

Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 17, 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/

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: 20260912T200038+0000

Ratings and Alerts

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

No alerts have been found for Slide-seq Pipeline.

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 [PRECISE-TBI](#) .

, 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.