# sklearn.model selection.GridSearchCV

class sklearn.model\_selection. GridSearchCV(estimator, param\_grid, scoring=None, n\_jobs=None, iid='deprecated', 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 "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**.

#### **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 (string) 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: string, callable, list/tuple, dict or None, default: None

A single string (see <u>The scoring parameter: defining model evaluation rules</u>) or a callable (see <u>Defining your scoring strategy from metric functions</u>) to evaluate the predictions on the test set.

For evaluating multiple metrics, either give a list of (unique) strings or a dict with names as keys and callables as values.

NOTE that when using custom scorers, each scorer should return a single value. Metric functions returning a list/array of values can be wrapped into multiple scorers that return one value each.

See <u>Specifying multiple metrics for evaluation</u> for an example.

If None, the estimator's score method is used.

### n\_jobs : int or None, optional (default=None)

Number of jobs to run in parallel. None means 1 unless in a <u>joblib.parallel\_backend</u> context. -1 means using all processors. See <u>Glossary</u> for more details.

### pre\_dispatch : int, or string, optional

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 string, giving an expression as a function of n\_jobs, as in '2\*n\_jobs'

### iid: boolean, default=False

If True, return the average score across folds, weighted by the number of samples in each test set. In this case, the data is assumed to be identically distributed across the folds, and the loss minimized is the total loss per sample, and not the mean loss across the folds.

Deprecated since version 0.22: Parameter iid is deprecated in 0.22 and will be removed in 0.24

# cv: int, cross-validation generator or an iterable, optional

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,

Toggle Menu erable 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.

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

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

### refit: boolean, string, 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 string 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\_parameters\_ 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.

Changed in version 0.20: Support for callable added.

#### verbose: integer

Controls the verbosity: the higher, the more messages.

### error\_score : 'raise' or numeric

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. Default is np.nan.

### return\_train\_score : boolean, 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.

### **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
ʻrbf'	0.1	_	0.80		3
ʻrbfʻ	0.2	_	0.93		1

will be represented by a cv\_results\_ dict of:

#### **NOTE**

The key '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.

### See also:

# <u>ParameterGrid</u>

generates all the combinations of a hyperparameter grid.

sklearn.model\_selection.train\_test\_split

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

sklearn.metrics.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**

```
>>>
>>> from sklearn import svm, datasets
>>> from sklearn.model_selection import GridSearchCV
>>> iris = datasets.load_iris()
>>> parameters = {'kernel':('linear', 'rbf'), 'C':[1, 10]}
>>> svc = svm.SVC()
>>> clf = GridSearchCV(svc, parameters)
>>> clf.fit(iris.data, iris.target)
GridSearchCV(estimator=SVC(),
             param_grid={'C': [1, 10], 'kernel': ('linear', 'rbf')})
>>> sorted(clf.cv_results_.keys())
['mean_fit_time', 'mean_score_time', 'mean_test_score',...
 'param_C', 'param_kernel', 'params',...
 'rank_test_score', 'split0_test_score',...
 'split2_test_score', ...
 'std_fit_time', 'std_score_time', 'std_test_score']
```

#### **Methods**

<pre>decision_function(self, X)</pre>	Call decision_function on the estimator with the best found parameters.
<pre>fit(self, X[, y, groups])</pre>	Run fit with all sets of parameters.
<pre>get_params(self[, deep])</pre>	Get parameters for this estimator.
<pre>inverse_transform(self, Xt)</pre>	Call inverse_transform on the estimator with the best found params.
<pre>predict(self, X)</pre>	Call predict on the estimator with the best found parameters.
<pre>predict_log_proba(self, X)</pre>	Call predict_log_proba on the estimator with the best found parameters.
<pre>predict_proba(self, X)</pre>	Call predict_proba on the estimator with the best found parameters.
<pre>score(self, X[, y])</pre>	Returns the score on the given data, if the estimator has been refit.
<pre>set_params(self, \*\*params)</pre>	Set the parameters of this estimator.
<pre>transform(self, X)</pre>	Call transform on the estimator with the best found parameters.

\_\_init\_\_(self, estimator, param\_grid, scoring=None, n\_jobs=None, iid='deprecated', refit=True, cv=None, verbose=0, pre\_dispatch='2\*n\_jobs', error\_score=nan, return\_train\_score=False) [source]

Initialize self. See help(type(self)) for accurate signature.

```
{\tt decision\_function}(self, 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.

```
fit(self, X, y=None, groups=None, **fit_params) [source]
```

Run fit with all sets of parameters.

### **Parameters:**

## X: array-like of shape (n\_samples, n\_features)

<u>Training</u> vector, where n\_samples is the number of samples and n\_features is the number of features.

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# y: array-like of shape (n\_samples, n\_output) or (n\_samples,), optional

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

### groups: array-like, with shape (n\_samples,), optional

Group labels for the samples used while splitting the dataset into train/test set. Only used in conjunction with a "Group" <u>cv</u> instance (e.g., <u>GroupKFold</u>).

# \*\*fit\_params : dict of string -> object

Parameters passed to the fit method of the estimator

get\_params(self, 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: mapping of string to any

Parameter names mapped to their values.

inverse\_transform(self, 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.

predict(self, X)

[source]

Call predict 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.

predict\_log\_proba(self, 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.

predict\_proba(self, 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.

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score(self, X, y=None) [source]

Returns 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,), optional

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

#### **Returns:**

score: float

set\_params(self, \*\*params)

[source]

Set the parameters of this estimator.

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

#### **Parameters:**

\*\*params : dict

Estimator parameters.

#### **Returns:**

### self : object

Estimator instance.

transform(self, X) [source]

Call transform on the estimator with the best found parameters.

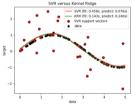
Only available if the underlying estimator supports transform and refit=True.

### **Parameters:**

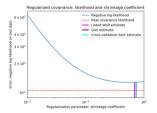
## X: indexable, length n\_samples

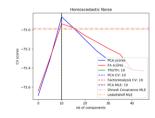
Must fulfill the input assumptions of the underlying estimator.

# Examples using sklearn.model\_selection.GridSearchCV







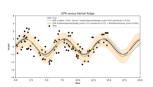




<u>Comparison of kernel</u> <u>ridge regression and SVR</u> Feature agglomeration vs. univariate selection

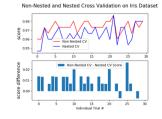
Shrinkage covariance
estimation: LedoitWolf vs
OAS and max-likelihood

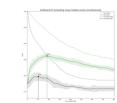
Model selection with Probabilistic PCA and Factor Analysis (FA) Faces recognition
example using
eigenfaces and SVMs











Parameter estimation using grid search with



Nested versus nonnested cross-validation

Demonstration of multimetric evaluation on

process regression

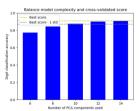
cross-validation

for hyperparameter estimation

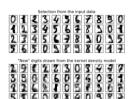
cross val score and GridSearchCV



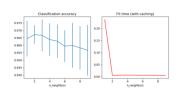
Sample pipeline for text feature extraction and evaluation



Balance model complexity and cross-validated score



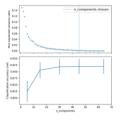
Kernel Density Estimation



<u>Caching nearest</u> <u>neighbors</u>



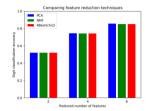
Concatenating multiple feature extraction methods



Pipelining: chaining a PCA and a logistic regression



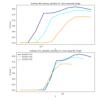
<u>Column Transformer</u> <u>with Mixed Types</u>



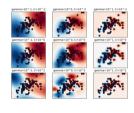
Selecting dimensionality reduction with Pipeline and GridSearchCV



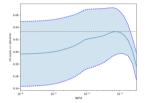
Feature discretization



Scaling the regularization parameter for SVCs



**RBF SVM parameters** 



<u>Cross-validation on</u> <u>diabetes Dataset</u> <u>Exercise</u>

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