

# Package ‘dynfeature’

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**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, tidyverse, tibble

**Suggests** caret, covr

**RoxxygenNote** 7.1.1

**NeedsCompilation** no

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**calculate\_branch\_feature\_importance**  
*Calculating feature importances across trajectories*

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## 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"><li>• If a matrix is provided, it is used as is.</li><li>• If a character is provided, <code>trajectory[[expression_source]]</code> should contain the matrix.</li><li>• If a function is provided, that function will be called in order to obtain the expression (useful for lazy loading).</li></ul>
fi_method	A feature importance method. Default: <code>fi_ranger_rf_lite()</code> . Check <code>?fi_methods</code> 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)
```

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## Description

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

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 fi\_ranger\_rf\_lite      *Feature Importance methods*


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## 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	( <i>fi_ranger_rf_lite</i> ) The number of trees to use
num_variables_per_split	( <i>fi_ranger_rf_lite</i> ) The number of variables to sample per split
num_samples_per_tree	( <i>fi_ranger_rf_lite</i> ) The number of samples to bootstrap per split
min_node_size	( <i>fi_ranger_rf_lite</i> ) 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	( <i>fi_caret</i> ) 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())
```

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