

Resource Summary Report

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Taiji

RRID:SCR_028464

Type: Tool

Proper Citation

Taiji (RRID:SCR_028464)

Resource Information

URL: <https://github.com/Taiji-pipeline/Taiji>

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Description: Software integrative analysis pipeline for analyzing bulk/single-cell ATAC-seq and RNA-seq data. Integrative multi-omics data analysis framework. It can be used as a standalone pipeline to analyze ATAC-seq, RNA-seq, single cell ATAC-seq or Drop-seq data. Used to integrate diverse datasets and use these information to construct regulatory network and identify candidate driver genes.

Resource Type: software toolkit, software resource, source code

Defining Citation: [PMID:30944857](#)

Keywords: integrative analysis, single-cell ATAC-seq data, RNA-seq data, identify driver transcription factors, construct regulatory network, identify candidate driver genes,

Funding: NHGRI U54HG006997;
NHGRI R01HG009626

Availability: Free, Available for download, Freely available

Resource Name: Taiji

Resource ID: SCR_028464

Alternate URLs: <https://taiji-pipeline.github.io/>

License: BSD-3-Clause

Record Creation Time: 20260519T054547+0000

Record Last Update: 20260804T043942+0000

Ratings and Alerts

No rating or validation information has been found for Taiji.

No alerts have been found for Taiji.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.