```
import math
from collections import defaultdict
import pandas as pd
import numpy as np
from sklearn.model_selection import train_test_split
import matplotlib.pyplot as plt
```

## 1. Evaluating the k-NN Algorithm

```
In [ ]: df = pd.read_csv("iris.csv", header=None)
        np.random.seed(0)
        #store results of iterations in dict
        test res = defaultdict(list)
        train res = defaultdict(list)
        test no norm = defaultdict(list)
        for iteration in range(20):
            #shuffle data
            ds = df.sample(frac=1)
            #split data into 80% train, 20% test
            train, test = train test split(ds,test size=0.2)
            train att = train[0,1,2,3]
            test att = test[[0,1,2,3]]
            #normalize data
            train norm = (train att - train att.min()) / (train att.max() - train att.min())
            test_norm = (test_att - train_att.min()) / (train_att.max() - train_att.min())
            train norm.fillna(0, inplace=True)
            test_norm.fillna(0, inplace=True)
            k - number of neighbors
            data - dataset
            instance - instance to classify
            def knn(k,data,instance):
                distances = data.apply(lambda row: math.dist(row,instance), axis=1)
                #get k closest instances (including the input instance)
                k_neighbors = distances.sort_values()[:k]
                #get class value with largest number of occurences
                 predicted class = df.loc[k neighbors.index,[4]][4].mode()
                return predicted class[0]
            train knn using training data
            return the accuracy of the model
            examples - specify training set
            def train knn(k,data,examples):
                predictions = data.apply(lambda row: knn(k,examples,row), axis=1)
                 actual labels = df.loc[predictions.index,[4]]
                 return predictions.eq(actual_labels[4].values).mean()
            for i in range(1,51,2):
                train_res[i].append(train_knn(i,train_norm,train_norm))
```

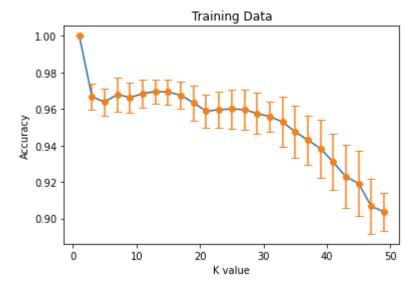
```
test_res[i].append(train_knn(i,test_norm,train_norm))
test_no_norm[i].append(train_knn(i,test_att,train_att))
```

**Q1.1 (10 Points)** In the first graph, you should show the value of k on the horizontal axis, and on the vertical axis, the average accuracy of models trained over the training set, given that particular value of k. Also show, for each point in the graph, the corresponding standard deviation; you should do this by adding error bars to each point. The graph should look like the one in Figure 2 (though the "shape" of the curve you obtain may be different, of course).

```
In []: train_iter = []
    train_acc = []
    train_std = []

for k_val,res in train_res.items():
        train_iter.append(k_val)
        train_acc.append(np.average(res))
        train_std.append(np.std(res))

plt.plot(train_iter,train_acc)
    plt.errorbar(train_iter,train_acc,yerr = train_std,fmt='o',capsize=4)
    plt.xlabel("K value")
    plt.ylabel("Accuracy")
    plt.title("Training Data")
    plt.show()
```

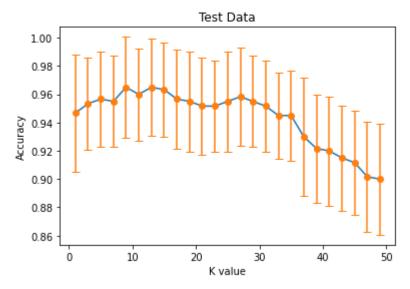


**Q1.2 (10 Points)** In the second graph, you should show the value of k on the horizontal axis, and on the vertical axis, the average accuracy of models trained over the testing set, given that particular value of k. Also show, for each point in the graph, the corresponding standard deviation by adding error bars to the point.

```
In [ ]: test_iter = []
    test_acc = []
    test_std = []

for k_val,res in test_res.items():
        test_iter.append(k_val)
        test_acc.append(np.average(res))
        test_std.append(np.std(res))
```

```
plt.plot(test_iter,test_acc)
plt.errorbar(test_iter,test_acc,yerr = test_std,fmt='o',capsize=4)
plt.xlabel("K value")
plt.ylabel("Accuracy")
plt.title("Test Data")
plt.show()
```



**Q1.3 (8 Points)** Explain intuitively why each of these curves look the way they do. First, analyze the graph showing performance on the training set as a function of k. Why do you think the graph looks like that? Next, analyze the graph showing performance on the testing set as a function of k. Why do you think the graph looks like that?

For the graph showing performance on the training set, for K=1 the accuracy is 1.00 due to the fact that we do not ignore the instance being tested in our KNN algorithm. This means that the algorithm will always group instances from the training sets with themselves. The std deviation for each point generally increases for increased values of K. This makes sense because as larger groups are formed, it is likelier that data points are incorrectly labeled. For the graph showing performance on the testing set, the accuracy maxes out at around 0.96 for K=9. There is a general decrease in accuracy as K increases but for low values of K there is slight growth. The std deviation for each point is roughly the same and is significantly larger than in the training set. This is likely due to the fact that both the training and testing sets are normalized according to the training set, and that the testing set is significantly smaller than the training set.

**Q1.4 (6 Points)** We say that a model is underfitting when it performs poorly on the training data (and most likely on the testing data as well). We say that a model is overfitting when it performs well on training data but it does not generalize to new instances. Identify and report the ranges of values of k for which k-NN is underfitting, and ranges of values of k for which k-NN is overfitting.

For K values from 1-15 the model is overfitting. For K values from 15-51, the model is still overfitting, but it becomes increasingly underfitting.

**Q1.5 (6 Points)** Based on the analyses made in the previous question, which value of k you would select if you were trying to fine-tune this algorithm so that it worked as well as possible in real life? Justify your answer.

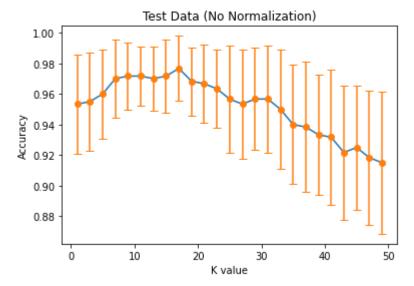
I would select a K value of about 15, as this is when the model starts to switch from overfitting to underfitting. In both the training and testing data, the accuracy is around 0.96 at K = 15, meaning the estimates for both sets are similarly accurate (although the testing data shows a significantly greater std deviation).

**Q1.6 (10 Points)** In the experiments conducted earlier, you normalized the features before running k-NN. This is the appropriate procedure to ensure that all features are considered equally important when computing distances. Now, you will study the impact of omitting feature normalization on the performance of the algorithm. To accomplish this, you will repeat Q1.2 and create a graph depicting the average accuracy (and corresponding standard deviation) of k-NN as a function of k, when evaluated on the testing set. However, this time you will run the algorithm without first normalizing the features. This means that you will run k-NN directly on the instances present in the original dataset without performing any pre-processing normalization steps to ensure that all features lie in the same range/interval. Now (a) present the graph you created;

```
In []: test_iter_raw = []
    test_acc_raw = []
    test_std_raw = []

for k_val,res in test_no_norm.items():
        test_iter_raw.append(k_val)
        test_acc_raw.append(np.average(res))
        test_std_raw.append(np.std(res))

plt.plot(test_iter_raw,test_acc_raw)
    plt.errorbar(test_iter_raw,test_acc_raw,yerr = test_std_raw,fmt='o',capsize=4)
    plt.xlabel("K value")
    plt.ylabel("Accuracy")
    plt.title("Test Data (No Normalization)")
    plt.show()
```



(b) based on this graph, identify the best value of k; that is, the value of k that results in k-NN performing the best on the testing set;

Based on this graph, I would choose a value of K closer to 20, either 17 or 19. This is because the accuracy is higher around these points and the std deviation is lower when compared to the normalized graphs.

and (c) describe how the performance of this version of k-NN (without feature normalization) compares with the performance of k-NN with feature normalization. Discuss intuitively the reasons why one may have performed better than the other.

This version of KNN seems to perform better than the version with normalization. The accuracy of the non-normalized KNN is slightly higher and the std deviation is lower in the K = 15-20 range. The reasons for this improved accuracy and deviation may be due to the fact that normalizing the test data on the training data could cause the model to overfit to the training data.

## 2. Evaluating the Decision Tree Algorithm

```
Function to create decision tree based on data
do ig - boolean, True to use info gain, False to use gini critereon
set_depth - float (default 1) to set as the proportions of instances to create a leaf
def learn_tree(data,attr_list,do_ig,set_depth=1.0):
    #if proportion of entries in the same target class > set depth, return leaf node of
    if (data['target'].value counts().iloc[0]/len(data)) >= set depth:
        return DecisionNode(None, None, None, None, data['target'].unique()[0])
    #if no more attr to be tested, return leaf w majority class
    if len(attr list) == 0:
        return DecisionNode(None, None, None, None, data['target'].mode()[0])
    #find best attr from list using info gain
    max gain = float('-inf')
    min gini = float('inf')
    best_attr = ''
    for attr in attr list:
        if do ig:
            ig_attr = get_info_gain(data,attr)
            if ig_attr > max_gain:
                max gain = ig attr
                best attr = attr
        else:
            gini_attr = get_critereon(data,attr)
            if gini_attr < min_gini:</pre>
                min_gini = gini_attr
                best attr = attr
    #remove attr from list
    attr_list.remove(best_attr)
    #create children, if no data with that value, create leaf node with majority class
    child_0 = learn_tree(data.loc[data[best_attr] == 0],attr_list,do_ig) if len(data.]
    child 1 = learn tree(data.loc[data[best attr] == 1],attr list,do ig) if len(data.]
    child_2 = learn_tree(data.loc[data[best_attr] == 2],attr_list,do_ig) if len(data.l
    return DecisionNode(best attr,child 0,child 1,child 2)
#Helper functions for decision tree
def get info gain(data,attr):
    data 0 = data.loc[data[attr] == 0]
    data_1 = data.loc[data[attr] == 1]
    data 2 = data.loc[data[attr] == 2]
    split entropy = (entropy(data 0)*len(data 0) + entropy(data 1)*len(data 1) + entro
    return entropy(data) - split entropy
def entropy(data):
    if len(data) == 0:
        return 0
    prob 0 = data['target'].value counts().iloc[0]/len(data)
    ent = -(prob_0 * math.log(prob_0,2))
    if len(data['target'].value counts()) > 1:
        prob_1 = data['target'].value_counts().iloc[1]/len(data)
        ent -= (prob_1 * math.log(prob_1,2))
    else:
```

```
return ent
                      def get_critereon(data,attr):
                                data_0 = data.loc[data[attr] == 0]
                                data 1 = data.loc[data[attr] == 1]
                                data 2 = data.loc[data[attr] == 2]
                                return (gini(data_0)*len(data_0) + gini(data_1)*len(data_1) + gini(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len(data_2)*len
                      def gini(data):
                                if len(data) == 0:
                                           return 0
                                prob_0 = data['target'].value_counts().iloc[0]/len(data)
                                crit = prob 0 ** 2
                                if len(data['target'].value_counts()) > 1:
                                           prob_1 = data['target'].value_counts().iloc[1]/len(data)
                                           crit += prob 1 ** 2
                                else:
                                           pass
                                return crit
                       #train tree on training data
                       voter_df = pd.read_csv('house_votes_84.csv')
In [ ]: #function to classify an instance based on decision tree
                      def classify(instance):
                                node = decision tree
                                while node.label is None:
                                           if instance[node.attr] == 0:
                                                     node = node.child 0
                                           elif instance[node.attr] == 1:
                                                     node = node.child 1
                                           else:
                                                     node = node.child_2
                                 guess = node.label
                                actual = instance['target']
                                return guess
                       #test accuracy on training data
                       def test decision(to train):
                                 predictions = to train.apply(lambda row: classify(row), axis=1)
                                actual labels = voter df.loc[predictions.index,['target']]
                                return predictions.eq(actual_labels['target'].values).mean()
                       accuracy test = []
                       accuracy_train = []
                       for i in range (100):
                                voter_train, voter_test = train_test_split(voter_df,test_size=0.2)
```

**Q2.1 (12 Points)** In the first histogram, you should show the accuracy distribution when the algorithm is evaluated over training data. The horizontal axis should show different accuracy values, and the vertical axis should show the frequency with which that accuracy was observed while conducting these 100 experiments/training processes. The histogram should look like the

attributes = list(voter train.columns.values)

decision\_tree = learn\_tree(voter\_train,attributes,True)
accuracy\_train.append(test\_decision(voter\_train))
accuracy\_test.append(test\_decision(voter\_test))

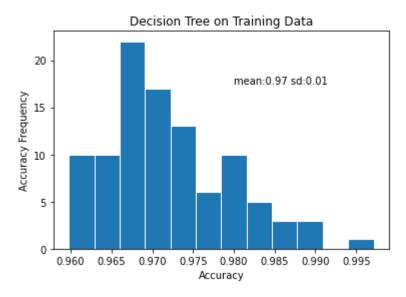
attributes.remove('target')

one in Figure 3 (though the "shape" of the histogram you obtain may be different, of course). You should also report the mean accuracy and its standard deviation.

Mean: 0.97, Standard Deviation: 0.01

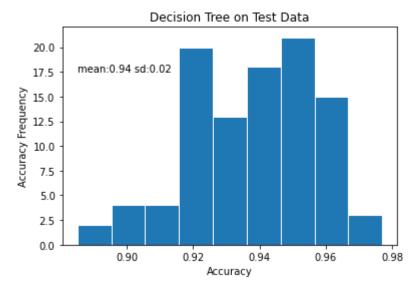
```
In []: num_bins = len(set(accuracy_train))
    plt.hist(accuracy_train,bins=num_bins,edgecolor='white',linewidth=1)
    plt.ylabel("Accuracy Frequency")
    plt.xlabel("Accuracy")
    plt.title("Decision Tree on Training Data")
    plt.text(0.98,17.5,f'mean:{np.mean(accuracy_train):.2f} sd:{np.std(accuracy_train):.2f}
    plt.show
```

Out[ ]: <function matplotlib.pyplot.show(close=None, block=None)>



**Q2.2 (12 Points)** In the second histogram, you should show the accuracy distribution when the algorithm is evaluated over testing data. The horizontal axis should show different accuracy values, and the vertical axis should show the frequency with which that accuracy was observed while conducting these 100 experiments/training processes. You should also report the mean accuracy and its standard deviation.

Mean: 0.94, Standard Deviation: 0.02



**Q2.3 (12 Points)** Explain intuitively why each of these histograms looks the way they do. Is there more variance in one of the histograms? If so, why do you think that is the case? Does one histogram show higher average accuracy than the other? If so, why do you think that is the case?

The test data histogram has a higher standard deviation and a lower accuracy than the training data histogram. This makes sense, as the training data was used to create the decision tree.

**Q2.4 (8 Points)** By comparing the two histograms, would you say that the Decision Trees algorithm, when used in this dataset, is underfitting, overfitting, or performing reasonably well? Explain your reasoning.

The algorithm is slightly overfitting, but it could be argued that it is performing reasonably well. The model scores an average 0.97 accuracy on the training data and 0.94 on the test data. This accuracy is high but adjustments to the conditions for determining leaf nodes could improve this accuracy.

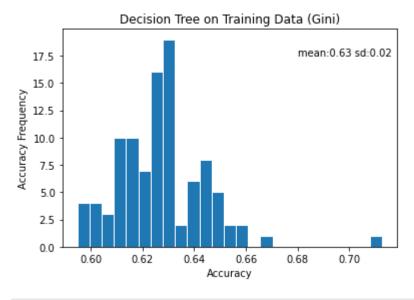
**Q2.5 (6 Points)** In class, we discussed how Decision Trees might be non-robust. Is it possible to experimentally confirm this property/tendency via these experiments, by analyzing the histograms you generated and their corresponding average accuracies and standard deviations? Explain your reasoning.

It is experimentally possible to confirm the issue of decision trees being non-robust. The histogram displaying the performance on the test data shows this somewhat due to its standard deviation of 0.02 (double that of the testing data). If the number of iterations were increased, it is likely that this issue would be more visible.

**[QE.1] Extra points (15 Points)** Repeat the experiments Q2.1 to Q2.4, but now use the Gini criterion for node splitting, instead of the Information Gain criterion.

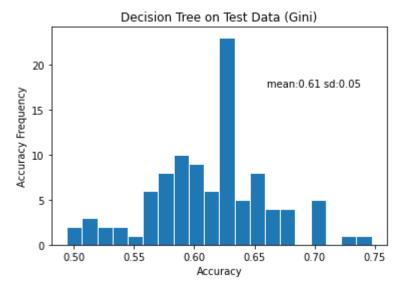
```
In [ ]: num_bins = len(set(accuracy_train_gini))
    plt.hist(accuracy_train_gini,bins=num_bins,edgecolor='white',linewidth=1)
    plt.ylabel("Accuracy Frequency")
    plt.xlabel("Accuracy")
    plt.title("Decision Tree on Training Data (Gini)")
    plt.text(0.68,17.5,f'mean:{np.mean(accuracy_train_gini):.2f} sd:{np.std(accuracy_train_gini):.2f}
```

Out[ ]: <function matplotlib.pyplot.show(close=None, block=None)>



```
In []: num_bins = len(set(accuracy_test_gini))
    plt.hist(accuracy_test_gini,bins=num_bins,edgecolor='white',linewidth=1)
    plt.ylabel("Accuracy Frequency")
    plt.xlabel("Accuracy")
    plt.title("Decision Tree on Test Data (Gini)")
    plt.text(0.66,17.5,f'mean:{np.mean(accuracy_test_gini):.2f} sd:{np.std(accuracy_test_gini):.2f} sd:{np.std(accuracy_test_gini):.2f}
```

Out[ ]: <function matplotlib.pyplot.show(close=None, block=None)>



Explain intuitively why each of these histograms looks the way they do. Is there more variance in one of the histograms? If so, why do you think that is the case? Does one histogram show higher average accuracy than the other? If so, why do you think that is the case?

The histogram for the training data has a higher average accuracy and lower standard deviation than the test data. Similarly to the graphs when using information gain, this is due to the fact that the training data is used to train the decision tree, so training data is likely to perform better than the testing data. In general the Gini critereon performs significantly worse than information gain.

By comparing the two histograms, would you say that the Decision Trees algorithm, when used in this dataset, is underfitting, overfitting, or performing reasonably well? Explain your reasoning.

The algorithm seems to be underfitting, as the accuracy of both the training and testing data is very low and has a high standard deviation.

**[QE.2] Extra points (15 Points)** Repeat the experiments Q2.1 to Q2.4 but now use a simple heuristic to keep the tree from becoming too "deep"; i.e., to keep it from testing a (possibly) excessive number of attributes, which is known to often cause overfitting. To do this, use an additional stopping criterion: whenever more than 85% of the instances associated with a decision node belong to the same class, do not further split this node. Instead, replace it with a leaf node whose class prediction is the majority class within the corresponding instances. E.g., if 85% of the instances associated with a given decision node have the label/class Democrat, do not further split this node, and instead directly return the prediction Democrat.

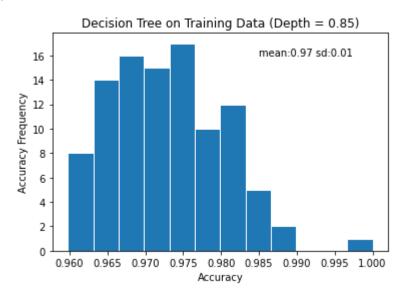
```
In [ ]: accuracy_test_depth = []
    accuracy_train_depth = []

for i in range (100):
    voter_train, voter_test = train_test_split(voter_df,test_size=0.2)
    attributes = list(voter_train.columns.values)
    attributes.remove('target')
```

```
decision_tree = learn_tree(voter_train,attributes,True,0.85)
accuracy_train_depth.append(test_decision(voter_train))
accuracy_test_depth.append(test_decision(voter_test))
```

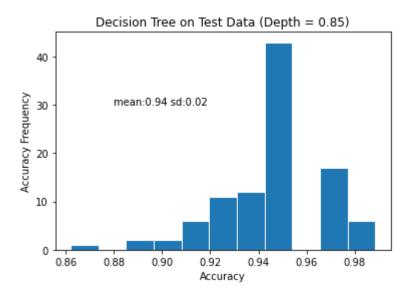
```
In [ ]: num_bins = len(set(accuracy_train_depth))
    plt.hist(accuracy_train_depth,bins=num_bins,edgecolor='white',linewidth=1)
    plt.ylabel("Accuracy Frequency")
    plt.xlabel("Accuracy")
    plt.title("Decision Tree on Training Data (Depth = 0.85)")
    plt.text(0.985,16,f'mean:{np.mean(accuracy_train_depth):.2f} sd:{np.std(accuracy_train_depth):.2f}
```

Out[ ]: <function matplotlib.pyplot.show(close=None, block=None)>



```
In [ ]: num_bins = len(set(accuracy_test_depth))
    plt.hist(accuracy_test_depth,bins=num_bins,edgecolor='white',linewidth=1)
    plt.ylabel("Accuracy Frequency")
    plt.xlabel("Accuracy")
    plt.title("Decision Tree on Test Data (Depth = 0.85)")
    plt.text(0.88,30,f'mean:{np.mean(accuracy_test_depth):.2f} sd:{np.std(accuracy_test_depth):.2f}
```

Out[ ]: <function matplotlib.pyplot.show(close=None, block=None)>



Explain intuitively why each of these histograms looks the way they do. Is there more variance in one of the histograms? If so, why do you think that is the case? Does one histogram show higher average accuracy than the other? If so, why do you think that is the case?

The histogram for the training data has a higher average accuracy and lower standard deviation than the test data. Similarly to the previous graphs, this is due to the fact that the training data is used to train the decision tree, so training data is likely to perform better than the testing data.

By comparing the two histograms, would you say that the Decision Trees algorithm, when used in this dataset, is underfitting, overfitting, or performing reasonably well? Explain your reasoning.

The algorithm has similar outputs to the original information gain model. The algorithm is slightly overfitting, but it could be argued that it is performing reasonably well. The model scores an average 0.97 accuracy on the training data and 0.94 on the test data.