

# Resource Summary Report

Generated by [RRID](#) on Sep 10, 2026

## scrattch taxonomy

RRID:SCR\_027910

Type: Tool

### Proper Citation

scrattch taxonomy (RRID:SCR\_027910)

### Resource Information

**URL:** <https://alleninstitute.github.io/scrattch.taxonomy/>

**Proper Citation:** scrattch taxonomy (RRID:SCR\_027910)

**Description:** Software R package from the Allen Institute designed to build, standardize, and analyze single-cell RNA-seq-based cell type taxonomies. It utilizes a structured Allen Institute schema (AIT) to organize cell annotations and metadata, enabling hierarchical, data-driven cell type classification

**Resource Type:** software resource, software toolkit, source code

**Keywords:** building cell type taxonomies, single cell RNA-seq data, Allen Institute (AIT) schema,

**Availability:** Free, Available for download, Freely available

**Resource Name:** scrattch taxonomy

**Resource ID:** SCR\_027910

**Alternate URLs:** <https://github.com/AllenInstitute/scrattch.taxonomy>

**Record Creation Time:** 20260131T071535+0000

**Record Last Update:** 20260905T133602+0000

### Ratings and Alerts

No rating or validation information has been found for scrattch taxonomy.

No alerts have been found for scrattch taxonomy.

### Data and Source Information

**Source:** [SciCrunch Registry](#)

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## Usage and Citation Metrics

We found 1 mentions in open access literature.

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

Ray PL, et al. (2026) A layered standards framework for integrating single-cell and spatial omics data into brain cell atlases. bioRxiv : the preprint server for biology.