

Resource Summary Report

Generated by RRID on Sep 10, 2026

scTHI

RRID:SCR_018918

Type: Tool

Proper Citation

scTHI (RRID:SCR_018918)

Resource Information

URL: <https://bioconductor.org/packages/scTHI/>

Proper Citation: scTHI (RRID:SCR_018918)

Description: Software R package to identify active pairs of ligand receptors from single cells in order to study,among others, tumor host interactions. Contains set of signatures to classify cells from tumor microenvironment.

Abbreviations: scTHI

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

Keywords: Ligand receptor, active pair, active pair ligand receptor, single cell, tumor interaction, tumor microenvironment, cells clasification

Availability: Free, Available for download, Freely available

Resource Name: scTHI

Resource ID: SCR_018918

License: GPL-2

Record Creation Time: 20220129T080342+0000

Record Last Update: 20260905T132844+0000

Ratings and Alerts

No rating or validation information has been found for scTHI.

No alerts have been found for scTHI.

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 [RRID](#) .

Caruso FP, et al. (2020) A map of tumor-host interactions in glioma at single-cell resolution. GigaScience, 9(10).