-seq Pipeline

RRID:SCR\_023379

Type: Tool

### Proper Citation

Slide-seq Pipeline (RRID:SCR\_023379)

### Resource Information

**URL:** [https://broadinstitute.github.io/warp/docs/Pipelines/SlideSeq\\_Pipeline/README/](https://broadinstitute.github.io/warp/docs/Pipelines/SlideSeq_Pipeline/README/)

**Proper Citation:** Slide-seq Pipeline (RRID:SCR\_023379)

**Description:** Software pipeline developed in collaboration with BRAIN Initiative Cell Census Network and BRAIN Initiative Cell Atlas Network. Supports processing of spatial transcriptomic data generated with Slide-seq commercialized as Curio Seeker assay.

**Synonyms:** Slide-seq, slideseq, SlideSeq, SlideSeq Pipeline

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

**Defining Citation:** [PMID:30923225](#)

**Keywords:** single cell data, spatial transcriptomics, barcode correction, reads alignment, aligns reads to genome,

**Funding:** Hertz Graduate Fellowship ;  
National Science Foundation Graduate Research Fellowship Program ;  
NIH Early Independence Award ;  
NIH New Innovator Award ;  
Schmidt Fellows Program at the Broad Institute

**Availability:** Free, Available for download, Freely available

**Resource Name:** Slide-seq Pipeline

**Resource ID:** SCR\_023379

**Alternate URLs:** <https://github.com/broadchenf/Slideseq/tree/master/BeadSeq%20Code>

**Old URLs:** <https://github.com/broadinstitute/warp/tree/master/pipelines/skylab/slideseq>

**Record Creation Time:** 20230321T180026+0000

**Record Last Update:** 20260903T235910+0000

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## Ratings and Alerts

No rating or validation information has been found for Slide-seq Pipeline.

No alerts have been found for Slide-seq Pipeline.

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## Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

, et al. (2026) A Community Standard Multispecies Cell Atlas of the Basal Ganglia. bioRxiv : the preprint server for biology.

Natarajan N, et al. (2024) Aberrant mitochondrial DNA synthesis in macrophages exacerbates inflammation and atherosclerosis. Nature communications, 15(1), 7337.