# **Resource Summary Report**

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# **ATAC Pipeline**

RRID:SCR\_024656 Type: Tool

#### **Proper Citation**

ATAC Pipeline (RRID:SCR\_024656)

### **Resource Information**

URL: https://github.com/broadinstitute/warp/blob/master/pipelines/skylab/multiome/atac.wdl

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**Description:** Software pipeline developed in collaboration with the BRAIN Initiative Cell Census Network, BRAIN Initiative Cell Atlas Network, and SCORCH. Supports processing of 10x single-nucleus data generated with 10x Multiome ATAC-seq (Assay for Transposase-Accessible Chromatin) assay.

Synonyms: ATAC, atac, atac pipeline

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

**Keywords:** data processing, 10x Genomics Multiome ATAC-seq assay data, single-cell data, single-cell chromatin accessibility assay data

#### Funding:

Availability: Free, Available for download, Freely available

Resource Name: ATAC Pipeline

Resource ID: SCR\_024656

Alternate URLs: https://broadinstitute.github.io/warp/docs/Pipelines/ATAC/README

Record Creation Time: 20231027T050251+0000

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## **Ratings and Alerts**

No rating or validation information has been found for ATAC Pipeline.

No alerts have been found for ATAC Pipeline.

Data and Source Information

Source: <u>SciCrunch Registry</u>

**Usage and Citation Metrics** 

We have not found any literature mentions for this resource.