

Package ‘dynfeature’

October 13, 2022

Type Package

Title Feature Importance for Dynamic Processes

Version 1.0.0

Description Calculating feature importance scores from trajectories using the random forests algorithm and more. Saelens and Cannoodt et al. (2019) <[doi:10.1038/s41587-019-0071-9](https://doi.org/10.1038/s41587-019-0071-9)>.

License GPL-3

Encoding UTF-8

Imports dplyr, dynutils (>= 1.0.2), dynwrap (>= 1.0.0), purrr, magrittr, methods, ranger, reshape2, testthat, tidyr, tibble

Suggests caret, covr

RoxygenNote 7.1.1

NeedsCompilation no

Author Robrecht Cannoodt [aut, cre] (<<https://orcid.org/0000-0003-3641-729X>>, rcannood),
Wouter Saelens [aut] (<<https://orcid.org/0000-0002-7114-6248>>, zouter)

Maintainer Robrecht Cannoodt <rcannood@gmail.com>

Repository CRAN

Date/Publication 2021-06-14 07:30:12 UTC

R topics documented:

calculate_branch_feature_importance	2
dynfeature	3
fi_ranger_rf_lite	4

Index	6
--------------	----------

`calculate_branch_feature_importance`*Calculating feature importances across trajectories*

Description

Uses the feature importance measures of [ranger](#) or [caret](#). `calculate_overall_feature_importance` calculates the importance for the whole trajectory, `calculate_milestone_feature_importance` calculates it for individual milestones (e.g. branching points)

Usage

```
calculate_branch_feature_importance(  
  trajectory,  
  expression_source = "expression",  
  fi_method = fi_ranger_rf_lite(),  
  verbose = FALSE  
)  
  
calculate_branching_point_feature_importance(  
  trajectory,  
  expression_source = "expression",  
  milestones_oi = trajectory$milestone_ids,  
  fi_method = fi_ranger_rf_lite(),  
  verbose = FALSE  
)  
  
calculate_cell_feature_importance(  
  trajectory,  
  expression_source = "expression",  
  fi_method = fi_ranger_rf_lite(),  
  verbose = FALSE  
)  
  
calculate_milestone_feature_importance(  
  trajectory,  
  expression_source = "expression",  
  milestones_oi = NULL,  
  fi_method = fi_ranger_rf_lite(),  
  verbose = FALSE  
)  
  
calculate_overall_feature_importance(  
  trajectory,  
  expression_source = "expression",  
  fi_method = fi_ranger_rf_lite(),  
  verbose = FALSE  
)
```

```

)

calculate_waypoint_feature_importance(
  trajectory,
  expression_source = "expression",
  waypoints = NULL,
  fi_method = fi_ranger_rf_lite(),
  verbose = FALSE
)

```

Arguments

trajectory	A trajectory object containing expression values and a trajectory.
expression_source	The expression data matrix, with features as columns. <ul style="list-style-type: none"> • If a matrix is provided, it is used as is. • If a character is provided, trajectory[[expression_source]] should contain the matrix. • If a function is provided, that function will be called in order to obtain the expression (useful for lazy loading).
fi_method	A feature importance method. Default: fi_ranger_rf_lite(). Check ?fi_methods for a full list of available feature importance methods.
verbose	Whether to print out extra information.
milestones_oi	The milestone(s) for which to calculate feature importance
waypoints	The waypoints, optional

Value

A data frame with two or more columns, feature_id, and importance. feature_id is a column in the trajectory expression matrix. Additional columns may be available depending on the function called.

Examples

```

library(dynwrap)
data(example_trajectory)

calculate_overall_feature_importance(example_trajectory)

```

dynfeature

Feature Importance for Dynamic Processes

Description

Calculating feature importance scores from trajectories using the random forests algorithm.

fi_ranger_rf_lite *Feature Importance methods*

Description

Feature Importance methods

Usage

```
fi_ranger_rf_lite(
  num_trees = 2000,
  num_variables_per_split = 50,
  num_samples_per_tree = 250,
  min_node_size = 20,
  ...
)

fi_ranger_rf(...)

fi_caret(caret_method, ...)

fi_ranger_rf_tiny(
  num_trees = 100,
  num_variables_per_split = 50,
  num_samples_per_tree = 250,
  min_node_size = 20,
  ...
)
```

Arguments

num_trees	(fi_ranger_rf_lite) The number of trees to use
num_variables_per_split	(fi_ranger_rf_lite) The number of variables to sample per split
num_samples_per_tree	(fi_ranger_rf_lite) The number of samples to bootstrap per split
min_node_size	(fi_ranger_rf_lite) The minimum node size, no split will be made if the node size is less than this value.
...	Extra parameters to pass onto the underlying feature importance function.
caret_method	(fi_caret) Which caret method to use for feature importance.

Value

A list containing a helper function for calling a feature importance function.

Examples

```
library(dynwrap)
data(example_trajectory)

calculate_overall_feature_importance(example_trajectory, fi_method = fi_ranger_rf())
```

Index

`calculate_branch_feature_importance`, [2](#)
`calculate_branching_point_feature_importance`
 (`calculate_branch_feature_importance`),
 [2](#)
`calculate_cell_feature_importance`
 (`calculate_branch_feature_importance`),
 [2](#)
`calculate_milestone_feature_importance`
 (`calculate_branch_feature_importance`),
 [2](#)
`calculate_overall_feature_importance`
 (`calculate_branch_feature_importance`),
 [2](#)
`calculate_waypoint_feature_importance`
 (`calculate_branch_feature_importance`),
 [2](#)

`dynfeature`, [3](#)

`fi_caret` (`fi_ranger_rf_lite`), [4](#)
`fi_ranger_rf` (`fi_ranger_rf_lite`), [4](#)
`fi_ranger_rf_lite`, [4](#)
`fi_ranger_rf_tiny` (`fi_ranger_rf_lite`), [4](#)

`ranger`, [2](#)