

Performance Evaluation, Ensemble Methods

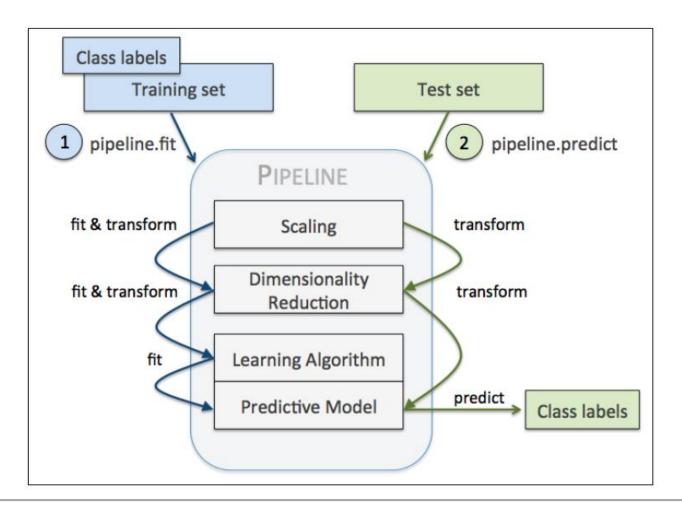
Machine Learning

Contents

- Model Evaluation & Hyperparameter Tuning
 - Pipelining
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 - Learning Curve
 - Validation Curve
 - Grid Search
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- Ensemble Methods
 - Bagging
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Pipelining : Streamlining workflows with pipelines



Loading the Breast Cancer Wisconsin dataset

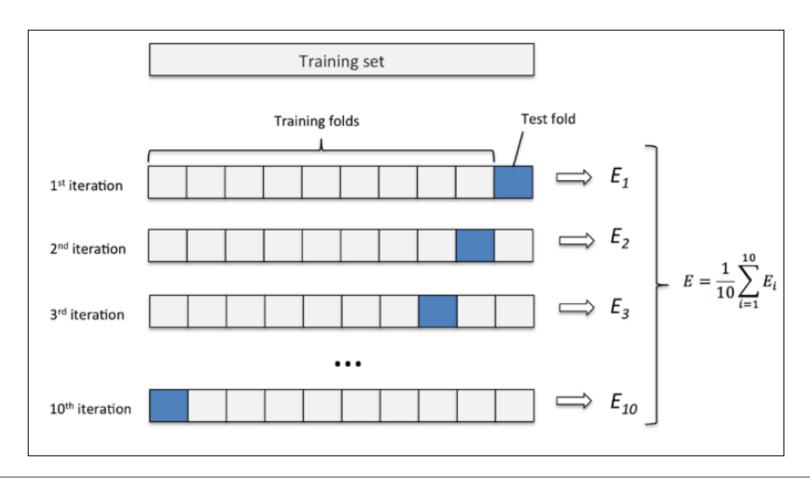
```
import pandas as pd
df = pd.read csv('https://archive.ics.uci.edu/ml/'
                  'machine-learning-databases'
                  '/breast-cancer-wisconsin/wdbc.data', header=None)
df.head()
df.shape
                                                                    842302 M 17.99 10.38 122.80 100
from sklearn.preprocessing import LabelEncoder
                                                                    842517 M 20.57 17.77 132.90 132
# Get X, y. Encoding class Tabel with scikit-learn
X = df.loc[:, 2:].values # remove 1st column(ID) & 2nd column 2 84300903 M
                                                                            19.69 21.25 130.00 120
y = df.loc[:, 1].values
                                                                3 84348301 M 11.42 20.38
                                                                                        77.58
                                                                  84358402 M 20.29 14.34 135.10 129
le = LabelEncoder()
y = le.fit transform(y)
                                                                (569, 32)
X.shape
                                                                (569, 30)
from sklearn.model selection import train test split
# train / test split
X train, X test, y train, y test = train test split(X, y,
                                             test size=0.20,
                                             stratify=y,
                                             random state=1)
X train.shape
                                                                 (455, 30)
```

Combining transformers and estimators in a pipeline

```
from sklearn.preprocessing import StandardScaler
from sklearn.decomposition import PCA
from sklearn.linear model import LogisticRegression
from sklearn.pipeline import make_pipeline, Pipeline
# make pipeline
pipe lr = make pipeline(StandardScaler(),
                      PCA(n components=2),
                      LogisticRegression(random state=1))
# Instead you can use Pipeline class
pipe lr = Pipeline(steps=[
     ('scale', StandardScaler()),
     ('PCA', PCA(n components=2)),
     ('LR', LogisticRegression(random state=1))
116
# process using pipeline
pipe lr.fit(X train, y train)
print('Test Accuracy: %.3f' % pipe lr.score(X test, y test))
```

Test Accuracy: 0.956

Using k-fold cross validation to assess model performance

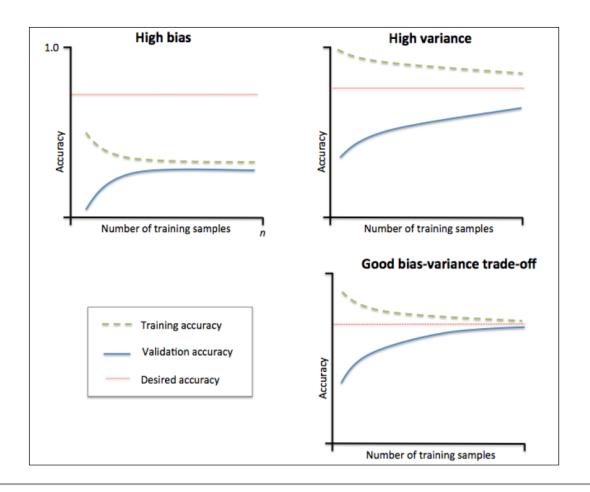


Using k-fold cross validation to assess model performance

```
import numpy as np
from sklearn.model selection import StratifiedKFold
# make indices for stratified k-fold cross validation
kfold = StratifiedKFold(n splits=5,
                      shuffle=False, # random state=1 sklean <= 0.22</pre>
                      ).split(X, y)
# train and compute test score for each set
scores = []
for k, (train, test) in enumerate(kfold):
    pipe lr.fit(X[train], y[train])
    score = pipe lr.score(X[test], y[test])
    scores.append(score)
    print('Fold: %2d, Class dist.: %s, Acc: %.3f' % (k+1,
          np.bincount(y[train]), score))
print('\nCV accuracy: %.3f +/- %.3f' % (np.mean(scores), np.std(scores)))
Fold: 1, Class dist.: [286 169], Acc: 0.939
Fold: 5, Class dist.: [286 170], Acc: 0.982
CV accuracy: 0.951 +/- 0.016
```

Using k-fold cross validation to assess model performance

Checking performance with learning curves



Checking performance with learning curves

```
import matplotlib.pyplot as plt
from sklearn.model selection import learning curve
pipe lr = make pipeline(StandardScaler(),
                      LogisticRegression(penalty='12',
                                      random state=1,
                                      max iter=500))
# accuracies for different training set size
train sizes, train scores, test scores =\
                     learning curve(estimator=pipe lr,
                     X=X,
                     y=y,
                     train sizes=np.linspace(0.1, 1.0, 10),
                     cv=10,
                     n jobs=1)
print(train sizes)
print(train scores.shape)
[ 51 102 153 204 256 307 358 409 460 512]
(10, 10)
```

Checking performance with learning curves

```
train mean = np.mean(train scores, axis=1)
test mean = np.mean(test scores, axis=1)
plt.plot(train sizes, train mean,
            color='blue', marker='o',
            label='training accuracy')
plt.plot(train sizes, test mean,
            color='green', linestyle='--', marker='s',
            label='validation accuracy')
plt.grid()
plt.xlabel('Number of training samples')
                                                         1.00
plt.ylabel('Accuracy')
plt.legend(loc='lower right')
                                                         0.95
plt.ylim([0.8, 1.03])
plt.tight layout()
                                                       4ccuracy
                                                         0.90
plt.show()
                                                         0.85
                                                                                              training accuracy
                                                                                              validation accuracy
                                                         0.80
                                                                  100
                                                                                              400
                                                                                                        500
                                                                            200
                                                                                     300
                                                                           Number of training samples
```

Checking overfitting and underfitting with validation curves

Checking overfitting and underfitting with validation curves

```
train mean = np.mean(train scores, axis=1)
test mean = np.mean(test scores, axis=1)
plt.plot(param range, train mean,
             color='blue', marker='o',
             label='training accuracy')
plt.plot(param range, test mean,
             color='green', linestyle='--', marker='s',
            label='validation accuracy')
plt.grid()
                                                                   1.000
plt.xscale('log')
plt.legend(loc='lower right')
                                                                   0.975
plt.xlabel('Parameter C')
                                                                   0.950
plt.ylabel('Accuracy')
                                                                   0.925
plt.ylim([0.8, 1.0])
                                                                  Accuracy
plt.tight layout()
                                                                   0.900
                                                                   0.875
plt.show()
                                                                   0.850
                                                                   0.825
                                                                                                      training accuracy
                                                                    0.800
                                                                        10^{-3}
                                                                                10^{-2}
                                                                                        10-1
                                                                                                 10°
                                                                                                        10<sup>1</sup>
                                                                                                                10^{2}
                                                                                          Parameter C
```

Tuning hyperparameters via grid search

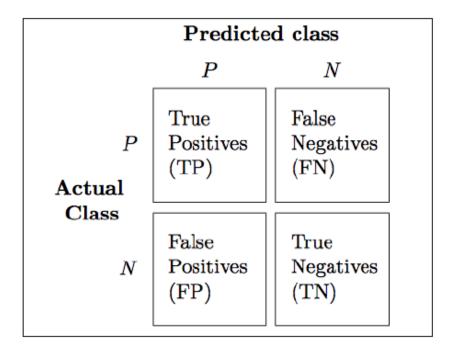
```
from sklearn.model selection import GridSearchCV
                                                        nan (+/-nan) for {'C': 0.01, 'penalty': 'l1'}
                                                        0.949 (+/-0.026) for {'C': 0.01, 'penalty': '12'}
                                                        nan (+/-nan) for {'C': 0.1, 'penalty': 'l1'}
stdsc = StandardScaler()
                                                        0.975 (+/-0.013) for {'C': 0.1, 'penalty': '12'}
X = stdsc.fit transform(X)
                                                        nan (+/-nan) for {'C': 1.0, 'penalty': 'l1'}
                                                        0.981 (+/-0.013) for {'C': 1.0, 'penalty': '12'}
# training with various parameter combinations
                                                        nan (+/-nan) for {'C': 10.0, 'penalty': 'l1'}
param grid = [\{'C': [0.01, 0.1, 1.0, 10.0, 100.0],
                                                        0.970 (+/-0.028) for {'C': 10.0, 'penalty': '12'}
                'penalty': ['l1', 'l2']}]
                                                        nan (+/-nan) for {'C': 100.0, 'penalty': 'l1'}
                                                        0.963 (+/-0.037) for {'C': 100.0, 'penalty': 'l2'}
gs = GridSearchCV(estimator=LogisticRegression(),
                 param grid=param grid,
                                                        Best score: 0.9806862288464524
                 scoring='accuracy',
                                                        Parameters: {'C': 1.0, 'penalty': '12'}
                 cv=5,
                 n jobs=-1
gs = gs.fit(X, y)
means = gs.cv results ['mean test score']
stds = gs.cv results ['std test score']
params = gs.cv results ['params']
for mean, std, params in zip(means, stds, params):
    print("%0.3f (+/-%0.3f) for %s" % (mean, std * 2, params))
print()
print("Best score:", gs.best score )
print("Parameters:", gs.best params )
```

Tuning hyperparameters via grid search

```
# the best model
clf = gs.best_estimator_
clf.fit(X_train, y_train)
print('Test accuracy: %.3f' % clf.score(X_test, y_test))
Test accuracy: 0.947
```

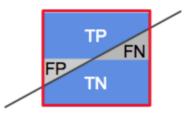
Confusion Matrix

 The confusion matrix is simply a square matrix that reports the counts of the true positive, true negative, false positive, and false negative predictions of a classifier



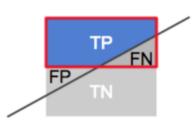
- ACC, TPR, FPR
 - Accuracy(ACC)

$$ACC = \frac{TP + TN}{FP + FN + TP + TN}$$



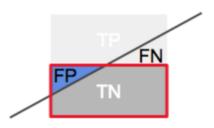
True Positive Rate(TPR)

$$TPR = \frac{TP}{P} = \frac{TP}{FN + TP}$$



False Positive Rate(FPR)

$$FPR = \frac{FP}{N} = \frac{FP}{FP + TN}$$

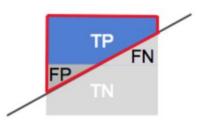


https://bcho.tistory.com/1206

Performance Evaluation Metrics

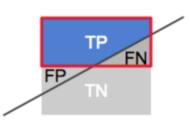
- Precision, Recall, F1
 - Precision(PRE)

$$PRE = \frac{TP}{TP + FP}$$



Recall(REC)

$$REC = TPR = \frac{TP}{P} = \frac{TP}{FN + TP}$$



• F1 - score

$$F1 = 2\frac{PRE \times REC}{PRE + REC}$$



Confusion Matrix

```
from sklearn.metrics import confusion_matrix

clf.fit(X_train, y_train)
y_pred = clf.predict(X_test)
confmat = confusion_matrix(y_true=y_test, y_pred=y_pred)
print(confmat)

[[68   4]
[ 2  40]]
```

Precision, Recall, and F1

```
from sklearn.metrics import precision_score, recall_score, f1_score

print('Precision: %.3f' % precision_score(y_true=y_test, y_pred=y_pred))
print('Recall: %.3f' % recall_score(y_true=y_test, y_pred=y_pred))
print('F1: %.3f' % f1_score(y_true=y_test, y_pred=y_pred))

Precision: 0.909
Recall: 0.952
F1: 0.930
```

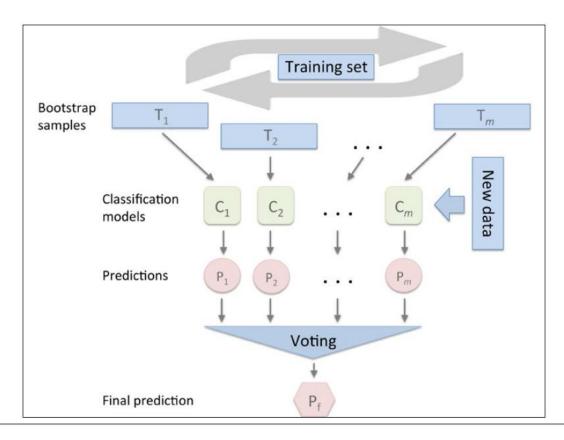
ROC (Receiver Operating Characteristic) Curve

```
from sklearn.metrics import roc curve, auc
from scipy import interp
pipe lr = make pipeline(StandardScaler(),
                     PCA(n components=2),
                      LogisticRegression(penalty='12',
                                      random state=1,
                                      C=0.1)
X train2 = X train[:, [4, 5]]
X \text{ test2} = X \text{ test[:, [4, 5]]}
# prediction probability of class 1
pipe lr.fit(X train2, y train)
probas = pipe lr.predict proba(X test2)
probas[:,1]
array([ 0.54725907, 0.16785314, 0.53630409, 0.64486632, 0.43203903,
        0.45691422, 0.86281954, 0.35614241, 0.32248996, 0.09339228,
        0.14273763, 0.13536069, 0.95496384, 0.26308029, 0.68040328,
        0.51767461, 0.43428621, 0.08286686, 0.37322008, 0.3374638,
        0.27978014, 0.14773322, 0.13018966, 0.23524479, 0.85951027,
```

ROC (Receiver Operating Characteristic) Curve

```
# FPR, TPR, AUC
fpr, tpr, thresholds = roc curve(y test, probas[:,1], pos label=1)
roc auc = auc(fpr, tpr)
plt.plot(fpr, tpr,
            label='ROC (area = %0.2f)' % roc auc)
plt.plot([0, 1], [0, 1],
            linestyle='--',
                                                                        ROC with 2 features 4, 5
            label='random guessing')
                                                     1.0
plt.xlim([-0.05, 1.05])
                                                     0.8
plt.ylim([-0.05, 1.05])
plt.title('ROC with 2 features 4, 5')
                                                   true positive rate
plt.xlabel('false positive rate')
                                                     0.6
plt.ylabel('true positive rate')
plt.legend(loc="lower right")
                                                     0.4
plt.tight layout()
                                                     0.2
plt.show()
                                                                                               ROC (area = 0.84)
                                                                                               random guessing
                                                     0.0
                                                                   0.2
                                                          0.0
                                                                             0.4
                                                                                       0.6
                                                                                                0.8
                                                                                                          1.0
                                                                            false positive rate
```

- Building an ensemble of classifiers from bootstrap samples
 - Bagging is an ensemble learning technique that is closely related to the Majority Vote Classifier, as illustrated in the following diagram



- Building an ensemble of classifiers from bootstrap samples
 - Instead of using the same training set to fit the individual classifiers in the ensemble, we draw bootstrap samples (random samples with replacement) from the initial training set.

Sample indices	Bagging round 1	Bagging round 2	
1	2	7	
2	2	3	
3	1	2	
4	3	1	
5	7	1	
6	2	7	
7	4	7	
	c_1	C ₂	c_m

Pros

- Improve the accuracy of unstable models
- Decrease the degree of overfitting

Loading Wine Dataset

```
import pandas as pd
df wine = pd.read csv('https://archive.ics.uci.edu/ml/'
                      'machine-learning-databases/wine/wine.data',
                      header=None)
df wine.columns = ['Class label', 'Alcohol', 'Malic acid', 'Ash',
                   'Alcalinity of ash', 'Magnesium', 'Total phenols',
                   'Flavanoids', 'Nonflavanoid phenols', 'Proanthocyanins',
                   'Color intensity', 'Hue', 'OD280/OD315 of diluted wines',
                   'Proline']
# drop 1 class
df wine = df wine[df wine['Class label'] != 1]
y = df wine['Class label'].values
X = df wine[['Alcohol', 'OD280/OD315 of diluted wines']].values
X.shape
(119, 2)
```

Loading Wine Dataset

Bagging

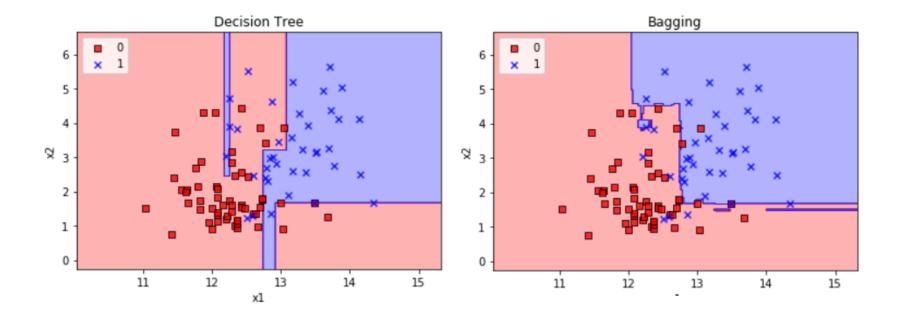
```
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import BaggingClassifier
# decision tree classifier
tree = DecisionTreeClassifier(criterion='entropy',
                                 max depth=4,
                                 random state=1)
tree.fit(X train, y train)
# bag of trees classifier
bag = BaggingClassifier(base estimator=tree,
                           n estimators=100,
                           max samples=0.5,
                           max features=1.0,
                           bootstrap=True,
                           bootstrap features=False,
                           n jobs=1,
                           random state=1)
                                                        Tree training/test accuracy: 0.92 / 0.79
bag.fit(X_train, y train)
                                                        Bag training/test accuracy: 0.92 / 0.88
print('Tree training/test accuracy: %.2f / %.2f'
     % (tree.score(X train, y train), tree.score(X test, y test)))
print('Bag training/test accuracy: %.2f / %.2f'
     % (bag.score(X_train, y_train), bag.score(X_test, y_test)))
```

Bagging

```
# decision boundary of the tree
plot_decision_regions(X_train, y_train, classifier=tree)
plt.xlabel('x1')
plt.ylabel('x2')
plt.title("Decision Tree")
plt.legend(loc='upper left')
plt.show()
```

```
# decision boundary of the bag of trees classifier
plot_decision_regions(X_train, y_train, classifier=bag)
plt.xlabel('x1')
plt.ylabel('x2')
plt.title("Bagging")
plt.legend(loc='upper left')
plt.show()
```

Bagging



- Leveraging weak learners via boosting
 - In contrast to bagging, the boosting algorithm learns simple classifiers k_j sequentially, and the final classifier is the weighted combination of k_j

$$C_m(x_i) = \alpha_1 k_1(x_i) + \alpha_2 k_2(x_i) + \dots + \alpha_m k_m(x_i)$$

- The key concept behind boosting is to focus on training samples that are hard to classify, that is, to let the weak learners subsequently learn from misclassified training samples to improve the performance of the ensemble
- Example
 - 1. Draw random subset of training samples d_1 d without replacement from the training set D to train a weak learner C_1
 - 2. Draw second random training subset d_2 without replacement from the training set and add 50 percent of the samples that were previously misclassified to train a weak learner C_2
 - 3. Find the training samples d_3 in the training set D on which C_1 and C_2 disagree to train a third weak learner C_3
 - 4. Combine the weak learners C_1 , C_2 , and C_3 via majority voting



Boosting

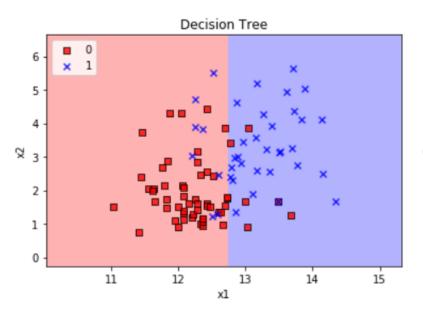
```
from sklearn.ensemble import AdaBoostClassifier
# decision tree classifier
tree = DecisionTreeClassifier(criterion='entropy',
                                 max depth=1,
                                 random state=1)
tree.fit(X train, y train)
# boosting classifier
ada = AdaBoostClassifier(base estimator=tree,
                                 n estimators=100,
                                 learning rate=0.1,
                                 random state=1)
ada.fit(X train, y train)
print('Tree training/test accuracy: %.2f / %.2f'
     % (tree.score(X train, y train), tree.score(X test, y test)))
print('AdaBoost training/test accuracy: %.2f / %.2f'
     % (ada.score(X train, y train), ada.score(X test, y test)))
 Tree training/test accuracy: 0.85 / 0.75
 AdaBoost training/test accuracy: 0.93 / 0.88
```

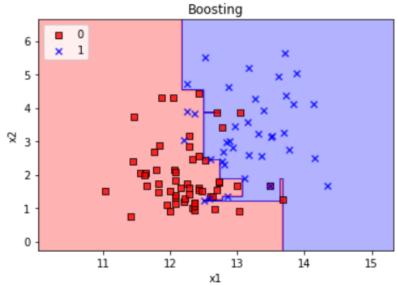
Boosting

```
# decision boundary of the tree
plot_decision_regions(X_train, y_train, classifier=tree)
plt.xlabel('x1')
plt.ylabel('x2')
plt.title("Decision Tree")
plt.legend(loc='upper left')
plt.show()
```

```
# decision boundary of the boosting classifier
plot_decision_regions(X_train, y_train, classifier=bag)
plt.xlabel('x1')
plt.ylabel('x2')
plt.title("Boosting")
plt.legend(loc='upper left')
plt.show()
```

Boosting





Submit

- To make sure if you have completed this practice, Submit your practice file(Week09 givencode.ipynb) to e-class.
- Deadline : tomorrow 11:59pm
- Modify your ipynb file name as "Week09_StudentNum_Name.ipynb"
 Ex) Week09_2020123456_홍일동.ipynb
- You can upload this file without taking the quiz, but homework will be provided like a quiz every three weeks, so it is recommended to take the quiz as well.

Quiz 1: Performance Evaluation

- Fine-tune your Logistic Regression model you build last week quiz
 - On the F1-Score,
 - Do K-fold Cross Validation
 - Plot Learning curve
 - Plot Validation curve
 - Find optimal hyperparameter using Grid Search



Quiz 2: Ensemble Methods

Build a Bagging Classifier using decision tree classifiers

 Build a AdaBoosting Classifier using decision tree classifiers