

## sklearn.ensemble.RandomForestRegressor

```
class sklearn.ensemble.RandomForestRegressor(n_estimators=100, *, criterion='squared_error', max_depth=None, min_samples_split=2, min_samples_leaf=1, min_weight_fraction_leaf=0.0, max_features='auto', max_leaf_nodes=None, min_impurity_decrease=0.0, bootstrap=True, oob_score=False, n_jobs=None, random_state=None, verbose=0, warm_start=False, ccp_alpha=0.0, max_samples=None)
```

A random forest regressor.

A random forest is a meta estimator that fits a number of classifying decision trees on various sub-samples of the dataset and uses averaging to improve the predictive accuracy and control over-fitting. The sub-sample size is controlled with the `max_samples` parameter if `bootstrap=True` (default), otherwise the whole dataset is used to build each tree.

### Parameters:

**n\_estimators : int, default=100**

The number of trees in the forest.

*Changed in version 0.22:* The default value of `n_estimators` changed from 10 to 100 in 0.22.

**criterion : {"squared\_error", "absolute\_error", "poisson"}, default="squared\_error"**

The function to measure the quality of a split. Supported criteria are "squared\_error" for the mean squared error, which is equal to variance reduction as feature selection criterion, "absolute\_error" for the mean absolute error, and "poisson" which uses reduction in Poisson deviance to find splits. Training using "absolute\_error" is significantly slower than when using "squared\_error".

*New in version 0.18:* Mean Absolute Error (MAE) criterion.

*New in version 1.0:* Poisson criterion.

*Deprecated since version 1.0:* Criterion "mse" was deprecated in v1.0 and will be removed in version 1.2. Use `criterion="squared_error"` which is equivalent.

*Deprecated since version 1.0:* Criterion "mae" was deprecated in v1.0 and will be removed in version 1.2. Use `criterion="absolute_error"` which is equivalent.

**max\_depth : int, default=None**

The maximum depth of the tree. If None, then nodes are expanded until all leaves are pure or until all leaves contain less than `min_samples_split` samples.

**min\_samples\_split : int or float, default=2**

The minimum number of samples required to split an internal node:

- If int, then consider `min_samples_split` as the minimum number.
- If float, then `min_samples_split` is a fraction and `ceil(min_samples_split * n_samples)` are the minimum number of samples for each split.

*Changed in version 0.18:* Added float values for fractions.

**min\_samples\_leaf : int or float, default=1**

The minimum number of samples required to be at a leaf node. A split point at any depth will only be considered if it leaves at least `min_samples_leaf` training samples in each of the left and right branches. This may have the effect of smoothing the model, especially in regression.

- If int, then consider `min_samples_leaf` as the minimum number.
- If float, then `min_samples_leaf` is a fraction and `ceil(min_samples_leaf * n_samples)` are the minimum number of samples for each node.

*Changed in version 0.18: Added float values for fractions.*

**min\_weight\_fraction\_leaf : float, default=0.0**

The minimum weighted fraction of the sum total of weights (of all the input samples) required to be at a leaf node. Samples have equal weight when `sample_weight` is not provided.

**max\_features : {"auto", "sqrt", "log2"}, int or float, default="auto"**

The number of features to consider when looking for the best split:

- If int, then consider `max_features` features at each split.
- If float, then `max_features` is a fraction and `round(max_features * n_features)` features are considered at each split.
- If "auto", then `max_features=n_features`.
- If "sqrt", then `max_features=sqrt(n_features)`.
- If "log2", then `max_features=log2(n_features)`.
- If None, then `max_features=n_features`.

Note: the search for a split does not stop until at least one valid partition of the node samples is found, even if it requires to effectively inspect more than `max_features` features.

**max\_leaf\_nodes : int, default=None**

Grow trees with `max_leaf_nodes` in best-first fashion. Best nodes are defined as relative reduction in impurity. If None then unlimited number of leaf nodes.

**min\_impurity\_decrease : float, default=0.0**

A node will be split if this split induces a decrease of the impurity greater than or equal to this value.

The weighted impurity decrease equation is the following:

$$N_t / N * (impurity - N_{t_R} / N_t * right\_impurity - N_{t_L} / N_t * left\_impurity)$$

where `N` is the total number of samples, `Nt` is the number of samples at the current node, `Nt_L` is the number of samples in the left child, and `Nt_R` is the number of samples in the right child.

`N`, `Nt`, `Nt_R` and `Nt_L` all refer to the weighted sum, if `sample_weight` is passed.

**bootstrap : bool, default=True**

Whether bootstrap samples are used when building trees. If False, the whole dataset is used to build each tree.

**oob\_score : bool, default=False**

Whether to use out-of-bag samples to estimate the generalization score. Only available if bootstrap=True.

**n\_jobs : int, default=None**

The number of jobs to run in parallel. `fit`, `predict`, `decision_path` and `apply` are all parallelized over the trees. `None` means 1 unless in a `joblib.parallel_backend` context. `-1` means using all processors. See [Glossary](#) for more details.

**random\_state : int, RandomState instance or None, default=None**

Controls both the randomness of the bootstrapping of the samples used when building trees (if `bootstrap=True`) and the sampling of the features to consider when looking for the best split at each node (if `max_features < n_features`). See [Glossary](#) for details.

**verbose : int, default=0**

Controls the verbosity when fitting and predicting.

**warm\_start : bool, default=False**

When set to `True`, reuse the solution of the previous call to fit and add more estimators to the ensemble, otherwise, just fit a whole new forest. See [the Glossary](#).

**ccp\_alpha : non-negative float, default=0.0**

Complexity parameter used for Minimal Cost-Complexity Pruning. The subtree with the largest cost complexity that is smaller than `ccp_alpha` will be chosen. By default, no pruning is performed. See [Minimal Cost-Complexity Pruning](#) for details.

*New in version 0.22.*

**max\_samples : int or float, default=None**

If bootstrap is True, the number of samples to draw from X to train each base estimator.

- If None (default), then draw `X.shape[0]` samples.
- If int, then draw `max_samples` samples.
- If float, then draw `max_samples * X.shape[0]` samples. Thus, `max_samples` should be in the interval `(0.0, 1.0]`.

*New in version 0.22.*

## Methods

|   |  |
|---|--|
| <code>apply(X)</code>                     | Apply trees in the forest to X, return leaf indices.       |
| <code>decision_path(X)</code>             | Return the decision path in the forest.                    |
| <code>fit(X, y[, sample_weight])</code>   | Build a forest of trees from the training set (X, y).      |
| <code>get_params([deep])</code>           | Get parameters for this estimator.                         |
| <code>predict(X)</code>                   | Predict regression target for X.                           |
| <code>score(X, y[, sample_weight])</code> | Return the coefficient of determination of the prediction. |
| <code>set_params(**params)</code>         | Set the parameters of this estimator.                      |

**Attributes:****base\_estimator\_ : *DecisionTreeRegressor***

The child estimator template used to create the collection of fitted sub-estimators.

**estimators\_ : *list of DecisionTreeRegressor***

The collection of fitted sub-estimators.

**feature\_importances\_ : *ndarray of shape (n\_features,)***

The impurity-based feature importances.

**n\_features\_ : *int***

DEPRECATED: Attribute `n_features_` was deprecated in version 1.0 and will be removed in 1.2.

**n\_features\_in\_ : *int***

Number of features seen during `fit`.

*New in version 0.24.*

**feature\_names\_in\_ : *ndarray of shape (n\_features\_in\_,)***

Names of features seen during `fit`. Defined only when `x` has feature names that are all strings.

*New in version 1.0.*

**n\_outputs\_ : *int***

The number of outputs when `fit` is performed.

**oob\_score\_ : *float***

Score of the training dataset obtained using an out-of-bag estimate. This attribute exists only when `oob_score` is `True`.

**oob\_prediction\_ : *ndarray of shape (n\_samples,) or (n\_samples, n\_outputs)***

Prediction computed with out-of-bag estimate on the training set. This attribute exists only when `oob_score` is `True`.