

Resource Summary Report

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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>

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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: 20260821T194120+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 [ASWG](#) .

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.