

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://broadinstitute.github.io/warp/docs/Pipelines/ATAC/README>

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Description: Software open-source, cloud-optimized pipeline developed in collaboration with members of the BRAIN Initiative (BICCN and BICAN Sequencing Working Group) and SCORCH. It supports the processing of 10x single-nucleus data generated with 10x Multiome ATAC-seq (Assay for Transposase-Accessible Chromatin), a technique used in molecular biology to assess genome-wide chromatin accessibility.

Synonyms: ATAC, atac, atac pipeline

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

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

Availability: Free, Available for download, Freely available

Resource Name: ATAC Pipeline

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Old URLs:

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

Record Creation Time: 20231027T050251+0000

Record Last Update: 20260820T055024+0000

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: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.