Programming for Data Science and Artificial Intelligence

Supervised Learning - Classification - K-Nearest Neighbors

Readings:

- [VANDER] Ch5
- [HASTIE] Ch9, 13

```
In [1]: Name = "Muhammad Omer Farooq Bhatti"
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In [1]: import matplotlib.pyplot as plt
    import numpy as np
    from sklearn.datasets import make_blobs
    from sklearn.model_selection import train_test_split
    from sklearn.preprocessing import StandardScaler
    from sklearn.metrics import average_precision_score, classification_report
    from sklearn.preprocessing import label_binarize
```

K-Nearest Neighbors

The intuition behind the KNN algorithm is one of the simplest of all the supervised machine learning algorithms. It simply calculates the distance of a new data point to all other training data points. The distance can be of any type e.g Euclidean or Manhattan etc. It then selects the K-nearest data points, where K can be any integer. Finally it assigns the data point to the class to which the majority of the K data points belong.

For example, given the red cross X, it simply get the majority class of neighbors, and assign to its own.

Scratch

Implementation steps:

- 1. Prepare your data
 - \$\mathbf{X}\$ and \$\mathbf{y}\$ in the right shape
 - \$\mathbf{X}\$ -> \$(m, n)\$
 - \$\mathbf{y}\$ -> \$(m,)\$
 - Why no w?
 - train-test split
 - feature scale
 - clean out any missing data
 - (optional) feature engineering
- 2. Write a function for computing pairwise distance between every points
- 3. Then, given set of X_test data, compute their distance to all other points, then argsort the distance matrix, and get the k-nearest indices
- 4. Get the majority class
- 1. Prepare your data
- 2. Function for pairwise distance

I have written three different ways in numpy assignment answer question 11

- 3. Argsort the pairwise distance matrix
- 4. Get the majority class

When to use KNN

I guess the only good thing about it is that KNN is super easy to implement, and generally work quite well on simple classification problems. However, it also comes with a price:

- Computational expense as feature grows, since it requires computing the distance for each feature, where for each feature, we have to compute the input points with every single points, then perform sort (which can be expensive), and then get the majority class from the nearest nth-neighbors. Very expensive!
- Can't work with categorical features since it is difficult to formulate distance formulas for categorial features
- Of course, it takes even more time to find the right n_neighbors (or commonly known as k)

```
===Task===
```

Your work: Let's modify the above scratch code to

• If the majority class of the first place is equal to the second place, then ask the algorithm to pick the next nearest neighbors as the

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decider

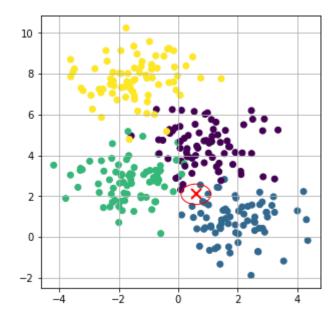
- Modify the code so it outputs the probability of the decision, where the probability is simply the class probability based on all the nearest neighbors
- Write a function which allows the program to receive a range of k, and output the cross validation score. Last, it shall inform us which k is the best to use from a predefined range
- Put everything into a class KNN(k=3) . It should have at least one method, predict(X_train, X_test, y_train)

```
In [2]:
        class KNN():
            def _init__(self, k=3):
                self.k=k
                self.X train=None
                self.X test=None
                self.y train=None
            def predict(self, X_train, X_test, y_train):
                self.X train=X train
                self.X test=X test
                self.y_train=y_train
                num_class=len(set(self.y_train))
                neighbours_idx = self.find_all_neighbours()
                prediction=np.zeros(X_test.shape[0])
                probability=np.zeros(X_test.shape[0])
                for idx, y in enumerate(self.y_train[neighbours_idx]):
                     prediction[idx], probability[idx] = self.get most common(y, num class)
                return prediction, probability
            def find k neighbours(self):
                 squared distance = np.square(self.X test[:,np.newaxis,:]-self.X train[np.newaxis,:,:])
                euclidian distance = np.sqrt(np.sum(squared distance, axis=2))
                neighbours idx = np.argsort(euclidian distance)
                return neighbours idx[:,:self.k]
            def find all neighbours(self):
                 squared distance = np.square(self.X test[:,np.newaxis,:]-self.X train[np.newaxis,:,:])
                euclidian distance = np.sqrt(np.sum(squared distance, axis=2))
                neighbours idx = np.argsort(euclidian distance)
                return neighbours idx
            def get_most_common(self, y, num_class):
                k \text{ nearest } y = y[:self.k]
                y count = np.bincount(k nearest y, minlength=num class)
                most_common_y = y_count.argmax()
                second_most_common_y = y_count.argsort()[-2]
                 #print(f"k nearest y: {k nearest y}")
                 #print(f"bincount: {y_count}")
                 #print(f"num classes: {num class}")
                 #print(f"most_common_y: {most_common_y}")
                 #print(f"second_most_common_y: {second_most_common_y}")
                 #print(f"y_count[most_common_y]: {y_count[most_common_y]}")
                 #print(f"y_count[second_most_common_y]: {y_count[second_most_common_y]}")
                if (y_count[most_common_y] == y_count[second_most_common_y]):
                     k_nearest_y = y[:self.k+1] ## add another y element/neighbour
                y_count = np.bincount(k_nearest_y, minlength=num_class)
                yhat = y count.argsort()[-1]
                p_yhat = y_count[yhat]/np.sum(y count)
                return yhat, p_yhat
            def k fold cross validation(self, X train, y train, k range, k=10 ):
                 \#implementing k fold cross validation for determining the best value of hyperparameter k=no. of neighborst
                 #We divide the training set into k parts. We withhold one part for testing and use the rest for trai
                 #Loop through all the folds, by keeping each one separate as a testing set and using the rest for tr
                 fold size = int(X train.shape[0]/k)
                 accuracy = {}
                 for k in k range:
                    self.k = k
                    accuracy[self.k]=[]
                    for i in range(0, X_train.shape[0], fold_size):
                        xtest = X_train[i:i+fold_size]
                         ytest = y_train[i:i+fold_size]
                        xtrain = np.concatenate((X_train[:i], X_train[i+fold_size:]), axis=0)
                         ytrain = np.concatenate((y_train[:i], y_train[i+fold_size:]), axis=0)
                         yhat, p_yhat = self.predict(xtrain, xtest, ytrain)
                         accuracy[self.k].append( np.sum(yhat==ytest)/ytest.shape[0] )
                 return accuracy
```

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```
X, y = make blobs(n samples=300, centers=4,
                  random_state=0, cluster_std=1.0)
xfit = np.linspace(-1, 3.5)
figure = plt.figure(figsize=(5, 5))
ax = plt.axes() #get the instance of axes from plt
ax.grid()
ax.scatter(X[:, 0], X[:, 1], c=y)
#where should this value be classified as?
ax.plot([0.6], [2.1], 'x', color='red', markeredgewidth=2, markersize=10)
#let's say roughly 5 neighbors
circle = plt.Circle((0.6, 2.1), 0.5, color='red', fill=False)
ax.add artist(circle)
```

<matplotlib.patches.Circle at 0x1ad19c91fd0> Out[3]:



```
In [4]:
        #standardize
        scaler = StandardScaler()
        X = scaler.fit_transform(X)
        #do train test split
        X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.3)
```

```
In [5]:
        model = KNN()
        yhat, yp = model.predict(X_train, X_test, y_train)
        print(yhat)
        print(f"Probability of yhat: {yp}")
```

```
[3. 2. 1. 1. 3. 3. 0. 1. 0. 0. 2. 1. 1. 2. 3. 0. 0. 1. 2. 3. 3. 2. 2. 1.
3. 0. 2. 3. 1. 3. 3. 0. 1. 0. 3. 0. 2. 0. 2. 0. 2. 0. 1. 1. 3. 0. 3. 2.
1. 1. 0. 3. 2. 3. 2. 0. 3. 3. 1. 0. 3. 0. 2. 2. 1. 0. 3. 3. 3. 0. 3. 2.
1. 0. 2. 1. 3. 0. 2. 0. 1. 1. 0. 3. 2. 1. 2. 3. 0. 1.]
Probability of yhat: [1.
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```

```
In [6]:
        n_classes = len(np.unique(y_test))
        print("Accuracy: ", np.sum(yhat == y_test)/len(y_test))
        print("============")
        y_test_binarized = label_binarize(y_test, classes=[0, 1, 2, 3])
        yhat binarized = label_binarize(yhat, classes=[0, 1, 2, 3])
        for i in range(n_classes):
           class_score = average_precision_score(y_test_binarized[:, i], yhat_binarized[:, i])
           print(f"Class {i} score: ", class score)
        print("======Classification report======")
       print("Report: ", classification_report(y_test, yhat))
```

Accuracy: 0.866666666666667 ======Average precision score===== Class 0 score: 0.66833333333333333 Class 1 score: 0.9352657004830918

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```
Class 2 score: 0.7558333333333332
       Class 3 score: 0.8129292929293
       ======Classification report======
                             precision
       Report:
                                        recall f1-score support
                       0.790.760.781.000.910.950.850.850.850.840.950.89
                  0
                                                        25
                                                         23
                  1
                                                         20
                  2
                                                        22
                                           0.87
                                                        90
           accuracy
                                          0.87
                     0.87 0.87
                                                         90
          macro avg
                                           0.87
                                                         90
       weighted avg
                        0.87
                                  0.87
In [8]:
        k_{range} = [3, 4, 5, 6, 7, 8, 9, 10]
        accuracy = model.k fold cross validation(X train, y train, k range, k=5)
        acc=np.zeros((len(k range),2))
        for idx, key in enumerate(k range):
            accuracy[key] = np.mean(accuracy[key])
           print(f"Mean accuracy for k={key}: {accuracy[key]}")
           acc[idx,0] = key
            acc[idx,1] = accuracy[key]
        #print(acc)
        print(f"The max accuracy achieved was \{acc[acc.argmax(axis=0)[1],1]\}\ for k = \{acc[acc.argmax(axis=0)[1],0]\}"
       Mean accuracy for k=3: 0.9523809523809523
       Mean accuracy for k=4: 0.9476190476190475
       Mean accuracy for k=5: 0.9523809523809522
       Mean accuracy for k=6: 0.9523809523809522
       Mean accuracy for k=7: 0.9523809523809522
       Mean accuracy for k=8: 0.9523809523809522
       Mean accuracy for k=9: 0.9523809523809522
       Mean accuracy for k=10: 0.9571428571428571
```

In []:

The max accuracy achieved was 0.9571428571428571 for k = 10.0

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