

# Resource Summary Report

Generated by [dkNET](#) on Sep 11, 2026

## Parsnip

RRID:SCR\_024744

Type: Tool

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### Proper Citation

Parsnip (RRID:SCR\_024744)

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### 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

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### Ratings and Alerts

No rating or validation information has been found for Parsnip.

No alerts have been found for Parsnip.

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### Data and Source Information

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

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## Usage and Citation Metrics

We found 3 mentions in open access literature.

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

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.