

# Resource Summary Report

Generated by [PRECISE-TBI](#) on Sep 8, 2026

## R package inTrees

RRID:SCR\_017299

Type: Tool

---

### Proper Citation

R package inTrees (RRID:SCR\_017299)

---

### Resource Information

**URL:** <https://CRAN.R-project.org/package=inTrees>

**Proper Citation:** R package inTrees (RRID:SCR\_017299)

**Description:** Software R package for tree ensembles such as random forests, regularized random forests and gradient boosted trees. Provides functions for extracting, measuring and pruning rules, selecting compact rule set, summarizing rules into learner, calculating frequent variable interactions, formatting rules in latex code.

**Synonyms:** inTrees

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

**Keywords:** tree, ensemble, random, forest, gradient, extracting, function, measuring, pruning, formatting, variable, interaction, latex, code

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

**Resource Name:** R package inTrees

**Resource ID:** SCR\_017299

**Alternate URLs:** <https://github.com/softwareeng/inTrees/issues>

**License:** GPL

**Record Creation Time:** 20220129T080334+0000

**Record Last Update:** 20260905T133302+0000

---

### Ratings and Alerts

No rating or validation information has been found for R package inTrees.

No alerts have been found for R package inTrees.

---

## Data and Source Information

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

---

## Usage and Citation Metrics

We found 2 mentions in open access literature.

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

Sariya S, et al. (2021) GARCOM: A user-friendly R package for genetic mutation counts. F1000Research, 10, 524.

Radebe L, et al. (2021) Predicting Malignancy in Pediatric Thyroid Nodules: Early Experience With Machine Learning for Clinical Decision Support. The Journal of clinical endocrinology and metabolism, 106(12), e5236.