

Resource Summary Report

Generated by [Kravitz](#) on Sep 12, 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

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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: data processing software, software application, software resource

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

Record Creation Time: 20240305T200904+0000

Record Last Update: 20260912T200436+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](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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