

EE2211 Tutorial 10

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- a) True
- b) False

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We have 3 parameter candidates for a classification model, and we would like to choose the optimal one for deployment. As such, we run 5-fold cross-validation.

Once we have completed the 5-fold cross-validation, in total, we have trained classifiers. Note that, we treat models with different parameters as different classifiers.

- A) 10
- B) 20
- C) 25
- D) 15

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Once we have completed the 5-fold cross-validation, in total, we have trained _____ classifiers. Note that, we treat models with different parameters as different classifiers.

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- C) 25
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In each fold, we train 3 classifiers, so 5 folds give 15 classifiers.

Suppose the binary classification problem, which you are dealing with, has highly imbalanced classes. The majority class has 99 hundred samples and the minority class has 1 hundred samples. Which of the following metric(s) would you choose for assessing the classification performance?

- a) Classification Accuracy
- b) Cost sensitive accuracy
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	$\widehat{\mathbf{P}}$ (predicted)	$\widehat{\mathbf{N}}$ (predicted)	
P (actual)	TP	FN	Recall TP/(TP+FN)
N (actual)	FP	TN	
	Precision TP/(TP+FP)	(TP+TN	Accuracy I)/(TP+TN+FP+FN

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The goal is to highlight the problems of the results!

In this case, we shall

- 1) Use cost matrix, assign different costs for each entry
- 2) Use Precision and Recall! Precision = 0.5 and Recall = 0.5

Given below is a scenario for Training error rate Tr, and Validation error rate Va for a machine learning algorithm. You want to choose a hyperparameter (P) based on Tr and Va. Which value of P will you choose based on the above table?

- a) 10
- b) 9
- c) 8
- d) 7
- e) 6

D	T _a	V/o
Р	Tr	Va
10	0.10	0.25
9	0.30	0.35
8	0.22	0.15
7	0.15	0.25
6	0.18	0.15

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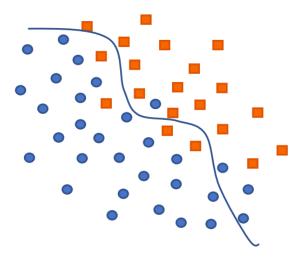
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(Binary and Multicategory Confusion Matrices)

Tabulate the confusion matrices for the following classification problems.

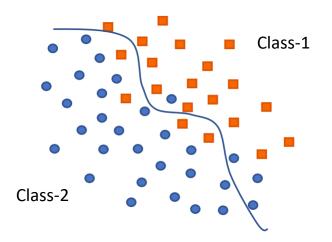
a) Binary problem (the class-1 and class-2 data points are respectively indicated by squares and circles)



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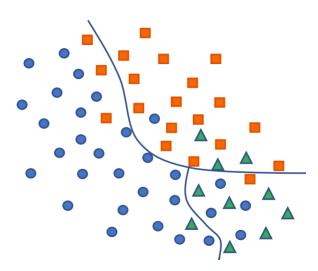
 a) Binary problem (the class-1 and class-2 data points are respectively indicated by squares and circles)



	$P_{\widehat{1}}$	$P_{\widehat{2}}$
P_1	16	4
P_2	4	26

Tabulate the confusion matrices for the following classification problems.

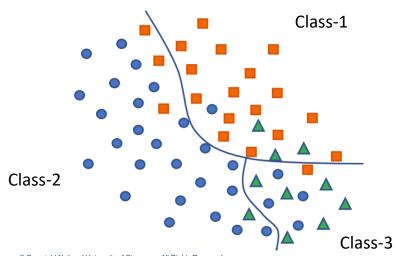
b) Three-category problem (the class-1, class-2 and class-3 data points are respectively indicated by squares, circles and triangles)



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Tabulate the confusion matrices for the following classification problems.

b) Three-category problem (the class-1, class-2 and class-3 data points are respectively indicated by squares, circles and triangles)



	$P_{\widehat{1}}$	$P_{\widehat{2}}$	$P_{\widehat{3}}$
P_1	16	3	1
P_2	1	25	4
P_3	3	1	6

Q6 (python)

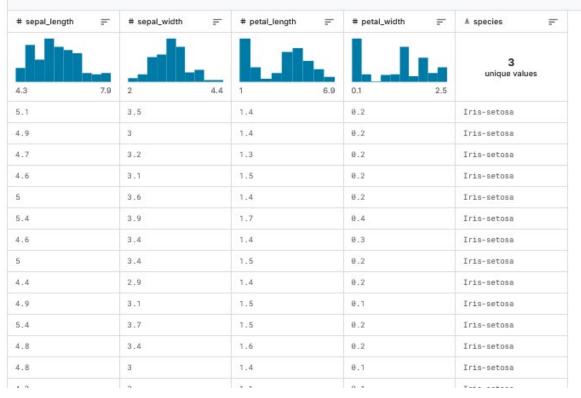
(5-fold Cross-Validation)

Get the data set "from sklearn.datasets import load_iris". Perform a 5-fold Cross-validation to observe the best polynomial order (among orders 1 to 10 and without regularization) for validation prediction. Note that, you will have to partition the whole dataset for training/validation/test parts, where the size of validation set is the same as that of test. Provide a plot of the average 5-fold training and validation error rates over the polynomial orders. The randomly partitioned data sets of the 5-fold shall be maintained for reuse in evaluation of future algorithms

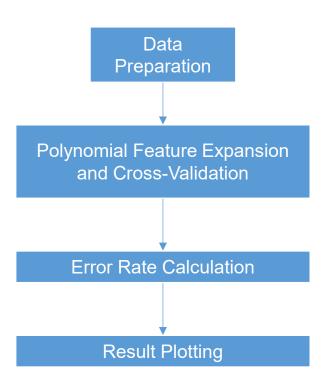


About this file

The dataset is a CSV file which contains a set of 150 records under 5 attributes - Petal Length, Petal Width, Sepal Length, Sepal width and Class(Species)

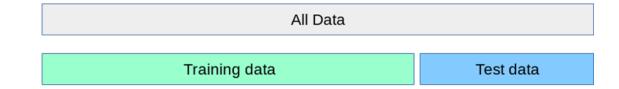


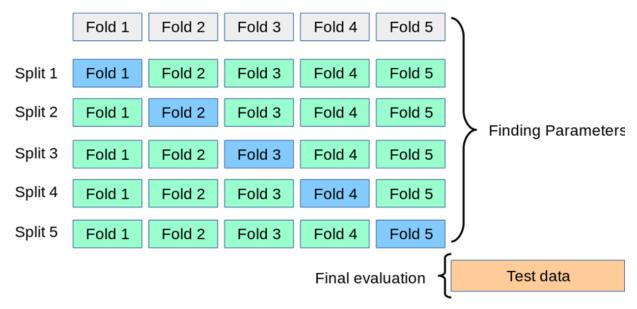
Block diagram



- One-Hot Encoding
- Data Splitting

- 5-Fold Splitting
- Feature Expansion
- Least-Squares Solution





https://scikit-learn.org/stable/modules/cross validation.html

```
##--- load data from scikit ---##
import numpy as np
import pandas as pd
print("pandas version: {}".format(pd.__version__))
import sklearn
print("scikit-learn version: {}".format(sklearn.__version__))
from sklearn.datasets import load iris

    Loads the Iris dataset, a classic dataset with three classes of flowers (Setosa,

iris dataset = load iris()
                                                      Versicolor, and Virginica).
X = np.array(iris dataset['data'])
                                                    • X: A NumPy array of shape (150, 4), containing four features for each sample.
y = np.array(iris_dataset['target'])
                                                    • y: A NumPy array of shape (150,), representing class labels (0, 1, or 2).
## one-hot encoding
Y = list()
for i in y:

    One-Hot Encoding: The target variable, y, is converted to a one-hot encoding format (Y) for

    letter = [0, 0, 0]
                                       each class (e.g., [1,0,0] for class 0, [0,1,0] for class 1, etc.).
    letter[i] = 1
    Y.append(letter)

    Y is now a NumPy array of shape (150, 3)

Y = np.array(Y)
test Idx = np.random.RandomState(seed=2).permutation(Y.shape[0])
X \text{ test} = X[\text{test } Idx[:25]]

    Uses a fixed seed (2) to create a reproducible random permutation of indices for splitting data.

Y_test = Y[test_Idx[:25]]

    Selects the first 25 samples for X test and Y test, leaving the remaining 125 samples in X and

X = X[test Idx[25:]]
                                        Y for training and validation.
Y = Y[test Idx[25:]]
```

```
from sklearn.preprocessing import PolynomialFeatures
error_rate_train_array = []
error rate val array = []
                                                      The code performs polynomial classification by expanding
##--- Loop for Polynomial orders 1 to 10 ---##
                                                       features to polynomial forms of varying degrees (1 to 10).
for order in range(1,11):
                                                      For each polynomial order, it uses 5-fold cross-validation:
    error rate train array fold = []
    error rate val array fold = []
   # Random permutation of data
                                                                   Creates a new random permutation of indices
    Idx = np.random.RandomState(seed=8).permutation(Y.shape[0])
                                                                   to assign data for 5-fold cross-validation.
   # Loop 5 times for 5-fold
   for k in range(0,5):
        ##--- Prepare training, validation, and test data for the 5-fold ---#
        # Prepare indexing for each fold
       X \text{ val} = X[Idx[k*25:(k+1)*25]]
                                                         Divides data into training and validation sets for each fold:
        Y_{val} = Y[Idx[k*25:(k+1)*25]]

    X val and Y val: Select the next 25 samples for validation.

        Idxtrn = np.setdiff1d(Idx, Idx[k*25:(k+1)*25])
                                                         • X train and Y train: Exclude the validation indices, using
        X train = X[Idxtrn]
                                                            the remaining samples for training.
        Y train = Y[Idxtrn]
```

```
##--- Polynomial Classification ---##
poly = PolynomialFeatures(order)
P = poly.fit transform(X train)
Pval = poly.fit transform(X val)
if P.shape[0] > P.shape[1]: # over-/under-determined cases
    reg L = 0.00*np.identity(P.shape[1])
    inv_PTP = np.linalg.inv(P.transpose().dot(P)+reg_L)
    pinv L = inv PTP.dot(P.transpose())
    wp = pinv L.dot(Y train)
else:
    reg_R = 0.00*np.identity(P.shape[0])
    inv PPT = np.linalg.inv(P.dot(P.transpose())+reg_R)
    pinv R = P.transpose().dot(inv PPT)
    wp = pinv R.dot(Y train)
```

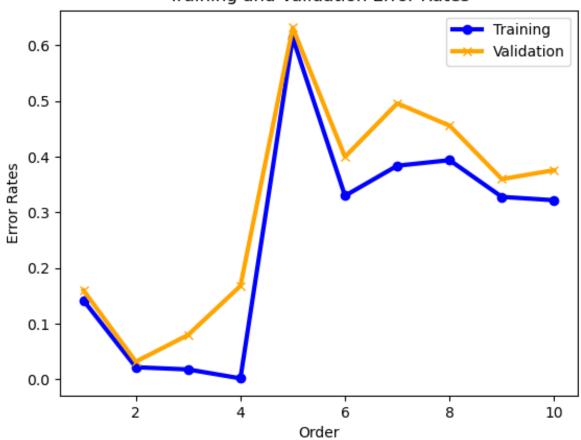
- Creates polynomial features of the specified order for X train and X val.
- P and Pval are the transformed features for training and validation, respectively.

- Checks if the system is over- or underdetermined (more rows than columns).
- Adds a small regularization term (reg_L or reg_R) for numerical stability in the pseudoinverse calculation.
- Least-Squares Solution: It calculates weights wp to fit the model by solving a system of equations based on whether the system is overdetermined or underdetermined.

```
##--- trained output ---##
    y_est_p = P.dot(wp);
    y_{cls_p} = [[1 \text{ if } y == max(x) \text{ else } 0 \text{ for } y \text{ in } x] \text{ for } x \text{ in } y_{est_p}]
    m1tr = np.matrix(Y train)
    m2tr = np.matrix(y cls p)
    # training classification error count and rate computation
    difference = np.abs(m1tr - m2tr)
    error_train = np.where(difference.any(axis=1))[0]
    error_rate_train = len(error_train)/len(difference)
    error rate train array fold += [error rate train]
    ##--- validation output ---##
    yval est p = Pval.dot(wp);
    yval cls p = [[1 \text{ if } y == max(x) \text{ else } 0 \text{ for } y \text{ in } x] \text{ for } x \text{ in } yval \text{ est } p]
    m1 = np.matrix(Y val)
    m2 = np.matrix(yval cls p)
    # validation classification error count and rate computation
    difference = np.abs(m1 - m2)
    error val = np.where(difference.any(axis=1))[0]
    error rate val = len(error val)/len(difference)
    error_rate_val_array_fold += [error_rate_val]
# store results for each polynomial order
error rate train array += [np.mean(error rate train array fold)]
error_rate_val_array += [np.mean(error_rate_val_array_fold)]
```

- y_est_p: Predicts continuous outputs by applying wp to the training data.
- y_cls_p: Converts y_est_p to a binary onehot format for classification (1 for the maximum value, 0 elsewhere).
- m1tr and m2tr represent the true and predicted one-hot encoded labels as matrices for easy comparison.
- Computes the training error rate by comparing y_cls_p to Y_train, identifying misclassified samples in each fold.
- Applies the same classification process to the validation set.
- Appends the validation error rate for this fold to error_rate_val_array_fo
- Stores the average training and validation error rates across the 5 folds for each polynomial order.

Training and Validation Error Rates



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THANK YOU