## **Model Tuning**

Each machine learning model has a set of options or configurations that can be tuned to optimize the model when fitting to data. These configurations are called **hyper-parameters**. Hence, for each hyper-parameter, there exist a range of values that can be chosen. Taking into consideration the number of hyper-parameters that an algorithm has, the entire space can become exponentially large and infeasible to explore all of them. Scikit-learn provides two convenient modules for searching through the hyper-parameter space of an algorithm to find the best values for each hyper-parameter that optimizes the model.

These modules are the

- Grid search
- Randomized search

## **Grid Search**

Grid search comprehensively explores all the specified hyper-parameter values for an estimator. It is implemented using the **GridSearchCV** module. Let's see an example using the Random forest for regression. The hyper-parameters we'll search over are

- The number of trees in the forest, n estimators
- The maximum depth of the tree, **max\_depth**
- The minimum number of samples required to split an internal node, min\_samples\_leaf

```
from sklearn.model_selection import GridSearchCV
from sklearn.ensemble import RandomForestRegressor
from sklearn import datasets

# load dataset
data = datasets.load_boston()

# separate features and target
X = data.data
y = data.target
```

```
# construct grid search parameters in a dictionary
parameters = {
    'n estimators': [2, 4, 6, 8, 10, 12, 14, 16],
    'max depth': [2, 4, 6, 8],
    'min samples leaf': [1,2,3,4,5]
# create the model
rf model = RandomForestRegressor()
# run the grid search
grid search = GridSearchCV(estimator=rf model, param grid=parameters)
# fit the model
grid_search.fit(X,y)
'Output':
GridSearchCV(cv=None, error score='raise',
       estimator=RandomForestRegressor(bootstrap=True, criterion='mse',
       max depth=None,
           max features='auto', max leaf nodes=None,
           min impurity decrease=0.0, min impurity split=None,
           min samples leaf=1, min samples split=2,
           min weight fraction leaf=0.0, n estimators=10, n jobs=1,
           oob score=False, random state=None, verbose=0, warm
           start=False),
       fit params=None, iid=True, n jobs=1,
       param grid={'n estimators': [2, 4, 6, 8, 10, 12, 14, 16],
       'max depth': [2, 4, 6, 8], 'min samples leaf': [1, 2, 3, 4, 5]},
       pre dispatch='2*n jobs', refit=True, return train score='warn',
       scoring=None, verbose=0)
# evaluate the model performance
print("Best Accuracy: %.3f%%" % (grid search.best score *100.0))
'Output':
Best Accuracy: 57.917%
```

## **Randomized Search**

As opposed to grid search, not all the provided hyper-parameter values are evaluated, but rather a determined number of hyper-parameter values are sampled from a random uniform distribution. The number of hyper-parameter values that can be evaluated is determined by the **n\_iter** attribute of the **RandomizedSearchCV** module.

In this example, we will use the same scenario as in the grid search case.

```
from sklearn.model_selection import RandomizedSearchCV
from sklearn.ensemble import RandomForestRegressor
from sklearn import datasets

# load dataset
data = datasets.load_boston()

# separate features and target
X = data.data
y = data.target

# construct grid search parameters in a dictionary
parameters = {
    'n_estimators': [2, 4, 6, 8, 10, 12, 14, 16],
    'max_depth': [2, 4, 6, 8],
    'min_samples_leaf': [1,2,3,4,5]
    }
```