3.2. Tuning the hyper-parameters of an estimator

Hyper-parameters are parameters that are not directly learnt within estimators. In scikit-learn they are passed as arguments to the constructor of the estimator classes. Typical examples include C, kernel and gamma for Support Vector Classifier, alpha for Lasso, etc.

It is possible and recommended to search the hyper-parameter space for the best <u>cross validation</u> score.

Any parameter provided when constructing an estimator may be optimized in this manner. Specifically, to find the names and current values for all parameters for a given estimator, use:

```
estimator.get_params()
```

A search consists of:

- an estimator (regressor or classifier such as sklearn.svm.SVC());
- a parameter space;
- a method for searching or sampling candidates;
- a cross-validation scheme; and
- a score function.

Two generic approaches to parameter search are provided in scikit-learn: for given values, <u>GridSearchCV</u> exhaustively considers all parameter combinations, while <u>RandomizedSearchCV</u> can sample a given number of candidates from a parameter space with a specified distribution. Both these tools have successive halving counterparts <u>HalvingGridSearchCV</u> and <u>HalvingRandomSearchCV</u>, which can be much faster at finding a good parameter combination.

After describing these tools we detail <u>best practices</u> applicable to these approaches. Some models allow for specialized, efficient parameter search strategies, outlined in <u>Alternatives to brute force parameter search</u>.

Note that it is common that a small subset of those parameters can have a large impact on the predictive or computation performance of the model while others can be left to their default values. It is recommended to read the docstring of the estimator class to get a finer understanding of their expected behavior, possibly by reading the enclosed reference to the literature.

3.2.1. Exhaustive Grid Search

The grid search provided by <u>GridSearchCV</u> exhaustively generates candidates from a grid of parameter values specified with the param_grid parameter. For instance, the following param_grid:

```
param_grid = [
    {'C': [1, 10, 100, 1000], 'kernel': ['linear']},
    {'C': [1, 10, 100, 1000], 'gamma': [0.001, 0.0001], 'kernel': ['rbf']},
]
```

specifies that two grids should be explored: one with a linear kernel and C values in [1, 10, 100, 1000], and the second one with an RBF kernel, and the cross-product of C values ranging in [1, 10, 100, 1000] and gamma values in [0.001, 0.0001].

The <u>GridSearchCV</u> instance implements the usual estimator API: when "fitting" it on a dataset all the possible combinations of parameter values are evaluated and the best combination is retained.

Examples:

- See <u>Custom refit strategy of a grid search with cross-validation</u> for an example of Grid Search computation on the digits dataset.
- See <u>Sample pipeline for text feature extraction and evaluation</u> for an example of Grid Search coupling parameters from a text documents feature extractor (n-gram count vectorizer and TF-IDF transformer) with a classifier (here a linear SVM trained with SGD with either elastic net or L2 penalty) using a pipeline.Pipeline instance.
- See <u>Nested versus non-nested cross-validation</u> for an example of Grid Search within a cross validation loop on the iris dataset. This is the best practice for evaluating the performance of a model with grid search.
- See <u>Demonstration of multi-metric evaluation on cross val score and GridSearchCV</u> for an example of <u>GridSearchCV</u> being used to evaluate multiple metrics simultaneously.
- See <u>Balance model complexity and cross-validated score</u> for an example of using refit=callable interface in <u>GridSearchCV</u>. The example shows how this interface adds certain amount of flexibility in identifying the "best" estimator. This interface can also be used in multiple metrics evaluation.
- See Statistical comparison of models using grid search for an example of how to do a statistical comparison on the outputs of GridSearchCV.

3.2.2. Randomized Parameter Optimization

While using a grid of parameter settings is currently the most widely used method for parameter optimization, other search methods have more favorable properties. RandomizedSearchCV implements a randomized search over parameters, where each setting is sampled from a distribution over possible parameter values. This has two main benefits over an exhaustive search:

- A budget can be chosen independent of the number of parameters and possible values.
- Adding parameters that do not influence the performance does not decrease efficiency.

Specifying how parameters should be sampled is done using a dictionary, very similar to specifying parameters for <u>GridSearchCV</u>. Additionally, a computation budget, being the number of sampled candidates or sampling iterations, is specified using the n_iter parameter. For each parameter, either a distribution over possible values or a list of discrete choices (which will be sampled uniformly) can be specified:

```
{'C': scipy.stats.expon(scale=100), 'gamma': scipy.stats.expon(scale=.1),
   'kernel': ['rbf'], 'class_weight':['balanced', None]}
```

This example uses the scipy.stats module, which contains many useful distributions for sampling parameters, such as expon, gamma, uniform or randint.

In principle, any function can be passed that provides a rvs (random variate sample) method to sample a value. A call to the rvs function should provide independent random samples from possible parameter values on consecutive calls.

Warning: The distributions in scipy.stats prior to version scipy 0.16 do not allow specifying a random state. Instead, they use the global numpy random state, that can be seeded via np.random.seed or set using np.random.set_state. However, beginning scikit-learn 0.18, the sklearn.model_selection module sets the random state provided by the user if scipy >= 0.16 is also available.

For continuous parameters, such as C above, it is important to specify a continuous distribution to take full advantage of the randomization. This way, increasing n_iter will always lead to a finer search.

A continuous log-uniform random variable is available through loguniform. This is a continuous version of log-spaced parameters. For example to specify C above, loguniform(1, 100) can be used instead of [1, 10, 100] or np.logspace(0, 2, num=1000). This is an alias to scipy.stats.loguniform.

Mirroring the example above in grid search, we can specify a continuous random variable that is log-uniformly distributed between 1e0 and 1e3:

```
from sklearn.utils.fixes import loguniform
{'C': loguniform(1e0, 1e3),
   'gamma': loguniform(1e-4, 1e-3),
   'kernel': ['rbf'],
   'class_weight':['balanced', None]}
```

Examples:

• <u>Comparing randomized search and grid search for hyperparameter estimation</u> compares the usage and efficiency of randomized search and grid search.

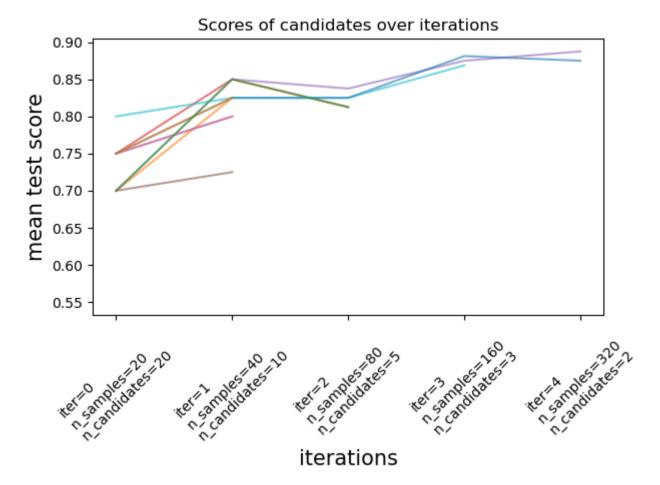
References:

• Bergstra, J. and Bengio, Y., Random search for hyper-parameter optimization, The Journal of Machine Learning Research (2012)

3.2.3. Searching for optimal parameters with successive halving

Scikit-learn also provides the HalvingGridSearchCV and HalvingRandomSearchCV estimators that can be used to search a parameter space using successive halving [1] [2]. Successive halving (SH) is like a tournament among candidate parameter combinations. SH is an iterative selection process where all candidates (the parameter combinations) are evaluated with a small amount of resources at the first iteration. Only some of these candidates are selected for the next iteration, which will be allocated more resources. For parameter tuning, the resource is typically the number of training samples, but it can also be an arbitrary numeric parameter such as n_estimators in a random forest.

As illustrated in the figure below, only a subset of candidates 'survive' until the last iteration. These are the candidates that have consistently ranked among the top-scoring candidates across all iterations. Each iteration is allocated an increasing amount of resources per candidate, here the number of samples.



We here briefly describe the main parameters, but each parameter and their interactions are described in more details in the sections below. The factor (> 1) parameter controls the rate at which the resources grow, and the rate at which the number of candidates decreases. In each iteration, the number of resources per candidate is multiplied by factor and the number of candidates is divided by the same factor. Along with resource and min_resources, factor is the most important parameter to control the search in our implementation, though a value of 3 usually works well. factor effectively controls the number of iterations in https://halvingRandomSearchCV and the number of available resources is small. More control is available through tuning the min_resources parameter.

These estimators are still **experimental**: their predictions and their API might change without any deprecation cycle. To use them, you need to explicitly import <code>enable_halving_search_cv</code>:

```
>>> # explicitly require this experimental feature
>>> from sklearn.experimental import enable_halving_search_cv # noqa
>>> # now you can import normally from model_selection
>>> from sklearn.model_selection import HalvingGridSearchCV
>>> from sklearn.model_selection import HalvingRandomSearchCV
```

Examples:

- Comparison between grid search and successive halving
- Successive Halving Iterations

3.2.3.1. Choosing min_resources and the number of candidates

Beside factor, the two main parameters that influence the behaviour of a successive halving search are the min_resources parameter, and the number of candidates (or parameter combinations) that are evaluated. min_resources is the amount of resources allocated at the first iteration for each candidate. The number of candidates is specified directly in HalvingRandomSearchCV, and is determined from the param_grid parameter of HalvingGridSearchCV.

Consider a case where the resource is the number of samples, and where we have 1000 samples. In theory, with min_resources=10 and factor=2, we are able to run **at most** 7 iterations with the following number of samples: [10, 20, 40, 80, 160, 320, 640].

But depending on the number of candidates, we might run less than 7 iterations: if we start with a **small** number of candidates, the last iteration might use less than 640 samples, which means not using all the available resources (samples). For example if we start with 5 candidates, we only need 2 iterations: 5 candidates for the first iteration, then 5 // 2 = 2 candidates at the second iteration, after which we know which candidate performs the best (so we don't need a third one). We would only be using at most 20 samples which is a waste since we have 1000 samples at our disposal. On the other hand, if we start with a **high** number of candidates, we might end up with a lot of candidates at the last iteration, which may not always be ideal: it means that many candidates will run with the full resources, basically reducing the procedure to standard search.

In the case of <u>HalvingRandomSearchCV</u>, the number of candidates is set by default such that the last iteration uses as much of the available resources as possible. For <u>HalvingGridSearchCV</u>, the number of candidates is determined by the param_grid parameter. Changing the value of min resources will impact the number of possible iterations, and as a result will also have an effect on the ideal number of candidates.

Another consideration when choosing min_resources is whether or not it is easy to discriminate between good and bad candidates with a small amount of resources. For example, if you need a lot of samples to distinguish between good and bad parameters, a high min_resources is recomtogle Menu be other hand if the distinction is clear even with a small amount of samples, then a small min_resources may be preferable since it

would speed up the computation.

Notice in the example above that the last iteration does not use the maximum amount of resources available: 1000 samples are available, yet only 640 are used, at most. By default, both HalvingRandomSearchCV and HalvingRandomSearchCV achieves this by sampling the right amount of candidates, while HalvingRandomSearchCV achieves this by sampling the right amount of candidates, while HalvingGridSearchCV achieves this by sampling the right amount of candidates, while HalvingGridSearchCV achieves this by properly setting min resources. Please see Example.com/Example.c

3.2.3.2. Amount of resource and number of candidates at each iteration

At any iteration i, each candidate is allocated a given amount of resources which we denote n_resources_i. This quantity is controlled by the parameters factor and min_resources as follows (factor is strictly greater than 1):

```
n_resources_i = factor**i * min_resources,
```

or equivalently:

```
n_resources_{i+1} = n_resources_i * factor
```

where min_resources == n_resources_0 is the amount of resources used at the first iteration. factor also defines the proportions of candidates that will be selected for the next iteration:

```
n_candidates_i = n_candidates // (factor ** i)
```

or equivalently:

```
n_candidates_0 = n_candidates
n_candidates_{i+1} = n_candidates_i // factor
```

So in the first iteration, we use min_resources resources n_candidates times. In the second iteration, we use min_resources * factor resources n_candidates // factor times. The third again multiplies the resources per candidate and divides the number of candidates. This process stops when the maximum amount of resource per candidate is reached, or when we have identified the best candidate. The best candidate is identified at the iteration that is evaluating factor or less candidates (see just below for an explanation).

Here is an example with min_resources=3 and factor=2, starting with 70 candidates:

n_resources_i	n_candidates_i	
3 (=min_resources)	70 (=n_candidates)	
3 * 2 = 6	70 // 2 = 35	
6 * 2 = 12	35 // 2 = 17	
12 * 2 = 24	17 // 2 = 8	
24 * 2 = 48	8 // 2 = 4	
48 * 2 = 96	4 // 2 = 2	
4		

We can note that:

- the process stops at the first iteration which evaluates factor=2 candidates: the best candidate is the best out of these 2 candidates. It is not necessary to run an additional iteration, since it would only evaluate one candidate (namely the best one, which we have already identified). For this reason, in general, we want the last iteration to run at most factor candidates. If the last iteration evaluates more than factor candidates, then this last iteration reduces to a regular search (as in RandomizedSearchCV or GridSearchCV).
- each n_resources_i is a multiple of both factor and min_resources (which is confirmed by its definition above).

The amount of resources that is used at each iteration can be found in the n_resources_ attribute.

3.2.3.3. Choosing a resource

By default, the resource is defined in terms of number of samples. That is, each iteration will use an increasing amount of samples to train on. You can however manually specify a parameter to use as the resource with the resource parameter. Here is an example where the resource is defined in terms of the number of estimators of a random forest:

```
>>>
>>> from sklearn.datasets import make_classification
>>> from sklearn.ensemble import RandomForestClassifier
>>> from sklearn.experimental import enable halving search cv # noga
>>> from sklearn.model_selection import HalvingGridSearchCV
>>> import pandas as pd
>>>
>>> param_grid = {'max_depth': [3, 5, 10],
                  'min_samples_split': [2, 5, 10]}
>>> base_estimator = RandomForestClassifier(random_state=0)
>>> X, y = make_classification(n_samples=1000, random_state=0)
>>> sh = HalvingGridSearchCV(base_estimator, param_grid, cv=5,
                             factor=2, resource='n_estimators',
. . .
                             max resources=30).fit(X, y)
>>> sh.best estimator
RandomForestClassifier(max_depth=5, n_estimators=24, random_state=0)
```

Note that it is not possible to budget on a parameter that is part of the parameter grid.

3.2.3.4. Exhausting the available resources

As mentioned above, the number of resources that is used at each iteration depends on the min_resources parameter. If you have a lot of resources available but start with a low number of resources, some of them might be wasted (i.e. not used):

The search process will only use 80 resources at most, while our maximum amount of available resources is $n_samples=1000$. Here, we have $min_resources = r_0 = 20$.

For <u>HalvingGridSearchCV</u>, by default, the min_resources parameter is set to 'exhaust'. This means that min_resources is automatically set such that the last iteration can use as many resources as possible, within the max_resources limit:

```
>>> sh = HalvingGridSearchCV(base_estimator, param_grid, cv=5,
... factor=2, min_resources='exhaust').fit(X, y)
>>> sh.n_resources_
[250, 500, 1000]
```

min_resources was here automatically set to 250, which results in the last iteration using all the resources. The exact value that is used depends on the number of candidate parameter, on max_resources and on factor.

For HalvingRandomSearchCV, exhausting the resources can be done in 2 ways:

- by setting min_resources='exhaust', just like for HalvingGridSearchCV;
- by setting n_candidates='exhaust'.

Both options are mutally exclusive: using min_resources='exhaust' requires knowing the number of candidates, and symmetrically n_candidates='exhaust' requires knowing min_resources.

In general, exhausting the total number of resources leads to a better final candidate parameter, and is slightly more time-intensive.

3.2.3.5. Aggressive elimination of candidates

Ideally, we want the last iteration to evaluate factor candidates (see <u>Amount of resource and number of candidates at each iteration</u>). We then just have to pick the best one. When the number of available resources is small with respect to the number of candidates, the last iteration may have to evaluate more than factor candidates:

```
>>> from sklearn.datasets import make classification
                                                                                                                            >>>
>>> from sklearn.svm import SVC
>>> from sklearn.experimental import enable_halving_search_cv # noqa
>>> from sklearn.model_selection import HalvingGridSearchCV
>>> import pandas as pd
>>>
>>>
>>> param_grid = {'kernel': ('linear', 'rbf'),
                  'C': [1, 10, 100]}
>>> base_estimator = SVC(gamma='scale')
>>> X, y = make_classification(n_samples=1000)
>>> sh = HalvingGridSearchCV(base_estimator, param_grid, cv=5,
                             factor=2, max_resources=40,
                             aggressive_elimination=False).fit(X, y)
. . .
>>> sh.n_resources_
[20, 40]
>>> sh.n_candidates_
[6, 3]
```

Since we cannot use more than max_resources=40 resources, the process has to stop at the second iteration which evaluates more than factor=2 candidates.

Using the aggressive_elimination parameter, you can force the search process to end up with less than factor candidates at the last iteration. To do this, the process will eliminate as many candidates as necessary using min_resources resources:

```
>>> sh = HalvingGridSearchCV(base_estimator, param_grid, cv=5,
... factor=2,
... max_resources=40,
... aggressive_elimination=True,
... ).fit(X, y)
>>> sh.n_resources_
[20, 20, 40]
>>> sh.n_candidates_
[6, 3, 2]
```

Notice that we end with 2 candidates at the last iteration since we have eliminated enough candidates during the first iterations, using $n_resources = min_resources = 20$.

3.2.3.6. Analyzing results with the cv_results_ attribute

The cv_results_ attribute contains useful information for analyzing the results of a search. It can be converted to a pandas dataframe with df = pd.DataFrame(est.cv_results_). The cv_results_ attribute of HalvingGridSearchCV and HalvingRandomSearchCV is similar to that of GridSearchCV and RandomizedSearchCV, with additional information related to the successive halving process.

Here is an example with some of the columns of a (truncated) dataframe:

	iter	n_resources	mean_test_score	params
0	0	125	0.983667	{'criterion': 'log_loss', 'max_depth': None, 'max_features': 9, 'min_samples_split': 5}
1	0	125	0.983667	{'criterion': 'gini', 'max_depth': None, 'max_features': 8, 'min_samples_split': 7}
2	0	125	0.983667	{'criterion': 'gini', 'max_depth': None, 'max_features': 10, 'min_samples_split': 10}
3	0	125	0.983667	{'criterion': 'log_loss', 'max_depth': None, 'max_features': 6, 'min_samples_split': 6}
		•••	•••	
15	2	500	0.951958	{'criterion': 'log_loss', 'max_depth': None, 'max_features': 9, 'min_samples_split': 10}
16	2	500	0.947958	{'criterion': 'gini', 'max_depth': None, 'max_features': 10, 'min_samples_split': 10}
17	2	500	0.951958	{'criterion': 'gini', 'max_depth': None, 'max_features': 10, 'min_samples_split': 4}
18	3	1000	0.961009	{'criterion': 'log_loss', 'max_depth': None, 'max_features': 9, 'min_samples_split': 10}
19	3	1000	0.955989	{'criterion': 'gini', 'max_depth': None, 'max_features': 10, 'min_samples_split': 4}
4				

Each row corresponds to a given parameter combination (a candidate) and a given iteration. The iteration is given by the iter column. The n_resources column tells you how many resources were used.

In the example above, the best parameter combination is {'criterion': 'log_loss', 'max_depth': None, 'max_features': 9, 'min_samples_split': 10} since it has reached the last iteration (3) with the highest score: 0.96.

References:

[1]

K. Jamieson, A. Talwalkar, Non-stochastic Best Arm Identification and Hyperparameter Optimization, in proc. of Machine Learning Research, 2016.

<u>[2]</u>

L. Li, K. Jamieson, G. DeSalvo, A. Rostamizadeh, A. Talwalkar, <u>Hyperband: A Novel Bandit-Based Approach to Hyperparameter Optimization</u>, in <u>Machine Learning</u> Research 18, 2018.

3.2.4. Tips for parameter search

3.2.4.1. Specifying an objective metric

By default, parameter search uses the score function of the estimator to evaluate a parameter setting. These are the sklearn.metrics.accuracy_score for classification and sklearn.metrics.r2_score for regression. For some applications, other scoring functions are better suited (for example in unbalanced classification, the accuracy score is often uninformative). An alternative scoring function can be specified via the scoring parameter of most parameter search tools. See These are the sklearn.metrics.accuracy_score for regression. For some applications, other scoring functions are better suited (for example in unbalanced classification, the accuracy score is often uninformative). An alternative scoring function can be specified via the scoring parameter of most parameter search tools. See These are the scoring function of the estimator of the scoring parameter search tools. The scoring parameter search tools are the scoring parameter search tools.

3.2.4.2. Specifying multiple metrics for evaluation

<u>GridSearchCV</u> and <u>RandomizedSearchCV</u> allow specifying multiple metrics for the scoring parameter.

Multimetric scoring can either be specified as a list of strings of predefined scores names or a dict mapping the scorer name to the scorer function and/or the predefined scorer name(s). See <u>Using multiple metric evaluation</u> for more details.

When specifying multiple metrics, the refit parameter must be set to the metric (string) for which the best_params_ will be found and used to build the best_estimator_ on the whole dataset. If the search should not be refit, set refit=False. Leaving refit to the default value None will result in an error when using multiple metrics.

See <u>Demonstration of multi-metric evaluation on cross val score and GridSearchCV</u> for an example usage.

HalvingRandomSearchCV and HalvingGridSearchCV do not support multimetric scoring.

3.2.4.3. Composite estimators and parameter spaces

<u>GridSearchCV</u> and <u>RandomizedSearchCV</u> allow searching over parameters of composite or nested estimators such as <u>Pipeline</u>, <u>ColumnTransformer</u>, <u>VotingClassifier</u> or <u>CalibratedClassifierCV</u> using a dedicated <estimator>__parameter syntax:

Here, <estimator> is the parameter name of the nested estimator, in this case estimator. If the meta-estimator is constructed as a collection of estimators as in pipeline. Pipeline, then <estimator> refers to the name of the estimator, see Nested parameters. In practice, there can be several levels of nesting:

Please refer to <u>Pipeline: chaining estimators</u> for performing parameter searches over pipelines.

3.2.4.4. Model selection: development and evaluation

Model selection by evaluating various parameter settings can be seen as a way to use the labeled data to "train" the parameters of the grid.

When evaluating the resulting model it is important to do it on held-out samples that were not seen during the grid search process: it is recommended to split the data into a **development set** (to be fed to the <u>GridSearchCV</u> instance) and an **evaluation set** to compute performance metrics.

This can be done by using the <u>train test split</u> utility function.

3.2.4.5. Parallelism

The parameter search tools evaluate each parameter combination on each data fold independently. Computations can be run in parallel by using the keyword n_jobs=-1. See function signature for more details, and also the Glossary entry for n_jobs.

3.2.4.6. Robustness to failure

Some parameter settings may result in a failure to fit one or more folds of the data. By default, this will cause the entire search to fail, even if some parameter settings could be fully evaluated. Setting error_score=0 (or =np.NaN) will make the procedure robust to such failure, issuing a warning and setting the score for that fold to 0 (or NaN), but completing the search.

3.2.5. Alternatives to brute force parameter search

3.2.5.1. Model specific cross-validation

Some models can fit data for a range of values of some parameter almost as efficiently as fitting the estimator for a single value of the parameter. This feature can be leveraged to perform a more efficient cross-validation used for model selection of this parameter.

The most common parameter amenable to this strategy is the parameter encoding the strength of the regularizer. In this case we say that we compute the **regularization path** of the estimator.

Here is the list of such models:

<pre>linear_model.ElasticNetCV(*[, 1_ratio,])</pre>	Elastic Net model with iterative fitting along a regularization path.
<pre>linear_model.LarsCV(*[, fit_intercept,])</pre>	Cross-validated Least Angle Regression model.
<pre>linear_model.LassoCV(*[, eps, n_alphas,])</pre>	Lasso linear model with iterative fitting along a regularization path.
<pre>linear_model.LassoLarsCV(*[, fit_intercept,</pre>]) Cross-validated Lasso, using the LARS algorithm.
<pre>linear_model.LogisticRegressionCV(*[, Cs,</pre>]) Logistic Regression CV (aka logit, MaxEnt) classifier.
<pre>linear_model.MultiTaskElasticNetCV(*[,])</pre>	Multi-task L1/L2 ElasticNet with built-in cross-validation.
<pre>linear_model.MultiTaskLassoCV(*[, eps,])</pre>	Multi-task Lasso model trained with L1/L2 mixed-norm as regularizer.
<pre>linear_model.OrthogonalMatchingPursuitC</pre>	<u>v</u> (*) Cross-validated Orthogonal Matching Pursuit model (OMP).
<pre>linear_model.RidgeCV([alphas,])</pre>	Ridge regression with built-in cross-validation.
<pre>linear_model.RidgeClassifierCV([alphas,</pre>	Ridge classifier with built-in cross-validation.
4	

3.2.5.2. Information Criterion

Some models can offer an information-theoretic closed-form formula of the optimal estimate of the regularization parameter by computing a single regularization path (instead of several when using cross-validation).

Here is the list of models benefiting from the Akaike Information Criterion (AIC) or the Bayesian Information Criterion (BIC) for automated model selection:

<u>linear_model.LassoLarsIC</u>([criterion, ...]) Lasso model fit with Lars using BIC or AIC for model selection.

3.2.5.3. Out of Bag Estimates

When using ensemble methods base upon bagging, i.e. generating new training sets using sampling with replacement, part of the training set remains unused. For each classifier in the ensemble, a different part of the training set is left out.

This left out portion can be used to estimate the generalization error without having to rely on a separate validation set. This estimate comes "for free" as no additional data is needed and can be used for model selection.

This is currently implemented in the following classes:

<pre>ensemble.RandomForestClassifier([])</pre>	A random forest classifier.
<pre>ensemble.RandomForestRegressor([])</pre>	A random forest regressor.
<pre>ensemble.ExtraTreesClassifier([])</pre>	An extra-trees classifier.
<pre>ensemble.ExtraTreesRegressor([n_estimators,])</pre>	An extra-trees regressor.
<pre>ensemble.GradientBoostingClassifier(*[,])</pre>	Gradient Boosting for classification.
<pre>ensemble.GradientBoostingRegressor(*[,])</pre>	Gradient Boosting for regression.
4	

© 2007 - 2023, scikit-learn developers (BSD License). Show this page source