

Resource Summary Report

Generated by [NIF](#) on Sep 2, 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: 20260829T183533+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](#)

Usage and Citation Metrics

We found 1 mentions in open access literature.

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

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.