## **Resource Summary Report**

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# snM3C Pipeline

RRID:SCR\_025041

Type: Tool

### **Proper Citation**

snM3C Pipeline (RRID:SCR\_025041)

#### **Resource Information**

URL: https://github.com/broadinstitute/warp/tree/master/pipelines/skylab/snM3C

**Proper Citation:** snM3C Pipeline (RRID:SCR\_025041)

**Description:** Software pipeline developed in collaboration with the BRAIN Initiative Cell Census Network and BRAIN Initiative Cell Atlas Network. Supports processing of single-nucleus methylome and chromatin contact (snM3C) sequencing data.

Abbreviations: snM3C

**Synonyms:** snM3C, Single Nucleus Methly-Seq and Chromatin Capture Pipeline, single-nucleus Methylome and Chromatin Contact

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

**Keywords:** data processing, single-nucleus methylome data, single-cell data, chromatin contact data, single-cell methylome and chromatin contact sequencing data,

**Funding:** 

Availability: Free, Freely available

Resource Name: snM3C Pipeline

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Alternate URLs: https://broadinstitute.github.io/warp/docs/Pipelines/snM3C/README

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## **Ratings and Alerts**

No rating or validation information has been found for snM3C Pipeline.

No alerts have been found for snM3C Pipeline.

### **Data and Source Information**

Source: SciCrunch Registry

# **Usage and Citation Metrics**

We have not found any literature mentions for this resource.