Implement RandomSearchCV with k fold cross validation on KNN

```
In [1]:
```

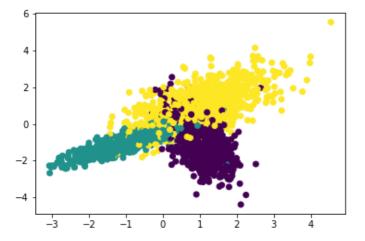
```
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
import numpy
from tqdm import tqdm
import numpy as np
from sklearn.metrics.pairwise import euclidean_distances

x,y = make_classification(n_samples=10000, n_features=2,n_classes=3, n_informative=2, n_
redundant= 0, n_clusters_per_class=1, random_state=60)
X_train, X_test, y_train, y_test = train_test_split(x,y,stratify=y,random_state=42)

# del X_train,X_test
```

In [2]:

```
%matplotlib inline
import matplotlib.pyplot as plt
#colors = {0:'orange', 1:'blue'}
plt.scatter(X_test[:,0], X_test[:,1],c=y_test)
plt.show()
```



Implementing Custom GridSearchCV

```
In [8]:
```

```
# it will take classifier and set of values for hyper prameter in dict type dict({hyper p
armeter: [list of values]})
# we are implementing this only for KNN, the hyper parameter should n neighbors
from sklearn.metrics import accuracy score
def randomly select 60 percent indices in range from 1 to len(x train):
   return random.sample(range(0, len(x train)), int(0.6*len(x train)))
def GridSearch(x train, y train, classifier, params, folds):
   trainscores = []
   testscores = []
    for k in tqdm(params['n neighbors']):
       trainscores folds = []
       testscores folds = []
        for j in range(0, folds):
            # check this out: https://stackoverflow.com/a/9755548/4084039
            train indices = randomly select 60 percent indices in range from 1 to len(x t
rain)
            test indices = list(set(list(range(1, len(x train)))) - set(train indices))
```

```
# selecting the data points based on the train_indices and test_indices
X_train = x_train[train_indices]
Y_train = y_train[train_indices]
X_test = x_train[test_indices]
Y_test = y_train[test_indices]

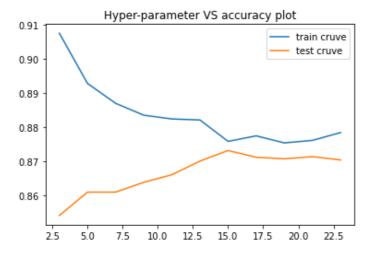
classifier.n_neighbors = k
classifier.fit(X_train,Y_train)

Y_predicted = classifier.predict(X_test)
testscores_folds.append(accuracy_score(Y_test, Y_predicted))

Y_predicted = classifier.predict(X_train)
trainscores_folds.append(accuracy_score(Y_train, Y_predicted))
trainscores.append(np.mean(np.array(trainscores_folds)))
testscores.append(np.mean(np.array(testscores_folds)))
return trainscores, testscores
```

In [9]:

```
from sklearn.metrics import accuracy score
from sklearn.neighbors import KNeighborsClassifier
import matplotlib.pyplot as plt
import random
import warnings
warnings.filterwarnings("ignore")
neigh = KNeighborsClassifier()
params = {'n neighbors':[3,5,7,9,11,13,15,17,19,21,23]}
folds = 3
trainscores, testscores = GridSearch(X train, y train, neigh, params, folds)
plt.plot(params['n neighbors'], trainscores, label='train cruve')
plt.plot(params['n neighbors'], testscores, label='test cruve')
plt.title('Hyper-parameter VS accuracy plot')
plt.legend()
plt.show()
             | 11/11 [00:04<00:00,
100%|
                                      2.51it/s]
```



In [5]:

```
# understanding this code line by line is not that importent
def plot_decision_boundary(X1, X2, y, clf):
    # Create color maps
    cmap_light = ListedColormap(['#FFAAAA', '#AAFFAA', '#AAAAFF'])
    cmap_bold = ListedColormap(['#FF0000', '#00FF00', '#0000FF'])

x_min, x_max = X1.min() - 1, X1.max() + 1
    y_min, y_max = X2.min() - 1, X2.max() + 1
```

```
xx, yy = np.meshgrid(np.arange(x_min, x_max, 0.02), np.arange(y_min, y_max, 0.02))
Z = clf.predict(np.c [xx.ravel(), yy.ravel()])
Z = Z.reshape(xx.shape)
plt.figure()
plt.pcolormesh(xx, yy, Z, cmap=cmap light)
# Plot also the training points
plt.scatter(X1, X2, c=y, cmap=cmap bold)
plt.xlim(xx.min(), xx.max())
plt.ylim(yy.min(), yy.max())
plt.title("2-Class classification (k = %i)" % (clf.n neighbors))
plt.show()
```

In [6]:

```
from matplotlib.colors import ListedColormap
neigh = KNeighborsClassifier(n neighbors = 21)
neigh.fit(X train, y train)
plot decision boundary(X train[:, 0], X train[:, 1], y train, neigh)
```

2-Class classification (k = 21) 2 0

In []:

```
from sklearn.metrics import accuracy score
def randomized search cv custom (x train total, y train total, classifier, param range, nu
m of total fold):
    # x train total: its numpy array of shape, (n,d)
    # y train total: its numpy array of shape, (n,) or (n,1)
    # classifier: its typically KNeighborsClassifier()
    # param range: Integer representing how many hyper-parameters I am considering for ea
ch iteration
    # num of total fold: an integer, represents number of num of total fold we need to de
vide the data and test our model
    # generating hyper-parameter range
    # generate 10 unique values(uniform random distribution) in the given range - startin
g 1 to "param range"
    # ex: if param range = 50, we need to generate 10 random numbers in range 1 to 50
    ten random values for param range = sorted(random.sample(range(1, param range), 10))
    train scores = []
    test scores = []
    classifier params = { 'n neighbors': ten random values for param range }
    # it will take classifier and set of values for hyper parameters in dict type
    # dict({hyper parmeter: [list of values]})
    # as we are implementing this only for KNN, the hyper parameter should be n neighbors
    # And I will return it from the function at the end so that I can use this same hyper
-param
    # while plotting the graph as well
    for k in tqdm(classifier params['n neighbors']):
```

```
trainscores_folds = []
        testscores_folds = []
        for fold in range(0, num_of_total_fold):
            \# divide numbers ranging from 0 to len(x train fold) into groups = num of t
otal fold
            # basically, splitting the data into k groups (k = len(x train fold) / num old)
f total fold)
            # It works by first training the algorithm on the k 1 group of the data and
            # evaluating it on the kth hold-out group as the test set. This is repeated
            # so that each of the k groups is given an opportunity to be held out and use
d as the test set.
            # ex: num of total fold=3, and len(x train total)=100, we can divide numbers
from 0 to 100 into 3 groups
            # group 1: 0-33, group 2:34-66, group 3: 67-100
            num of elements in each fold = int(len(x train total) / num of total fold)
            # for each hyperparameter that we generated in step 1:
            # and using the above groups we have created in step 2 you will do cross-vali
dation as follows
            # first we will keep group 1+group 2 i.e. 0-66 as train data and
            # group 3: 67-100 as test data, and find train and test accuracies
            # second we will keep group 1+group 3 i.e. 0-33, 67-100 as train data and
            # group 2: 34-66 as test data, and find train and test accuracies
            # third we will keep group 2+group 3 i.e. 34-100 as train data and
            # group 1: 0-33 as test data, and find train and test accuracies
            # based on the 'num of total fold' value we will do the same procedure
            # NOW IMPLEMENTATION OF THE CONCEPT OF 'fold' as below
            # For each of this inner loop running for values of fold (where fold represen
ts num of total fold of 0, 1, 2, 3 ...)
            # each of the test indices will have the data of a single fold ( which is =
num of elements in each fold )
            # i.e. the test indices will be the range starting at
            # num of elements in each fold * fold and ending at num of elements in each
fold * (fold + 1)
            # And this whole range needs to be converted to list => then apply set() to t
he list
            # => and then again converted to list
            test indices = list(set(list(range((num of elements in each fold * fold), (n
um of elements in each fold*(fold+1))))))
            # print('test indices ', test indices)
            # And the rest of the indices of the dataset will be the train indices
            train indices = list(set(list(range(0, len(x train total)))) - set(test indi
ces) )
            # print('train indices ', train indices)
            ''' So for a dataset of total 100 rows, one loop of fold, will have the follo
wing test indices and train indices
            test indices [0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17,
18, 19, 20, 21, 22, 23, 24]
            train_indices [25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 4
0, 41, 42, 43, 4<del>4</del>, 45, 46, 47, 48, 49, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, 68, 69, 70, 71, 72, 73, 74]
           And the next loop of j will have as below (for the same dataset of total 100
rows )
            test indices [25, 26, 27, 28, 29, 30, 31, 32, 33, 34, 35, 36, 37, 38, 39, 40
, 41, 42, 43, 44, 45, 46, 47, 48, 49]
            train_indices [0, 1, 2, 3, 4, 5, 6, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17,
18, 19, 20, 21, 22, 23, 24, 50, 51, 52, 53, 54, 55, 56, 57, 58, 59, 60, 61, 62, 63, 64, 6
5, 66, 67, 68, 69, 70, 71, 72, 73, 74]
            # after we have above, now select datapoints based on test indices and train
```

indices

```
x_train_fold = x_train_total[train_indices]
            y_train_fold = y_train_total[train_indices]
            x test fold = x train total[test indices]
            y test fold = y train total[test indices]
            # Now based on our classifier assign corresponding parameter values
            # and also fit() and predict()
            classifier.n neighbors = k
            classifier.fit(x train fold, y train fold)
            # First predict based on x test fold and keep the accuracy score in the tests
cores_folds
            y predicted = classifier.predict(x test fold)
            testscores folds.append(accuracy score(y test fold, y predicted))
            # Now run prediction based on x train fold and append the accuracy score in t
he trainscores folds
            y predicted = classifier.predict(x train fold)
            trainscores folds.append(accuracy score(y train fold, y predicted))
        train scores.append(np.mean(np.array(trainscores folds)))
        test scores.append(np.mean(np.array(testscores folds)))
    return train scores, test scores, classifier params
```

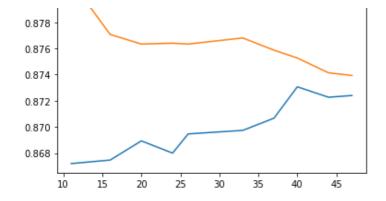
In [8]:

```
from sklearn.neighbors import KNeighborsClassifier
from matplotlib.colors import ListedColormap
import matplotlib.pyplot as plt
import warnings
warnings.filterwarnings("ignore")
# Our Classifier is KNN, hence assign a variable to it.
neigh = KNeighborsClassifier()
params range = 50
number of total folds = 3
# Now invoking our custom function randomized search cv custom(x train, y train, classifier
, param range, num of total fold) and store the returned values
testscores, trainscores, params = randomized_search_cv_custom(X_train, y_train, neigh, p
arams range, number of total folds)
print('trainscores are: ', trainscores)
print('testscores are: ', testscores)
      | 10/10 [00:04<00:00, 2.45it/s]
trainscores are: [0.860799999999999, 0.86773333333332, 0.87, 0.868666666666666, 0.86
7733333333334, 0.868933333333332, 0.8693333333334, 0.86920000000001, 0.8698666666
66667, 0.8730666666666668]
testscores are: [0.888933333333332, 0.880266666666666, 0.8802, 0.87693333333335, 0.8
776, 0.87633333333333, 0.876333333333333, 0.8773333333333, 0.8760666666666667, 0.87
52666666666661
```

Plotting hyper-parameter vs accuracy plot

```
In [10]:
```

```
# Now plotting hyper-parameter vs accuracy plot to choose the best hyperparameter
plt.plot(params['n_neighbors'], trainscores, label='train curve')
plt.plot(params['n_neighbors'], testscores, label='test curve')
plt.title('Hyper-parameter VS accuracy plot')
plt.legend()
plt.show()
```



Observation:

1-from the above curve we an conclude k value near to 30 will give good accuracy.

2-Because of taking random value of k, this graph is changing for each iteration

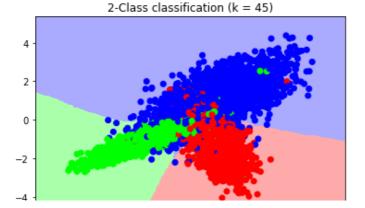
```
In [11]:
```

```
def plot decision boundary(X1, X2, y, clf):
    # Create color maps
    cmap light = ListedColormap(['#FFAAAA', '#AAFFAA', '#AAAAFF'])
    cmap_bold = ListedColormap(['#FF0000', '#00FF00', '#0000FF'])
    x \min, x \max = X1.\min() - 1, X1.\max() + 1
    y \min, y \max = X2.\min() - 1, X2.\max() + 1
   xx, yy = np.meshgrid(np.arange(x min, x max, 0.02), np.arange(y min, y max, 0.02))
    Z = clf.predict(np.c [xx.ravel(), yy.ravel()])
    Z = Z.reshape(xx.shape)
    plt.figure()
    plt.pcolormesh(xx, yy, Z, cmap=cmap light)
    # Plot also the training points
   plt.scatter(X1, X2, c=y, cmap=cmap bold)
   plt.xlim(xx.min(), xx.max())
    plt.ylim(yy.min(), yy.max())
    plt.title("2-Class classification (k = %i)" % (clf.n neighbors))
    plt.show()
```

Plotting the decision boundaries for the model initialized with the best hyperparameter

```
In [13]:
```

```
# plotting the decision boundaries for the model initialized with the best hyperparameter
(45)
neigh = KNeighborsClassifier(n_neighbors = 45)
neigh.fit(X_train, y_train)
plot_decision_boundary(X_train[:, 0], X_train[:, 1], y_train, neigh)
```



-4 -2 0 2 4

In []: