

Resource Summary Report

Generated by RRID on Sep 10, 2026

Parsnip

RRID:SCR_024744

Type: Tool

Proper Citation

Parsnip (RRID:SCR_024744)

Resource Information

URL: <https://www.tidyverse.org/blog/2018/11/parsnip-0-0-1/>

Proper Citation: Parsnip (RRID:SCR_024744)

Description: Software R package to create and manipulate functions commonly used during modeling e.g. fitting the model, making predictions, etc.

Resource Type: software resource, software toolkit

Keywords: create and manipulate functions, modeling functions, fitting the model, making predictions,

Availability: Free, Available for download, Freely available

Resource Name: Parsnip

Resource ID: SCR_024744

Alternate URLs: <https://CRAN.R-project.org/package=parsnip>,
<https://github.com/tidymodels/parsnip>

License: MIT license

Record Creation Time: 20231204T190629+0000

Record Last Update: 20260905T133440+0000

Ratings and Alerts

No rating or validation information has been found for Parsnip.

No alerts have been found for Parsnip.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 3 mentions in open access literature.

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

Ikegami M, et al. (2026) Germline Pathogenic Variant Prediction Model for Tumor-Only Sequencing Based on Japanese Clinicogenomic Database. Clinical cancer research : an official journal of the American Association for Cancer Research, 32(4), 770.

Hoekstra ME, et al. (2024) Distinct spatiotemporal dynamics of CD8+ T??cell-derived cytokines in the tumor microenvironment. Cancer cell, 42(1), 157.

Jessa S, et al. (2024) FOXR2 targets LHX6+/DLX+ neural lineages to drive CNS neuroblastoma. Cancer research.