

# Resource Summary Report

Generated by T1D on Aug 9, 2026

## Paired-Tag Pipeline

RRID:SCR\_025042

Type: Tool

### Proper Citation

Paired-Tag Pipeline (RRID:SCR\_025042)

### Resource Information

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

**Proper Citation:** Paired-Tag Pipeline (RRID:SCR\_025042)

**Description:** Software open-source, cloud-optimized pipeline developed in collaboration with the BRAIN Initiative Cell Census Network (BICCN) and the BRAIN Initiative Cell Atlas Network (BICAN). It supports the processing of 3' single-nucleus histone modification data (generated with the paired-tag protocol)] and 10x gene expression (GEX) data generated with the 10x Chromium Multiome assay.

**Synonyms:** Paired-Tag Pipeline, paired\_tag, Paired-Tag

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

**Keywords:** data processing, 10x Genomics Multiome assay data, single-cell data, single nucleus gene expression data, single-nucleus histone modification data,

**Availability:** Free, Freely available

**Resource Name:** Paired-Tag Pipeline

**Resource ID:** SCR\_025042

**Old URLs:** [https://github.com/broadinstitute/warp/tree/master/pipelines/skylab/paired\\_tag](https://github.com/broadinstitute/warp/tree/master/pipelines/skylab/paired_tag)

**Record Creation Time:** 20240305T200904+0000

**Record Last Update:** 20260808T190718+0000

### Ratings and Alerts

No rating or validation information has been found for Paired-Tag Pipeline.

No alerts have been found for Paired-Tag Pipeline.

## Data and Source Information

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

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

**Listed below are recent publications.** The full list is available at [T1D](#) .

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