sklearn.model_selection.GridSearchCV

class sklearn.model_selection.GridSearchCV(estimator, param_grid, *, scoring=None, n_jobs=None, refit=True, cv=None,
verbose=0, pre_dispatch='2*n_jobs', error_score=nan, return_train_score=False)
[source]

Exhaustive search over specified parameter values for an estimator.

Important members are fit, predict.

GridSearchCV implements a "fit" and a "score" method. It also implements "score_samples", "predict", "predict_proba", "decision_function", "transform" and "inverse_transform" if they are implemented in the estimator used.

The parameters of the estimator used to apply these methods are optimized by cross-validated grid-search over a parameter grid.

Read more in the **User Guide**.

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Parameters:

estimator: estimator object

This is assumed to implement the scikit-learn estimator interface. Either estimator needs to provide a score function, or scoring must be passed.

param_grid: dict or list of dictionaries

Dictionary with parameters names (str) as keys and lists of parameter settings to try as values, or a list of such dictionaries, in which case the grids spanned by each dictionary in the list are explored. This enables searching over any sequence of parameter settings.

scoring: str, callable, list, tuple or dict, default=None

Strategy to evaluate the performance of the cross-validated model on the test set.

If scoring represents a single score, one can use:

- a single string (see The scoring parameter: defining model evaluation rules);
- a callable (see <u>Defining your scoring strategy from metric functions</u>) that returns a single value.

If scoring represents multiple scores, one can use:

- a list or tuple of unique strings;
- a callable returning a dictionary where the keys are the metric names and the values are the metric scores;
- a dictionary with metric names as keys and callables a values.

See Specifying multiple metrics for evaluation for an example.

n_jobs : int, default=None

Number of jobs to run in parallel. None means 1 unless in a joblib.parallel backend context. -1 means using all processors. See Glossary for more details.

Changed in version v0.20: n_jobs default changed from 1 to None

refit: bool, str, or callable, default=True

Refit an estimator using the best found parameters on the whole dataset.

For multiple metric evaluation, this needs to be a str denoting the scorer that would be used to find the best parameters for refitting the estimator at the end.

Where there are considerations other than maximum score in choosing a best estimator, refit can be set to a function which returns the selected best_index_ given cv_results_. In that case, the best_estimator_ and best_params_ will be set according to the returned best_index_ while the best_score_ attribute will not be available.

The refitted estimator is made available at the best_estimator_ attribute and permits using predict directly on this GridSearchCV instance.

Also for multiple metric evaluation, the attributes best_index_, best_score_ and best_params_ will only be available if refit is set and all of them will be determined w.r.t this specific scorer.

See scoring parameter to know more about multiple metric evaluation.

See <u>Custom refit strategy of a grid search with cross-validation</u> to see how to design a custom selection strategy using a callable via refit.

Changed in version 0.20: Support for callable added.

cv: int, cross-validation generator or an iterable, default=None

Determines the cross-validation splitting strategy. Possible inputs for cv are:

- None, to use the default 5-fold cross validation,
- integer, to specify the number of folds in a (Stratified)KFold,
- CV splitter,
- An iterable yielding (train, test) splits as arrays of indices.

For integer/None inputs, if the estimator is a classifier and y is either binary or multiclass, **StratifiedKFold** is used. In all other cases, **KFold** is used. These splitters are instantiated with **shuffle=False** so the splits will be the same across calls.

Refer <u>User Guide</u> for the various cross-validation strategies that can be used here.

Toggle Menu I in version 0.22: cv default value if None changed from 3-fold to 5-fold.

verbose: int

Controls the verbosity: the higher, the more messages.

- >1 : the computation time for each fold and parameter candidate is displayed;
- >2 : the score is also displayed;
- >3: the fold and candidate parameter indexes are also displayed together with the starting time of the computation.

pre_dispatch : int, or str, default='2*n_jobs'

Controls the number of jobs that get dispatched during parallel execution. Reducing this number can be useful to avoid an explosion of memory consumption when more jobs get dispatched than CPUs can process. This parameter can be:

- None, in which case all the jobs are immediately created and spawned. Use this for lightweight and fast-running jobs, to avoid delays due to on-demand spawning of the jobs
- An int, giving the exact number of total jobs that are spawned
- A str, giving an expression as a function of n_jobs, as in '2*n_jobs'

error_score : 'raise' or numeric, default=np.nan

Value to assign to the score if an error occurs in estimator fitting. If set to 'raise', the error is raised. If a numeric value is given, FitFailedWarning is raised. This parameter does not affect the refit step, which will always raise the error.

return_train_score : bool, default=False

If False, the cv_results_ attribute will not include training scores. Computing training scores is used to get insights on how different parameter settings impact the overfitting/underfitting trade-off. However computing the scores on the training set can be computationally expensive and is not strictly required to select the parameters that yield the best generalization performance.

New in version 0.19.

Changed in version 0.21: Default value was changed from True to False

Attributes:

cv_results_: dict of numpy (masked) ndarrays

A dict with keys as column headers and values as columns, that can be imported into a pandas DataFrame.

For instance the below given table

param_kernel	param_gamma	param_degree	split0_test_score	 rank_t
'poly'	_	2	0.80	 2
'poly'	_	3	0.70	 4
'rbf'	0.1	_	0.80	 3
ʻrbf'	0.2	_	0.93	 1

will be represented by a cv_results_ dict of:

```
{
'param_kernel': masked_array(data = ['poly', 'poly', 'rbf', 'rbf'],
                            mask = [False False False False]...)
'param_gamma': masked_array(data = [-- -- 0.1 0.2],
                           mask = [ True True False False]...),
'param_degree': masked_array(data = [2.0 3.0 -- --],
                            mask = [False False True True]...),
'split0_test_score' : [0.80, 0.70, 0.80, 0.93],
'split1_test_score' : [0.82, 0.50, 0.70, 0.78],
'mean_test_score' : [0.81, 0.60, 0.75, 0.85],
'std_test_score'
                   : [0.01, 0.10, 0.05, 0.08],
'rank_test_score'
                    : [2, 4, 3, 1],
'split0_train_score': [0.80, 0.92, 0.70, 0.93],
'split1_train_score' : [0.82, 0.55, 0.70, 0.87],
'mean_train_score' : [0.81, 0.74, 0.70, 0.90],
                    : [0.01, 0.19, 0.00, 0.03],
'std_train_score'
'mean_fit_time'
                    : [0.73, 0.63, 0.43, 0.49],
'std_fit_time'
                    : [0.01, 0.02, 0.01, 0.01],
'mean_score_time'
                    : [0.01, 0.06, 0.04, 0.04],
'std_score_time'
                    : [0.00, 0.00, 0.00, 0.01],
                    : [{'kernel': 'poly', 'degree': 2}, ...],
'params'
}
```

NOTE

Toggle Menu params' is used to store a list of parameter settings dicts for all the parameter candidates.

The mean_fit_time, std_fit_time, mean_score_time and std_score_time are all in seconds.

For multi-metric evaluation, the scores for all the scorers are available in the cv_results_ dict at the keys ending with that scorer's name ('_<scorer_name>') instead of '_score' shown above. ('split0_test_precision', 'mean_train_precision' etc.)

best_estimator_: estimator

Estimator that was chosen by the search, i.e. estimator which gave highest score (or smallest loss if specified) on the left out data. Not available if refit=False.

See refit parameter for more information on allowed values.

best_score_ : float

Mean cross-validated score of the best estimator

For multi-metric evaluation, this is present only if refit is specified.

This attribute is not available if refit is a function.

best_params_: dict

Parameter setting that gave the best results on the hold out data.

For multi-metric evaluation, this is present only if refit is specified.

best_index_: int

The index (of the cv_results_ arrays) which corresponds to the best candidate parameter setting.

The dict at search.cv_results_['params'][search.best_index_] gives the parameter setting for the best model, that gives the highest mean score (search.best_score_).

For multi-metric evaluation, this is present only if refit is specified.

scorer_: function or a dict

Scorer function used on the held out data to choose the best parameters for the model.

For multi-metric evaluation, this attribute holds the validated scoring dict which maps the scorer key to the scorer callable.

n_splits_: int

The number of cross-validation splits (folds/iterations).

refit_time_ : float

Seconds used for refitting the best model on the whole dataset.

This is present only if refit is not False.

New in version 0.20.

multimetric_: bool

Whether or not the scorers compute several metrics.

classes_ : ndarray of shape (n_classes,)

Class labels.

n_features_in_: int

Number of features seen during fit.

feature_names_in_: ndarray of shape (n_features_in_,)

Names of features seen during <u>fit</u>. Only defined if <u>best_estimator_</u> is defined (see the documentation for the <u>refit</u> parameter for more details) and that <u>best_estimator_</u> exposes <u>feature_names_in_</u> when fit.

New in version 1.0.

See also:

ParameterGrid

Generates all the combinations of a hyperparameter grid.

train_test_split

Utility function to split the data into a development set usable for fitting a GridSearchCV instance and an evaluation set for its final evaluation.

Toggle Menu rics.make scorer

Make a scorer from a performance metric or loss function.

Notes

The parameters selected are those that maximize the score of the left out data, unless an explicit score is passed in which case it is used instead.

If n_jobs was set to a value higher than one, the data is copied for each point in the grid (and not n_jobs times). This is done for efficiency reasons if individual jobs take very little time, but may raise errors if the dataset is large and not enough memory is available. A workaround in this case is to set pre_dispatch. Then, the memory is copied only pre_dispatch many times. A reasonable value for pre_dispatch is 2 * n_jobs.

Examples

Methods

<pre>decision_function(X)</pre>	Call decision_function on the estimator with the best found parameters.
<pre>fit(X[, y])</pre>	Run fit with all sets of parameters.
<pre>get metadata routing()</pre>	Get metadata routing of this object.
<pre>get params([deep])</pre>	Get parameters for this estimator.
<pre>inverse transform(Xt)</pre>	Call inverse_transform on the estimator with the best found params.
<pre>predict(X)</pre>	Call predict on the estimator with the best found parameters.
<pre>predict log proba(X)</pre>	Call predict_log_proba on the estimator with the best found parameters.
<pre>predict_proba(X)</pre>	Call predict_proba on the estimator with the best found parameters.
<pre>score(X[, y])</pre>	Return the score on the given data, if the estimator has been refit.
<pre>score samples(X)</pre>	Call score_samples on the estimator with the best found parameters.
<pre>set params(**params)</pre>	Set the parameters of this estimator.
transform(X)	Call transform on the estimator with the best found parameters.

property classes_

Class labels.

Only available when refit=True and the estimator is a classifier.

decision_function(X)

Call decision_function on the estimator with the best found parameters.

Only available if refit=True and the underlying estimator supports decision_function.

Parameters:

X: indexable, length n_samples

Must fulfill the input assumptions of the underlying estimator.

Returns:

y_score : $ndarray of shape (n_samples,) or (n_samples, n_classes) or (n_samples, n_classes * (n_classes-1) / 2)$ Result of the decision function for \times based on the estimator with the best found parameters.

fit(X, y=None, **params) [source]

Toggle Menu all sets of parameters.

Parameters:

X : array-like of shape (n_samples, n_features)

Training vector, where n_samples is the number of samples and n_features is the number of features.

y: array-like of shape (n_samples, n_output) or (n_samples,), default=None

Target relative to X for classification or regression; None for unsupervised learning.

**params : dict of str -> object

Parameters passed to the fit method of the estimator, the scorer, and the CV splitter.

If a fit parameter is an array-like whose length is equal to $num_samples$ then it will be split across CV groups along with x and y. For example, the <u>sample_weight</u> parameter is split because $len(sample_weights) = len(X)$.

Returns:

self: object

Instance of fitted estimator.

get_metadata_routing()

[source]

Get metadata routing of this object.

Please check <u>User Guide</u> on how the routing mechanism works.

New in version 1.4.

Returns:

routing: MetadataRouter

A <u>MetadataRouter</u> encapsulating routing information.

get_params(deep=True)

[source]

Get parameters for this estimator.

Parameters:

deep: bool, default=True

If True, will return the parameters for this estimator and contained subobjects that are estimators.

Returns:

params: dict

Parameter names mapped to their values.

inverse_transform(Xt)

[source]

Call inverse_transform on the estimator with the best found params.

Only available if the underlying estimator implements inverse_transform and refit=True.

Parameters:

Xt : indexable, length n_samples

Must fulfill the input assumptions of the underlying estimator.

Returns:

X: {ndarray, sparse matrix} of shape (n_samples, n_features)

Result of the inverse_transform function for Xt based on the estimator with the best found parameters.

property n_features_in_

Number of features seen during fit.

Only available when refit=True.

predict(X)

[source]

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on the estimator with the best found parameters.

Only available if refit=True and the underlying estimator supports predict.

Parameters:

X: indexable, length n_samples

Must fulfill the input assumptions of the underlying estimator.

Returns:

y_pred : ndarray of shape (n_samples,)

The predicted labels or values for x based on the estimator with the best found parameters.

predict_log_proba(X)
[source]

Call predict_log_proba on the estimator with the best found parameters.

Only available if refit=True and the underlying estimator supports predict_log_proba.

Parameters:

X : indexable, length n_samples

Must fulfill the input assumptions of the underlying estimator.

Returns:

y_pred : ndarray of shape (n_samples,) or (n_samples, n_classes)

Predicted class log-probabilities for x based on the estimator with the best found parameters. The order of the classes corresponds to that in the fitted attribute <u>classes</u>.

predict_proba(X)
[source]

Call predict_proba on the estimator with the best found parameters.

Only available if refit=True and the underlying estimator supports predict_proba.

Parameters:

X: indexable, length n_samples

Must fulfill the input assumptions of the underlying estimator.

Returns:

y_pred : ndarray of shape (n_samples,) or (n_samples, n_classes)

Predicted class probabilities for x based on the estimator with the best found parameters. The order of the classes corresponds to that in the fitted attribute <u>classes</u>.

score(X, y=None, **params) [source]

Return the score on the given data, if the estimator has been refit.

This uses the score defined by scoring where provided, and the best_estimator_.score method otherwise.

Parameters:

X: array-like of shape (n_samples, n_features)

Input data, where n_samples is the number of samples and n_features is the number of features.

y: array-like of shape (n_samples, n_output) or (n_samples,), default=None

Target relative to X for classification or regression; None for unsupervised learning.

**params : dict

Parameters to be passed to the underlying scorer(s).

..versionadded:: 1.4

Only available if enable_metadata_routing=True. See Metadata Routing User Guide for more details.

Returns:

score: float

The score defined by scoring if provided, and the best_estimator_.score method otherwise.

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score_samples(X)

Call score_samples on the estimator with the best found parameters.

Only available if refit=True and the underlying estimator supports score_samples.

New in version 0.24.

Parameters:

X: iterable

Data to predict on. Must fulfill input requirements of the underlying estimator.

Returns:

y_score : ndarray of shape (n_samples,)

The best_estimator_.score_samples method.

set_params(**params) [source]

Set the parameters of this estimator.

The method works on simple estimators as well as on nested objects (such as Pipeline). The latter have parameters of the form component> so that it's possible to update each component of a nested object.

Parameters:

**params: dict

Estimator parameters.

Returns:

self: estimator instance

Estimator instance.

transform(X) [source]

Call transform on the estimator with the best found parameters.

Only available if the underlying estimator supports transform and refit=True.

Parameters:

X: indexable, length n_samples

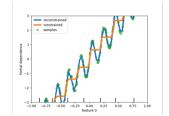
Must fulfill the input assumptions of the underlying estimator.

Returns:

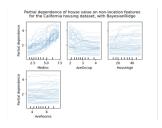
Xt: {ndarray, sparse matrix} of shape (n_samples, n_features)

x transformed in the new space based on the estimator with the best found parameters.

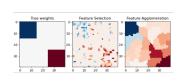
Examples using sklearn.model_selection.GridSearchCV



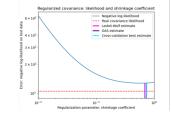
Release Highlights for scikit-learn 1.4



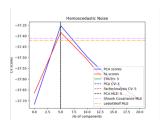
Release Highlights for scikit-learn 0.24



Feature agglomeration vs. univariate selection



Shrinkage covariance estimation: LedoitWolf vs OAS and max-likelihood

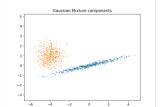


Model selection with Probabilistic PCA and Factor Analysis (FA)

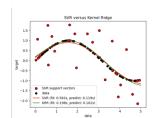
Toggle Menu



Comparing Random
Forests and Histogram
Gradient Boosting
models



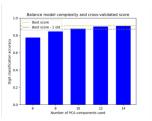
Gaussian Mixture Model Selection



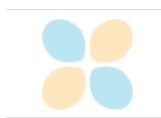
Comparison of kernel ridge regression and SVR



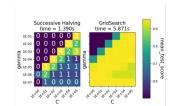
Displaying Pipelines



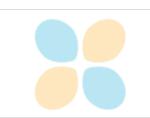
Balance model complexity and crossvalidated score



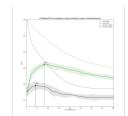
Comparing randomized search and grid search for hyperparameter estimation



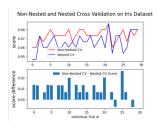
Comparison between grid search and successive halving



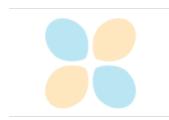
Custom refit strategy of a grid search with cross-validation



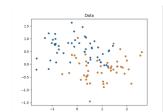
Demonstration of multimetric evaluation on cross_val_score and GridSearchCV



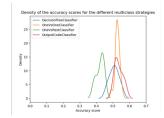
Nested versus nonnested cross-validation



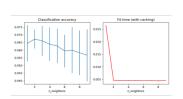
Sample pipeline for text feature extraction and evaluation



Statistical comparison of models using grid search



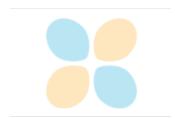
Overview of multiclass training meta-estimators



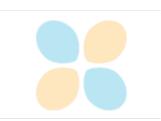
Caching nearest neighbors



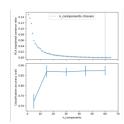
Kernel Density Estimation



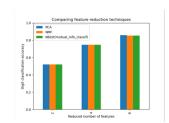
Column Transformer with Mixed Types



Concatenating multiple feature extraction methods



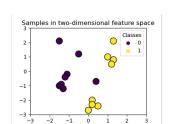
Pipelining: chaining a PCA and a logistic regression



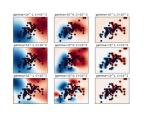
Selecting dimensionality reduction with Pipeline and GridSearchCV



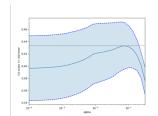
Feature discretization



Plot classification boundaries with different SVM Kernels



RBF SVM parameters



Cross-validation on diabetes Dataset Exercise

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