```
In [27]:
```

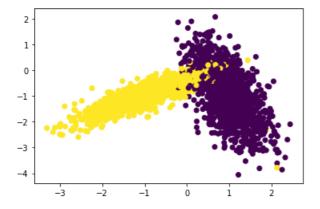
```
from sklearn.datasets import make_classification
from sklearn.model_selection import train_test_split
from sklearn.preprocessing import StandardScaler
import numpy
from tqdm import tqdm
import numpy as np
from sklearn.metrics.pairwise import euclidean_distances

x,y = make_classification(n_samples=10000, n_features=2, n_informative=2, n_redundant= 0, n_clusters_pe
r_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 [28]:

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



# Implementing Custom RandomSearchCV

```
def RandomSearchCV(x_train,y_train,classifier, param_range, folds):
    # x train: its numpy array of shape, (n,d)
    # y train: its numpy array of shape, (n,) or (n,1)
    # classifier: its typically KNeighborsClassifier()
    # param range: its a tuple like (a,b) a < b
    # folds: an integer, represents number of folds we need to devide the data and test our mode
    #1.generate 10 unique values(uniform random distribution) in the given range "param range" a
nd store them as "params"
    \# ex: if param_range = (1, 50), we need to generate 10 random numbers in range 1 to 50
    #2.devide numbers ranging from 0 to len(X train) into groups= folds
    # ex: folds=3, and len(x train)=100, we can devide numbers from 0 to 100 into 3 groups
     group 1: 0-33, group 2:34-66, group 3: 67-100
    #3.for each hyperparameter that we generated in step 1:
        # and using the above groups we have created in step 2 you will do cross-validation as f
ollows
        # 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 # based on the 'folds' value we will do the same procedure
- # find the mean of train accuracies of above 3 steps and store in a list "train\_scores"
- # find the mean of test accuracies of above 3 steps and store in a list "test scores"
- #4. return both "train scores" and "test scores"
- #5. call function RandomSearchCV(x\_train,y\_train,classifier, param\_range, folds) and store the r eturned values into "train score", and "cv scores"
- #6. plot hyper-parameter vs accuracy plot as shown in reference notebook and choose the best hyper-parameter
- #7. plot the decision boundaries for the model initialized with the best hyperparameter, as show n in the last cell of reference notebook

# In [29]:

```
#Select data for training. This function directly returns training data rather than indices
def selection of data for train(x train, k, folds):
   #Splits the data into (folds+1) parts.
   #[1:folds] considers the parts other than 0 and last value
   a1 = np.linspace(0,len(x train),folds + 1)[1:folds]
    #Slicing of data is done based on the iteration.
    #For eg. in 1st iteration, the 1st slice of data will be considered for testing and remaining data
for training.
    #For first iteration, training data is all the values except the first slice
   if k is 0:
       return x train[int(a1[k]):]
   #For last iteration, training data is all the values except the last slice
   elif k is (folds-1):
       return x train[:(int(a1[k-1]))]
    #during the remaining iterations the slice of iertaion number will be excluded
       return np.concatenate([x train[:int(a1[k-1])] , x train[int(a1[k]):]])
```

# In [30]:

```
#this function returns the data for testing based in the folds and iteration value

def selection_of_data_for_test(x_test,k,folds):
    #Splits the data into (folds+1) parts.
    #[1:folds] considers the parts other than 0 and last value
    al = np.linspace(0,len(x_test),folds + 1)[1:folds]

#for first iteration, only first slice will be considered for testing
    if k is 0:
        return x_test[:int(al[k])]

#for last iteration, only last slice will be considered for testing
    elif k is (folds-1):
        return x_test[(int(al[k-1])):]

#for nth iteration, ith slice will be considered
    else:
        return np.concatenate([x test[int(al[k-1]):int(al[k])]])
```

In [33]:

```
#implmentation of RandomSearchCV
def RandomSearchCV(x_train,y_train,classifier, param_range, folds):
   trainscore = []
    testscore = []
    for k in tqdm(param_range['n_neighbour']):
        trainscores folds = []
        testscores folds = []
        for j in range(folds):
             #selection of training and test data
            X train = selection of data for train(x train, j, folds)
            Y_train = selection_of_data_for_train(y_train,j,folds)
X_test = selection_of_data_for_test(x_train,j,folds)
            Y_test = selection_of_data_for_test(y_train,j,folds)
            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))
        trainscore.append(np.mean(np.array(trainscores folds)))
        testscore.append(np.mean(np.array(testscores folds)))
    return trainscore, testscore
```

#### In [38]:

```
from sklearn.metrics import accuracy score
from sklearn.neighbors import KNeighborsClassifier
import matplotlib.pyplot as plt
import random
import warnings
warnings.filterwarnings("ignore")
#KNN classifier object
neigh = KNeighborsClassifier()
#a random uniform distributed values between a and b are considered in sorted order
list1 = list(range(1,50))
params = {'n neighbour':sorted(random.sample(list1,10))}
print ("Parameters")
print (params)
folds = 3
trainscores, testscores = RandomSearchCV(X train, y train, neigh, params, folds)
print("Training scores")
print(trainscores)
plt.plot(params['n_neighbour'], trainscores, label='train cruve')
plt.plot(params['n_neighbour'], testscores, label='test cruve')
plt.title('Hyper-parameter VS accuracy plot')
plt.legend()
plt.show()
```

#### Parameters

```
{'n_neighbour': [4, 21, 23, 29, 30, 31, 33, 37, 45, 48]}
```

```
0% | | 0/10 [00:00<?, ?it/s]

10% | | 1/10 [00:00<00:07, 1.17it/s]

20% | | 2/10 [00:01<00:07, 1.09it/s]

30% | | 3/10 [00:03<00:07, 1.04s/it]

40% | | 4/10 [00:04<00:06, 1.07s/it]

50% | | 5/10 [00:05<00:05, 1.14s/it]

60% | | 6/10 [00:06<00:04, 1.15s/it]

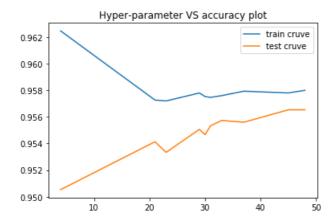
70% | | 7/10 [00:08<00:03, 1.18s/it]

80% | | 8/10 [00:09<00:02 1 24s/it]
```

```
90%| | 9/10 [00:10<00:01, 1.24s/it] | 10/10 [00:12<00:00, 1.33s/it]
```

Training scores

[0.96246666666667, 0.9572666666666666, 0.957199999999999, 0.95780000000001, 0.957533333333333, 0.95746666666666, 0.95759999999999, 0.95793333333333, 0.957800000000001, 0.9579999999999]



## In [40]:

```
# 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', '#AAAFF'])
cmap_bold = ListedColormap(['#FF0000', '#00FF00', '#000FF'])
    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 [41]:

```
from matplotlib.colors import ListedColormap
neigh = KNeighborsClassifier(n_neighbors = 48)
neigh.fit(X_train, y_train)
plot_decision_boundary(X_train[:, 0], X_train[:, 1], y_train, neigh)
```

