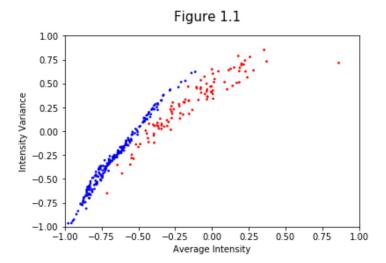
```
In [88]: #From the console, run the following
         #pip install numpy
         #pip install scipy
         #pip install scikit-learn
         #pip install matplotlib
         # Import required packages here (after they are installed)
         import numpy as np
         from sklearn.neighbors import KNeighborsClassifier
         import matplotlib.pyplot as mp
         from pylab import show
         from sklearn.model_selection import cross_val_score
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.ensemble import RandomForestClassifier
         from sklearn.ensemble import AdaBoostClassifier
         # Load data. csv file should be in the same folder as the notebook for this to wor
         k, otherwise
         # give data path.
         data = np.loadtxt("data.csv")
In [89]: #shuffle the data and select training and test data
         np.random.seed(100)
         np.random.shuffle(data)
         features = []
         digits = []
         for row in data:
             #import the data and select only the 1's and 5's
             if(row[0]==1 or row[0]==5):
                 features.append(row[1:])
                 digits.append(str(row[0]))
         #Select the proportion of data to use for training.
         #Notice that we have set aside 80% of the data for testing
         numTrain = int(len(features)*.2)
         trainFeatures = features[:numTrain]
         testFeatures = features[numTrain:]
         trainDigits = digits[:numTrain]
         testDigits = digits[numTrain:]
         #print(trainFeatures)
         #trainFeatures[0]
```

```
In [90]: #Convert the 256D data (trainFeatures) to 2D data
         \#We need X and Y for plotting and simpleTrain for building the model.
         #They contain the same points in a different arrangement
         X = []
         Y = []
         simpleTrain = []
         #Colors will be passed to the graphing library to color the points.
         #1's are blue: "b" and 5's are red: "r"
         colors = []
         \#legends = []
         for index in range(len(trainFeatures)):
             #print(index)
             #produce the 2D dataset for graphing/training and scale the data so it is in th
         e [-1,1] square
             xNew = 2*np.average(trainFeatures[index])+.75
             yNew = 3*np.var(trainFeatures[index])-1.5
             X.append(xNew)
             Y.append(yNew)
             simpleTrain.append([xNew,yNew])
             #trainDigits will still be the value we try to classify. Here it is the string
         "1.0" or "5.0"
             if(trainDigits[index] == "1.0"):
                 colors.append("b")
                 #legends.append("1")
                 colors.append("r")
                 #legends.append("5")
         #plot the data points
         ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
         fig = mp.figure()
         mp.scatter(X,Y,s=3,c=colors)
         #specify the axes
         mp.xlim(-1,1)
         mp.xlabel("Average Intensity")
         mp.ylim(-1,1)
         mp.ylabel("Intensity Variance")
         #Labeling the plot
         #mp.legend(['1'])
         #mp.legend(legends)
         fig.suptitle('Figure 1.1', fontsize=15)
         #display the current graph
         show()
```



DECISON TREES

1a) DECISION TREES - maxleaf nodes

```
In [91]: #create a new decision tree model
         max_leaf_nodes_list = [5,10,15,20,30,40,50,75,100,200,500,1000,10000]
         decisiontree_models = []
         cv_scores_list = []
         cv errors = []
         # iterating each number in list
         for num in max leaf nodes list:
             dt clf = DecisionTreeClassifier(criterion='entropy', max leaf nodes = num, rand
             decisiontree models.append(dt clf)
         #train model with cv of 10
         for dt in decisiontree models:
             cv_scores = cross_val_score(dt, simpleTrain, trainDigits, cv=10)
             cv_scores_list.append(cv_scores)
         #print each cv score (accuracy) and average them
         for i in range(len(max leaf nodes list)):
            print('Max leaf nodes:', max_leaf_nodes_list[i])
             print(decisiontree_models[i])
             print(cv_scores_list[i])
             print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
             error = 1-np.mean(cv_scores_list[i])
             cv_errors.append(error)
             print('Cross validation error:', error)
```

4/23/2020, 11:37 PM

```
Max leaf nodes: 5
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=5,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
          0.875
                      0.9375
                                           0.80645161 0.80645161
                              1.
0.87096774 0.90322581 0.93548387 0.83333333]
cv scores mean: 0.8905913978494624
Cross validation error: 0.1094086021505376
Max leaf nodes: 10
DecisionTreeClassifier(class_weight=None, criterion='entropy', max_depth=None,
                      max features=None, max leaf nodes=10,
                      min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction_leaf=0.0, presort=False,
                      random_state=0, splitter='best')
[0.9375
          0.96875
                     0.96875
                              0.96774194 0.87096774 0.87096774
                     0.90322581 0.9
           1.
cv scores mean: 0.9387903225806452
Cross validation error: 0.06120967741935479
Max leaf nodes: 15
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=15,
                      min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
[0.9375 0.96875
                    0.96875 0.96774194 0.83870968 0.87096774
           1.
                     0.96774194 0.9
                                         1
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Max leaf nodes: 20
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=20,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, presort=False,
                     random_state=0, splitter='best')
1.
                     0.96774194 0.9
                                         ]
cv scores mean: 0.9387903225806452
Cross validation error: 0.06120967741935479
Max leaf nodes: 30
DecisionTreeClassifier(class_weight=None, criterion='entropy', max_depth=None,
                      max_features=None, max_leaf_nodes=30,
                      min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
[0.9375
          0.96875
                     0.96875
                              0.96774194 0.83870968 0.83870968
                      0.96774194 0.9
           1.
cv scores mean: 0.9387903225806452
Cross validation error: 0.06120967741935479
Max leaf nodes: 40
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=40,
                      min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
[0.9375
           0.96875
                     0.96875 0.96774194 0.83870968 0.83870968
                      0.96774194 0.9
1.
           1.
                                          1
```

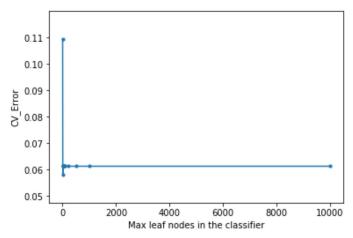
```
In [92]: #plot the data points
    ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
    fig = mp.figure()
    mp.scatter(max_leaf_nodes_list,cv_errors,s=10)
    mp.plot(max_leaf_nodes_list,cv_errors)

#specify the axes
    mp.xlabel("Max leaf nodes in the classifier")
    mp.ylabel("CV_Error")

#Labeling the plot
    fig.suptitle('Decision Tree Models with varying max leaf nodes', fontsize=15)

#display the current graph
    show()
```

Decision Tree Models with varying max leaf nodes



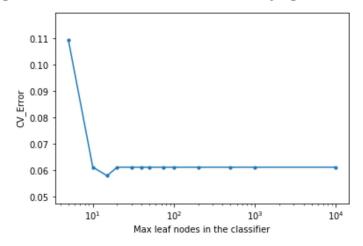
```
In [93]: #plot the data points
    ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
    fig = mp.figure()
    mp.scatter(max_leaf_nodes_list,cv_errors,s=10)
    mp.xscale('log')
    mp.plot(max_leaf_nodes_list,cv_errors)

#specify the axes
    mp.xlabel("Max leaf nodes in the classifier")
    mp.ylabel("CV_Error")

#Labeling the plot
    fig.suptitle('Figure 4.1 - Decision Tree Models with varying max leaf nodes', fonts ize=15)

#display the current graph
    show()
```

Figure 4.1 - Decision Tree Models with varying max leaf nodes



1b) Evidence of overfitting or underfitting

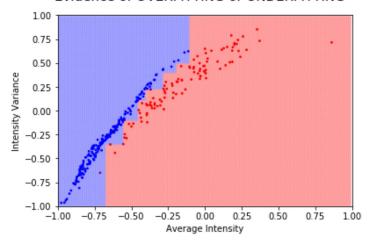
```
In [94]:
         #depth of the tree
         depth = []
         leaves = []
         path = []
         for dt in decisiontree_models:
             dt.fit(simpleTrain,trainDigits)
             dpt = dt.get_depth()
             depth.append(dpt)
             lfs = dt.get n leaves()
             leaves.append(lfs)
             pth = dt.decision path(simpleTrain)
             path.append(pth)
         print("Depth:", depth)
         print("Leaves:", leaves)
         Depth: [3, 5, 6, 7, 7, 7, 7, 7, 7, 7, 7, 7]
         Leaves: [5, 10, 15, 16, 16, 16, 16, 16, 16, 16, 16, 16]
```

```
In [95]: #table
         figb t = mp.figure()
         table_vals_b = []
         np array = np.array(cv errors)
         np round to tenths = np.around(np array, 5)
         round to tenths = list(np round to tenths)
         table vals b.append(round to tenths)
         np array = np.array(depth)
         np round to tenths = np.around(np array, 5)
         round to tenths = list(np round to tenths)
         table_vals_b.append(round_to_tenths)
         np array = np.array(leaves)
         np_round_to_tenths = np.around(np_array, 5)
         round_to_tenths = list(np_round_to_tenths)
         table vals b.append(round to tenths)
         row labels = ['cv error', 'Depth', 'leaves']
         col_labels = max_leaf_nodes_list
         the table = mp.table(cellText=table vals b,
                               rowLabels=row labels,
                               colLabels=col_labels,
                               loc='center')
         the_table.auto_set_font_size(False)
         the table.set fontsize(14)
         the table.scale(3,3)
         # Removing ticks and spines enables you to get the figure only with table
         mp.tick_params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
         mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
         for pos in ['right','top','bottom','left']:
             mp.gca().spines[pos].set visible(False)
         mp.savefig('matplotlib-table.png', bbox inches='tight', pad inches=0.05)
         #display the current graph
         mp.show()
```

	5	10	15	20	30	40	50	75	100	200	500	1000	10000
cv_error	0.10941	0.06121	0.05798	0.06121	0.06121	0.06121	0.06121	0.06121	0.06121	0.06121	0.06121	0.06121	0.06121
Depth	3	5	6	7	7	7	7	7	7	7	7	7	7
leaves	5	10	15	16	16	16	16	16	16	16	16	16	16

```
In [96]: # create the model
         # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCla
         ssifier.html
         # Declare Model
         model = DecisionTreeClassifier(criterion='entropy', max leaf nodes = 10000, random
         state=0)
         # Fit model to our data
         model.fit(simpleTrain, trainDigits)
         # Lists to hold inpoints, predictions and assigned colors
         xPred = []
         yPred = []
         cPred = []
         # Use input points to get predictions here
         for xP in range(-100,100):
             xP = xP/100.0
             for yP in range(-100,100):
                 yP = yP/100.0
                 xPred.append(xP)
                 yPred.append(yP)
                 if (model.predict([[xP,yP]]) == "1.0"):
                     cPred.append("b")
                 else:
                     cPred.append("r")
```

Evidence of OVERFITTING or UNDERFITTING

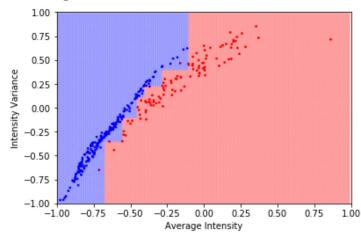


1b) ANSWER: Even though we have max leaf nodes till 10,000, the actual leaf nodes were only 16 at max. From the 4th model i.e., max leaf nodes = 20, all the trees had same depth and same leaf nodes. So we can say that from max leaf nodes 20, there can be a chance of overfitting since the model has reached maximum depth and can lead to overfitting. Similarly we can see that the cross validation errors are also increasing and similar.

1c) 2D REGION FOR OPTIMAL DECISION TREE MODEL

```
In [98]: # create the model
         # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCla
         ssifier.html
         # Declare Model
         model = DecisionTreeClassifier(criterion='entropy', max_leaf_nodes = 15, random_sta
         te=0)
         # Fit model to our data
         model.fit(simpleTrain,trainDigits)
         # Lists to hold inpoints, predictions and assigned colors
         xPred = []
         yPred = []
         cPred = []
         # Use input points to get predictions here
         for xP in range(-100,100):
             xP = xP/100.0
             for yP in range(-100,100):
                 yP = yP/100.0
                 xPred.append(xP)
                 yPred.append(yP)
                 if (model.predict([[xP,yP]]) == "1.0"):
                     cPred.append("b")
                 else:
                     cPred.append("r")
```

Figure 4.2 - OPTIMAL DECISION TREE MODEL



1c) ANSWER: The optimal model I have choosen is for max leaf nodes with 15. This trees actual leaf nodes in the model were 15 and the depth of the tree was 6 and it has the lowest cross validation error.

1d) GRADUATE STUDENT QUESTION - min impurity decrease & min impurity split

```
In [100]: #create a new decision tree model - min impurity decrease
          min impurity decrease list = [0,0.0001,0.001,0.01,0.05,0.1,0.2,0.3,0.4,0.5,0.6,0.
          7,0.8,0.9,1]
          decisiontree_models = []
          cv scores list = []
          cv errors = []
          # iterating each number in list
          for imp in min impurity decrease list:
              dt clf = DecisionTreeClassifier(criterion='entropy', min impurity decrease=im
          p, random state=0)
              decisiontree models.append(dt clf)
          #train model with cv of 10
          for dt in decisiontree models:
              cv_scores = cross_val_score(dt, simpleTrain, trainDigits, cv=10)
              cv_scores_list.append(cv_scores)
          #print each cv score (accuracy) and average them
          for i in range(len(min_impurity_decrease_list)):
              print('Min impurity decrease value:', min_impurity_decrease_list[i])
              print(decisiontree models[i])
              print(cv scores list[i])
              print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
              error = 1-np.mean(cv_scores_list[i])
              cv_errors.append(error)
              print('Cross validation error:', error)
```

```
Min impurity decrease value: 0
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=None,
                      min_impurity_decrease=0, min_impurity_split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
[0.9375
           0.96875
                      1.
                      0.96774194 0.9
          1.
                                          ]
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity decrease value: 0.0001
DecisionTreeClassifier(class_weight=None, criterion='entropy', max_depth=None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0001, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction_leaf=0.0, presort=False,
                      random_state=0, splitter='best')
[0.9375
           0.96875
                      0.96875 0.96774194 0.83870968 0.87096774
1.
           1.
                      0.96774194 0.9
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity decrease value: 0.001
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max_features=None, max_leaf_nodes=None,
                      min impurity decrease=0.001, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
[0.9375 0.96875
                    0.96875 0.96774194 0.83870968 0.87096774
                      0.96774194 0.9
           1.
                                          1
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity decrease value: 0.01
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=None,
                      min_impurity_decrease=0.01, min_impurity_split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, presort=False,
                      random_state=0, splitter='best')
[0.9375
          0.96875 0.96875 0.96774194 0.87096774 0.83870968
           1.
                      0.93548387 0.9
                                          ]
cv_scores mean:0.9387903225806452
Cross validation error: 0.06120967741935479
Min impurity decrease value: 0.05
DecisionTreeClassifier(class_weight=None, criterion='entropy', max_depth=None,
                      max_features=None, max_leaf_nodes=None,
                      min_impurity_decrease=0.05, min_impurity_split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
                      0.96875
                               0.90322581 0.77419355 0.80645161
[0.9375
         0.9375
0.90322581 0.90322581 0.93548387 0.9
cv scores mean: 0.8969556451612902
Cross validation error: 0.10304435483870977
Min impurity decrease value: 0.1
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.1, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
           0.84375
                      0.9375
                                           0.80645161 0.83870968
                               1.
                      0.87096774 0.86666667]
 0.87096774 1.
```

```
In [101]: #depth of the tree
    depth = []
    leaves = []
    path = []

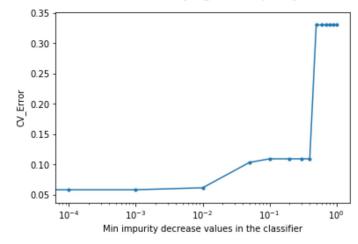
for dt in decisiontree_models:
        dt.fit(simpleTrain,trainDigits)
        dpt = dt.get_depth()
        depth.append(dpt)
        lfs = dt.get_n_leaves()
        leaves.append(lfs)
        pth = dt.decision_path(simpleTrain)
        path.append(pth)

print("Depth:", depth)
    print("Leaves:", leaves)
```

Depth: [7, 7, 7, 5, 3, 1, 1, 1, 1, 0, 0, 0, 0, 0, 0] Leaves: [16, 16, 16, 12, 6, 2, 2, 2, 2, 1, 1, 1, 1, 1, 1]

```
In [102]: #plot the data points
          ### https://matplotlib.org/api/ as gen/matplotlib.pyplot.scatter.html
          fig = mp.figure()
          mp.scatter(min_impurity_decrease_list,cv_errors,s=10)
          mp.xscale('log')
          mp.plot(min impurity decrease list,cv errors)
          #specify the axes
          mp.xlabel("Min impurity decrease values in the classifier")
          mp.ylabel("CV Error")
          #Labeling the plot
          fig.suptitle('Decision Tree Models with varying min impurity decrease values', fon
          tsize=15)
          #table
          figb t = mp.figure()
          table vals b = []
          np_array = np.array(cv_errors)
          np_round_to_tenths = np.around(np_array, 5)
          round_to_tenths = list(np_round_to_tenths)
          table_vals_b.append(round_to_tenths)
          np_array = np.array(depth)
          np round to tenths = np.around(np array, 5)
          round to tenths = list(np round to tenths)
          table vals b.append(round to tenths)
          np array = np.array(leaves)
          np_round_to_tenths = np.around(np_array, 5)
          round_to_tenths = list(np_round_to_tenths)
          table vals b.append(round to tenths)
          row labels = ['cv error', 'Depth', 'leaves']
          col labels = min impurity decrease list
          the table = mp.table(cellText=table vals b,
                                rowLabels=row labels,
                                 colLabels=col labels,
                                 loc='center')
          the table.auto set font size (False)
          the table.set fontsize(14)
          the table.scale (3,3)
          # Removing ticks and spines enables you to get the figure only with table
          mp.tick params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
          mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
          for pos in ['right','top','bottom','left']:
              mp.gca().spines[pos].set_visible(False)
          mp.savefig('matplotlib-table.png', bbox_inches='tight', pad_inches=0.05)
          #display the current graph
          mp.show()
```

Decision Tree Models with varying min impurity decrease values



	0	0.0001	0.001	0.01	0.05	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8	0.9	1
cv_error	0.05798	0.05798	0.05798	0.06121	0.10304	0.109	0.109	0.109	0.109	0.33001	0.33001	0.33001	0.33001	0.33001	0.33001
Depth	7	7	7	5	3	1	1	1	1	0	0	0	0	0	0
leaves	16	16	16	12	6	2	2	2	2	1	1	1	1	1	1

```
In [111]: #create a new decision tree model - min impurity split
          \min_{\text{impurity}} \text{split\_list} = [0,0.0001,0.001,0.01,0.05,0.1,0.2,0.3,0.4,0.5,0.6,0.7,0.]
          8,0.9,1]
          decisiontree_models = []
          cv scores list = []
          cv errors = []
          # iterating each number in list
          for imp in min impurity split list:
              dt clf = DecisionTreeClassifier(criterion='entropy', min impurity split=imp, r
          andom state=0)
              decisiontree models.append(dt clf)
          #train model with cv of 10
          for dt in decisiontree models:
              cv_scores = cross_val_score(dt, simpleTrain, trainDigits, cv=10)
              cv_scores_list.append(cv_scores)
          #print each cv score (accuracy) and average them
          for i in range(len(min_impurity_split_list)):
              print('Min impurity split value:', min_impurity_split_list[i])
              print(decisiontree models[i])
              print(cv scores list[i])
              print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
              error = 1-np.mean(cv_scores_list[i])
              cv_errors.append(error)
              print('Cross validation error:', error)
```

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

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DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

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DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

```
Min impurity split value: 0
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0, min impurity split=0,
                      min_samples_leaf=1, min_samples_split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
           0.96875
                      0.96875
                              0.96774194 0.83870968 0.87096774
1.
          1.
                     0.96774194 0.9
                                         ]
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity split value: 0.0001
DecisionTreeClassifier(class_weight=None, criterion='entropy', max_depth=None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0, min impurity split=0.0001,
                      min samples leaf=1, min samples split=2,
                      min weight fraction_leaf=0.0, presort=False,
                      random_state=0, splitter='best')
[0.9375
          0.96875
                     0.96875
                              0.96774194 0.83870968 0.87096774
1.
           1.
                     0.96774194 0.9
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity split value: 0.001
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0, min impurity split=0.001,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                     random state=0, splitter='best')
[0.9375 0.96875
                     0.96875
                              0.96774194 0.83870968 0.87096774
                     0.96774194 0.9
           1.
                                         1
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity split value: 0.01
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0, min impurity split=0.01,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, presort=False,
                     random_state=0, splitter='best')
[0.9375
         1.
                     0.96774194 0.9
cv_scores mean:0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity split value: 0.05
DecisionTreeClassifier(class_weight=None, criterion='entropy', max_depth=None,
                      max_features=None, max_leaf_nodes=None,
                      min_impurity_decrease=0.0, min_impurity_split=0.05,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
[0.9375
           0.96875
                    0.96875
                              0.96774194 0.83870968 0.87096774
                      0.96774194 0.9
           1.
cv scores mean: 0.9420161290322581
Cross validation error: 0.057983870967741935
Min impurity split value: 0.1
DecisionTreeClassifier(class weight=None, criterion='entropy', max depth=None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0, min impurity split=0.1,
                      min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, presort=False,
                      random state=0, splitter='best')
[0.9375
           0.96875
                     0.96774194 0.9
1.
           1.
                                         1
```

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

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DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

```
In [112]: #depth of the tree
    depth = []
    leaves = []
    path = []

for dt in decisiontree_models:
        dt.fit(simpleTrain,trainDigits)
        dpt = dt.get_depth()
        depth.append(dpt)
        lfs = dt.get_n_leaves()
        leaves.append(lfs)
        pth = dt.decision_path(simpleTrain)
        path.append(pth)

print("Depth:", depth)
print("Leaves:", leaves)
```

Depth: [7, 7, 7, 7, 7, 7, 7, 7, 7, 7, 1, 1, 1, 0]
Leaves: [16, 16, 16, 16, 16, 16, 12, 12, 8, 8, 8, 2, 2, 2, 1]

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

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DeprecationWarning)

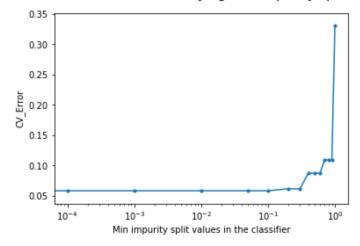
C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min _impurity_decrease parameter instead.

DeprecationWarning)

C:\Users\Anu\Anaconda3\lib\site-packages\sklearn\tree\tree.py:297: DeprecationWa rning: The min_impurity_split parameter is deprecated. Its default value will ch ange from 1e-7 to 0 in version 0.23, and it will be removed in 0.25. Use the min impurity decrease parameter instead.

```
In [113]: #plot the data points
          ### https://matplotlib.org/api/ as gen/matplotlib.pyplot.scatter.html
          fig = mp.figure()
          mp.scatter(min_impurity_split_list,cv_errors,s=10)
          mp.xscale('log')
          mp.plot(min impurity split list,cv errors)
          #specify the axes
          mp.xlabel("Min impurity split values in the classifier")
          mp.ylabel("CV Error")
          #Labeling the plot
          fig.suptitle('Decision Tree Models with varying min impurity split values', fontsi
          ze=15)
          #table
          figb t = mp.figure()
          table vals b = []
          np_array = np.array(cv_errors)
          np_round_to_tenths = np.around(np_array, 5)
          round to tenths = list(np round to tenths)
          table_vals_b.append(round_to_tenths)
          np_array = np.array(depth)
          np round to tenths = np.around(np array, 5)
          round to tenths = list(np round to tenths)
          table vals b.append(round to tenths)
          np array = np.array(leaves)
          np_round_to_tenths = np.around(np_array, 5)
          round_to_tenths = list(np_round_to_tenths)
          table vals b.append(round to tenths)
          row labels = ['cv error', 'Depth', 'leaves']
          col labels = min impurity split list
          the table = mp.table(cellText=table vals b,
                                rowLabels=row labels,
                                 colLabels=col labels,
                                 loc='center')
          the table.auto set font size (False)
          the table.set fontsize(14)
          the table.scale (3,3)
          # Removing ticks and spines enables you to get the figure only with table
          mp.tick params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
          mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
          for pos in ['right','top','bottom','left']:
              mp.gca().spines[pos].set_visible(False)
          mp.savefig('matplotlib-table.png', bbox_inches='tight', pad_inches=0.05)
          #display the current graph
          mp.show()
```

Decision Tree Models with varying min impurity split values



	0	0.0001	0.001	0.01	0.05	0.1	0.2	0.3	0.4	0.5	0.6	0.7	0.8	0.9	1
cv_error	0.05798	0.05798	0.05798	0.05798	0.05798	0.05798	0.06121	0.06121	0.08692	0.08692	0.08692	0.109	0.109	0.109	0.33001
Depth	7	7	7	7	7	7	7	7	7	7	7	1	1	1	0
leaves	16	16	16	16	16	16	12	12	8	8	8	2	2	2	1

Answer: The optimal values as per cross validation errors for min impurity decrease are 0, 0.0001, 0.001 and are similar for min impurity split since both of them almost explain the same thing. But if we want to reduce overfitting, we need to introduce some amount of impurity in the nodes. So for min impurity decrease the optimal value can be 0.001 if we want least impurity or we can consider 0.001 which has cross validation error as 0.06121 that can reduce both the depth and overfitting in the classifier which can reduce variance. Here, considering 0.1 as 'min impurity decrease' would underfit the data so much since the depth and leaves produced are just 1 and 2. So this means data is classified as either one side or the other. So, we cannot choose this value as this underfits the data and similarly for other values from there on. Similarly min impurity split can be considered as '0.1' as this has the lowest cross validation error and also doesn't underfit the data.

RANDOM FORESTS

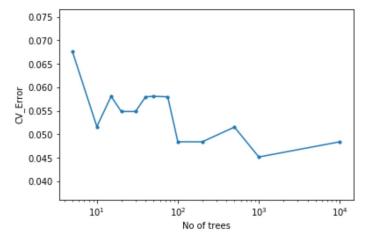
1e) PART 1 : RANDOM FORESTS - max leaf nodes = 10

```
In [91]: #create a new decision tree model
         n_{estimators_list} = [5, 10, 15, 20, 30, 40, 50, 75, 100, 200, 500, 1000, 10000]
         randomforests_models = []
         cv_scores_list = []
         cv errors = []
         # iterating each number in list
         for num in n estimators list:
             rf clf = RandomForestClassifier(n estimators=num, max leaf nodes = 10, random s
             randomforests models.append(rf clf)
         #train model with cv of 10
         for rf in randomforests models:
             cv scores = cross val score(rf, simpleTrain, trainDigits, cv=10)
             cv_scores_list.append(cv_scores)
         #print each cv score (accuracy) and average them
         for i in range(len(n estimators list)):
             print('n estimators:', n_estimators_list[i])
             print(randomforests_models[i])
             print(cv_scores_list[i])
             print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
             error = 1-np.mean(cv_scores_list[i])
             cv_errors.append(error)
             print('Cross validation error:', error)
```

```
n estimators: 5
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=10,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min weight fraction leaf=0.0, n estimators=5,
                      n jobs=None, oob score=False, random state=0, verbose=0,
                      warm start=False)
[0.96875 0.9375
                    0.90625 0.96774194 0.83870968 0.90322581
0.96774194 1.
                     0.96774194 0.86666667]
cv scores mean: 0.9324327956989247
Cross validation error: 0.06756720430107532
n estimators: 10
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max depth=None, max features='auto', max leaf nodes=10,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=10,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[0.96875 0.96875
                      0.96875
                               0.96774194 0.90322581 0.90322581
0.96774194 1.
                      0.93548387 0.9
cv scores mean: 0.9483669354838711
Cross validation error: 0.051633064516128924
n estimators: 15
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max depth=None, max features='auto', max leaf nodes=10,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=15,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[0.96875 0.9375
                     1. 0.96774194 0.87096774 0.90322581
                      0.96774194 0.9
0.90322581 1.
cv scores mean: 0.9419153225806453
Cross validation error: 0.05808467741935475
n estimators: 20
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=10,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min samples leaf=1, min samples split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=20,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[0.96875 0.96875
                      0.96875 0.96774194 0.87096774 0.90322581
0.93548387 1.
                      0.96774194 0.9
cv scores mean:0.945141129032258
Cross validation error: 0.054858870967742
n estimators: 30
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=10,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=30,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
                      0.96875 0.96774194 0.83870968 0.90322581
[0.96875 0.96875
0.96774194 1.
                      0.96774194 0.9
cv scores mean: 0.945141129032258
Cross validation error: 0.054858870967742
n estimators: 40
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=10,
                      min_impurity_decrease=0.0, min_impurity_split=None,
```

```
In [93]: #plot the data points
         ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
         fig = mp.figure()
         mp.scatter(n_estimators_list,cv_errors,s=10)
         mp.xscale('log')
         mp.plot(n estimators list,cv errors)
         #specify the axes
         mp.xlabel("No of trees")
         mp.ylabel("CV Error")
         #Labeling the plot
         fig.suptitle('Figure 4.3 - Random Forests with varying n estimators with max leaf n
         odes=10', fontsize=15)
         #table
         figb t = mp.figure()
         table vals b = []
         np_array = np.array(cv_errors)
         np_round_to_tenths = np.around(np_array, 5)
         round to tenths = list(np round to tenths)
         table vals b.append(round to tenths)
         row_labels = ['cv_error']
         col_labels = n_estimators_list
         the table = mp.table(cellText=table vals b,
                               rowLabels=row labels,
                               colLabels=col_labels,
                               loc='center')
         the table.auto set font size(False)
         the_table.set_fontsize(14)
         the_table.scale(3,3)
         # Removing ticks and spines enables you to get the figure only with table
         mp.tick_params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
         mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
         for pos in ['right','top','bottom','left']:
             mp.gca().spines[pos].set visible(False)
         mp.savefig('matplotlib-table.png', bbox inches='tight', pad inches=0.05)
         #display the current graph
         mp.show()
```

Figure 4.3 - Random Forests with varying n_estimators with max leaf nodes=10 $\,$



	5	10	15	20	30	40	50	75	100	200	500	1000	10000
cv_error	0.06757	0.05163	0.05808	0.05486	0.05486	0.05798	0.05808	0.05798	0.04841	0.04841	0.05153	0.04518	0.04841

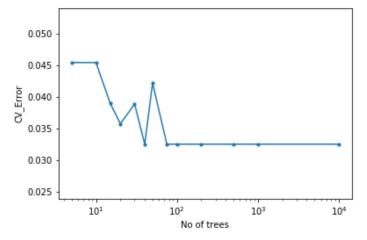
1e) PART 2 : RANDOM FORESTS - max leaf nodes = 100

```
In [94]: #create a new decision tree model
         n_{estimators_list} = [5, 10, 15, 20, 30, 40, 50, 75, 100, 200, 500, 1000, 10000]
         randomforests_models = []
         cv_scores_list = []
         cv errors = []
         # iterating each number in list
         for num in n estimators list:
             rf clf = RandomForestClassifier(n estimators=num, max leaf nodes = 100, random
             randomforests models.append(rf clf)
         #train model with cv of 10
         for rf in randomforests models:
             cv scores = cross val score(rf, simpleTrain, trainDigits, cv=10)
             cv_scores_list.append(cv_scores)
         #print each cv score (accuracy) and average them
         for i in range(len(n estimators list)):
             print('n estimators:', n_estimators_list[i])
             print(randomforests_models[i])
             print(cv_scores_list[i])
             print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
             error = 1-np.mean(cv_scores_list[i])
             cv_errors.append(error)
             print('Cross validation error:', error)
```

```
n estimators: 5
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=100,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min weight fraction leaf=0.0, n estimators=5,
                      n jobs=None, oob score=False, random state=0, verbose=0,
                      warm start=False)
[0.96875 1.
                      0.96875 0.96774194 0.87096774 0.93548387
          1.
                      0.96774194 0.86666667]
cv scores mean: 0.9546102150537635
Cross validation error: 0.04538978494623647
n estimators: 10
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                      max depth=None, max features='auto', max leaf nodes=100,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=10,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[1.
           1.
                      0.96875
                                 0.96774194 0.83870968 0.93548387
           1.
                      0.93548387 0.9
1.
cv scores mean: 0.954616935483871
Cross validation error: 0.045383064516128946
n estimators: 15
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max depth=None, max features='auto', max leaf nodes=100,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=15,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[1.
           0.96875
                      1. 0.96774194 0.87096774 0.93548387
                      0.96774194 0.9
           1.
cv scores mean: 0.9610685483870969
Cross validation error: 0.03893145161290312
n estimators: 20
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=100,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min samples leaf=1, min samples split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=20,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
                           0.96774194 0.90322581 0.93548387
           0.96875
                      0.96774194 0.9
           1.
cv scores mean: 0.9642943548387096
Cross validation error: 0.03570564516129038
n estimators: 30
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=100,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=30,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[0.96875 0.96875
                            0.96774194 0.90322581 0.93548387
          1.
                      0.96774194 0.9
cv scores mean: 0.9611693548387097
Cross validation error: 0.03883064516129031
n estimators: 40
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=100,
                      min_impurity_decrease=0.0, min_impurity_split=None,
```

```
In [95]: #plot the data points
         ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
         fig = mp.figure()
         mp.scatter(n_estimators_list,cv_errors,s=10)
         mp.xscale('log')
         mp.plot(n estimators list,cv errors)
         #specify the axes
         mp.xlabel("No of trees")
         mp.ylabel("CV Error")
         #Labeling the plot
         fig.suptitle('Figure 4.4 - Random Forests with varying n estimators with max leaf n
         odes=100', fontsize=15)
         #table
         figb t = mp.figure()
         table vals b = []
         np_array = np.array(cv_errors)
         np_round_to_tenths = np.around(np_array, 5)
         round to tenths = list(np round to tenths)
         table vals b.append(round to tenths)
         row_labels = ['cv_error']
         col_labels = n_estimators_list
         the table = mp.table(cellText=table vals b,
                               rowLabels=row labels,
                               colLabels=col_labels,
                               loc='center')
         the table.auto set font size(False)
         the_table.set_fontsize(14)
         the_table.scale(3,3)
         # Removing ticks and spines enables you to get the figure only with table
         mp.tick_params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
         mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
         for pos in ['right','top','bottom','left']:
             mp.gca().spines[pos].set visible(False)
         mp.savefig('matplotlib-table.png', bbox inches='tight', pad inches=0.05)
         #display the current graph
         mp.show()
```

Figure 4.4 - Random Forests with varying n_estimators with max leaf nodes=100



	5	10	15	20	30	40	50	75	100	200	500	1000	10000
cv_error	0.04539	0.04538	0.03893	0.03571	0.03883	0.03248	0.04216	0.03248	0.03248	0.03248	0.03248	0.03248	0.03248

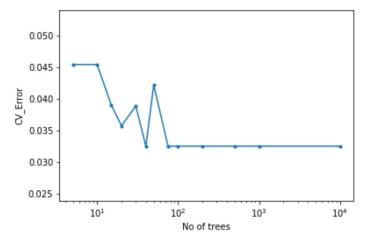
1e) PART 3: RANDOM FORESTS - max leaf nodes = 1000

```
In [96]: #create a new decision tree model
         n_{estimators_list} = [5, 10, 15, 20, 30, 40, 50, 75, 100, 200, 500, 1000, 10000]
         randomforests_models = []
         cv_scores_list = []
         cv errors = []
         # iterating each number in list
         for num in n estimators list:
             rf clf = RandomForestClassifier(n estimators=num, max leaf nodes = 1000, random
             randomforests models.append(rf clf)
         #train model with cv of 10
         for rf in randomforests models:
             cv scores = cross val score(rf, simpleTrain, trainDigits, cv=10)
             cv_scores_list.append(cv_scores)
         #print each cv score (accuracy) and average them
         for i in range(len(n estimators list)):
             print('n estimators:', n_estimators_list[i])
             print(randomforests_models[i])
             print(cv_scores_list[i])
             print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
             error = 1-np.mean(cv_scores_list[i])
             cv_errors.append(error)
             print('Cross validation error:', error)
```

```
n estimators: 5
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=1000,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min weight fraction leaf=0.0, n estimators=5,
                      n jobs=None, oob score=False, random state=0, verbose=0,
                      warm start=False)
[0.96875 1.
                      0.96875 0.96774194 0.87096774 0.93548387
          1.
                      0.96774194 0.86666667]
cv scores mean: 0.9546102150537635
Cross validation error: 0.04538978494623647
n estimators: 10
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                      max depth=None, max features='auto', max leaf nodes=1000,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=10,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[1.
           1.
                      0.96875
                                 0.96774194 0.83870968 0.93548387
           1.
                      0.93548387 0.9
1.
cv scores mean: 0.954616935483871
Cross validation error: 0.045383064516128946
n estimators: 15
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max depth=None, max features='auto', max leaf nodes=1000,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=15,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[1.
           0.96875
                      1. 0.96774194 0.87096774 0.93548387
                      0.96774194 0.9
           1.
cv scores mean: 0.9610685483870969
Cross validation error: 0.03893145161290312
n estimators: 20
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=1000,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min samples leaf=1, min samples split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=20,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
                           0.96774194 0.90322581 0.93548387
           0.96875
                      0.96774194 0.9
           1.
cv scores mean: 0.9642943548387096
Cross validation error: 0.03570564516129038
n estimators: 30
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=1000,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=30,
                      n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[0.96875 0.96875
                            0.96774194 0.90322581 0.93548387
          1.
                      0.96774194 0.9
cv scores mean: 0.9611693548387097
Cross validation error: 0.03883064516129031
n estimators: 40
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max_depth=None, max_features='auto', max_leaf_nodes=1000,
                      min_impurity_decrease=0.0, min_impurity_split=None,
```

```
In [97]: #plot the data points
         ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
         fig = mp.figure()
         mp.scatter(n_estimators_list,cv_errors,s=10)
         mp.xscale('log')
         mp.plot(n estimators list,cv errors)
         #specify the axes
         mp.xlabel("No of trees")
         mp.ylabel("CV Error")
         #Labeling the plot
         fig.suptitle('Figure 4.5 - Random Forests with varying n estimators with max leaf n
         odes=1000', fontsize=15)
         #table
         figb t = mp.figure()
         table vals b = []
         np_array = np.array(cv_errors)
         np_round_to_tenths = np.around(np_array, 5)
         round to tenths = list(np round to tenths)
         table vals b.append(round to tenths)
         row_labels = ['cv_error']
         col_labels = n_estimators_list
         the table = mp.table(cellText=table vals b,
                               rowLabels=row labels,
                               colLabels=col_labels,
                               loc='center')
         the table.auto set font size(False)
         the_table.set_fontsize(14)
         the_table.scale(3,3)
         # Removing ticks and spines enables you to get the figure only with table
         mp.tick_params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
         mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
         for pos in ['right','top','bottom','left']:
             mp.gca().spines[pos].set visible(False)
         mp.savefig('matplotlib-table.png', bbox inches='tight', pad inches=0.05)
         #display the current graph
         mp.show()
```

Figure 4.5 - Random Forests with varying n_estimators with max leaf nodes=1000



	5	10	15	20	30	40	50	75	100	200	500	1000	10000
cv_error	0.04539	0.04538	0.03893	0.03571	0.03883	0.03248	0.04216	0.03248	0.03248	0.03248	0.03248	0.03248	0.03248

1f) Max leaf nodes - Most impacted by bagging

The max leaf nodes with 10 has been most impacted by the bagging approach. The max leaf nodes for 100 and 1000 do not have much impact after the n_estimators reach 75. If max_leaf nodes is less, there is a high chance of having different decision trees that give different results in each bag of n_estimators. This is because the depth is limited and we might not able able to reach the end solution with perfect class labels and hence the results might vary. But if we see for 100 and 1000 max leaf nodes, the results don't vary much after n_estimators reach 75 i.e., the cross validation errors are same. This might be because the trees are fully grown and fully grown trees in all bags are able to predict data similarly than half grown trees which can yield varying results.

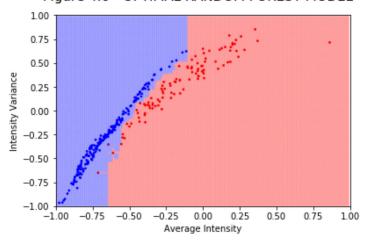
1g) N Estimators - Impact

Even though we find that the graph vary much with n_estimators, in real there is actually very slight difference between them and the difference reduces more as we reach higher values of n_estimators. But we can say that as n_estimators increase and reach a ceratin point, the cross validation errors do not change much. Yes, there was a point were n_estimators didn't have effect that is after n_estimators = 75 for both 100 and 1000 max_leaf_nodes. This is beacuse, once we build certain maximum nuber of trees, we reach a point were the different possible trees are built and including more trees doesn't change our results. This holds even more true if the max leaf nodes count is more.

1h) Optimal model - Random forests

```
In [200]: # create the model
          # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCl
          assifier.html
          # Declare Model
          model = RandomForestClassifier(n estimators=75, max leaf nodes=100, random state=
          0, n jobs=-1)
          # Fit model to our data
          model.fit(simpleTrain, trainDigits)
          # Lists to hold inpoints, predictions and assigned colors
          xPred = []
          yPred = []
          cPred = []
          # Use input points to get predictions here
          for xP in range(-100,100):
              xP = xP/100.0
              for yP in range(-100,100):
                  yP = yP/100.0
                  xPred.append(xP)
                  yPred.append(yP)
                  if (model.predict([[xP, yP]]) == "1.0"):
                       cPred.append("b")
                  else:
                       cPred.append("r")
```

Figure 4.6 - OPTIMAL RANDOM FOREST MODEL



Answer: The optimal Model for Random forest I've taken as max_leaf_nodes = 100 and n_estimators = 200. Max leaf nodes 100 and 1000 have same results and hence I've choosen 100 as the value. Having more leaf nodes can increase the depth and overfit the model (even though it's not the case here)so I chose 100 as an optimal value. And n_estimators from 75 till 10000 have same values. Having more number of bags doesn't overfit the model in bagging. But having too many bags(i.e., n_estimators) can increase the run_time. So I've taken 200 as an optimal value between 75 and 10000.

1i) Random forests - Max features

```
In [193]: #create a new decision tree model
          max_leaf_nodes = [10, 100, 1000]
          n_estimators_list = [1,10,100,100]
          max_features = [0.1, 0.2, 0.3, 0.4, 0.5, 0.6, 0.7, 0.8, 0.9, 1.0]
          models = []
          actual max leaf nodes = []
          actual n estimators = []
          actual max features = []
          randomforests models = []
          cv scores list = []
          cv errors = []
          # iterating each number in list
          for leaf in max leaf nodes:
              for estimator in n_estimators_list:
                  for feature in max_features:
                      rf clf = RandomForestClassifier(max leaf nodes = leaf, n estimators=es
          timator, max features= feature, random state=0)
                      randomforests models.append(rf clf)
                      actual_max_leaf_nodes.append(leaf)
                      actual_n_estimators.append(estimator)
                      actual max features.append(feature)
          #train model with cv of 10
          for rf in randomforests_models:
              models.append(randomforests_models.index(rf))
              cv_scores = cross_val_score(rf, simpleTrain, trainDigits, cv=10)
              cv_scores_list.append(cv_scores)
          #print each cv score (accuracy) and average them
          for i in range(len(randomforests models)):
              print(randomforests_models[i])
              print(cv_scores_list[i])
              print('cv scores mean:{}'.format(np.mean(cv scores list[i])))
              error = 1-np.mean(cv scores list[i])
              cv_errors.append(error)
              print('Cross validation error:', error)
```

4/23/2020, 11:37 PM

```
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max depth=None, max features=0.1, max leaf nodes=10,
                      min impurity decrease=0.0, min impurity split=None,
                      min_samples_leaf=1, min_samples_split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=1,
                       n_jobs=None, oob_score=False, random_state=0, verbose=0,
                       warm start=False)
                                0.96774194 0.90322581 0.90322581
0.96774194 0.93548387 0.96774194 0.86666667]
cv scores mean: 0.9355577956989247
Cross validation error: 0.06444220430107528
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                      max_depth=None, max_features=0.2, max_leaf_nodes=10,
                      min impurity decrease=0.0, min impurity split=None,
                       min samples leaf=1, min samples split=2,
                       min weight fraction leaf=0.0, n estimators=1,
                       n_jobs=None, oob_score=False, random_state=0, verbose=0,
                       warm start=False)
[0.96875 1.
                      0.875
                                 0.96774194 0.90322581 0.90322581
0.96774194 0.93548387 0.96774194 0.86666667]
cv scores mean: 0.9355577956989247
Cross validation error: 0.06444220430107528
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                       max depth=None, max features=0.3, max leaf nodes=10,
                       min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                       min weight fraction leaf=0.0, n estimators=1,
                       n jobs=None, oob score=False, random state=0, verbose=0,
                       warm start=False)
[0.96875 1.
                              0.96774194 0.90322581 0.90322581
                      0.875
0.96774194 0.93548387 0.96774194 0.86666667]
cv scores mean: 0.9355577956989247
Cross validation error: 0.06444220430107528
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                       max depth=None, max features=0.4, max leaf nodes=10,
                       min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=2,
                      min_weight_fraction_leaf=0.0, n_estimators=1,
                       n_jobs=None, oob_score=False, random_state=0, verbose=0,
                      warm start=False)
[0.96875 1.
                      0.875
                             0.96774194 0.90322581 0.90322581
0.96774194 0.93548387 0.96774194 0.86666667]
cv_scores mean:0.9355577956989247
Cross validation error: 0.06444220430107528
RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
                       max_depth=None, max_features=0.5, max_leaf_nodes=10,
                      min_impurity_decrease=0.0, min_impurity_split=None,
                       min samples leaf=1, min samples split=2,
                       min weight fraction leaf=0.0, n estimators=1,
                       n_jobs=None, oob_score=False, random_state=0, verbose=0,
                       warm_start=False)
                      0.875
[0.96875 1.
                               0.96774194 0.90322581 0.90322581
0.96774194 0.93548387 0.96774194 0.86666667]
cv scores mean: 0.9355577956989247
Cross validation error: 0.06444220430107528
RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
                      max depth=None, max features=0.6, max leaf nodes=10,
                       min impurity decrease=0.0, min impurity split=None,
                       min samples leaf=1, min samples split=2,
                      min weight fraction leaf=0.0, n estimators=1,
                       n jobs=None, oob score=False, random state=0, verbose=0,
                       warm start=False)
[0.96875
                      0.875
                               0.96774194 0.90322581 0.90322581
           1.
 0.96774194 0.93548387 0.96774194 0.86666667]
```

Out[194]:

	cv_errors	actual_max_leaf_nodes	actual_n_estimators	actual_max_features
0	0.064442	10	1	0.1
1	0.064442	10	1	0.2
2	0.064442	10	1	0.3
3	0.064442	10	1	0.4
4	0.064442	10	1	0.5
5	0.064442	10	1	0.6
6	0.064442	10	1	0.7
7	0.064442	10	1	0.8
8	0.064442	10	1	0.9
9	0.051633	10	1	1.0
10	0.051633	10	10	0.1
11	0.051633	10	10	0.2
12	0.051633	10	10	0.3
13	0.051633	10	10	0.4
14	0.051633	10	10	0.5
15	0.051633	10	10	0.6
16	0.051633	10	10	0.7
17	0.051633	10	10	0.8
18	0.051633	10	10	0.9
19	0.035605	10	10	1.0
20	0.048407	10	100	0.1
21	0.048407	10	100	0.2
22	0.048407	10	100	0.3
23	0.048407	10	100	0.4
24	0.048407	10	100	0.5
25	0.048407	10	100	0.6
26	0.048407	10	100	0.7
27	0.048407	10	100	0.8
28	0.048407	10	100	0.9
29	0.038730	10	100	1.0
90	0.045383	1000	10	0.1
91	0.045383	1000	10	0.2
92	0.045383	1000	10	0.3
93	0.045383	1000	10	0.4
94	0.045383	1000	10	0.5
95	0.045383	1000	10	0.6
96	0.045383	1000	10	0.7
97	0.045383	1000	10	0.8
98	0.045383	1000	10	0.9

The lowest cross validation error occured at max_leaf_nodes = 100, n_estimators = 100, max_features = 0.1

ADABOOST

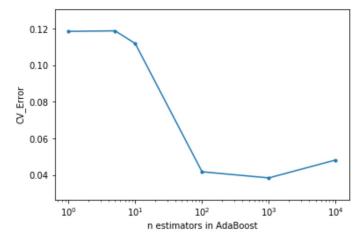
1j) ADABOOST CLASSIFIER - DecisionTreeClassifier(max_depth=1)

```
In [20]: #create a new decision tree model
         n_{estimators_list} = [1, 5, 10, 100, 1000, 10000]
         adaboost_models = []
         cv_scores_list = []
         cv errors = []
         # iterating each number in list
         for num in n estimators list:
             ada clf = AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=
         1), n estimators=num, random state=0)
             adaboost models.append(ada clf)
          #train model with cv of 10
         for ada in adaboost models:
             cv_scores = cross_val_score(ada, simpleTrain, trainDigits, cv=10)
             cv_scores_list.append(cv_scores)
          #print each cv score (accuracy) and average them
         for i in range(len(n estimators list)):
             print('n estimators:', n_estimators_list[i])
             print(adaboost_models[i])
             print(cv_scores_list[i])
             print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
             error = 1-np.mean(cv_scores_list[i])
             cv_errors.append(error)
             print('Cross validation error:', error)
```

```
n estimators: 1
AdaBoostClassifier(algorithm='SAMME.R',
                   base estimator=DecisionTreeClassifier(class weight=None,
                                                          criterion='gini',
                                                         max_depth=1,
                                                         max features=None,
                                                         max leaf nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min impurity split=Non
                                                         min samples leaf=1,
                                                         min_samples_split=2,
                                                         min_weight_fraction_lea
f=0.0,
                                                          presort=False,
                                                         random state=None,
                                                          splitter='best'),
                   learning_rate=1.0, n_estimators=1, random_state=0)
            0.84375 0.90625 1.
[0.875
                                             0.80645161 0.83870968
0.87096774 0.96774194 0.83870968 0.86666667]
cv_scores mean:0.8814247311827957
Cross validation error: 0.11857526881720426
n estimators: 5
AdaBoostClassifier(algorithm='SAMME.R',
                   base estimator=DecisionTreeClassifier(class weight=None,
                                                          criterion='gini',
                                                         max depth=1,
                                                         max features=None,
                                                         max leaf nodes=None,
                                                         min impurity decrease=
0.0,
                                                         min impurity split=Non
e,
                                                         min samples leaf=1,
                                                         min samples split=2,
                                                         min weight fraction lea
f=0.0,
                                                          presort=False,
                                                          random state=None,
                                                          splitter='best'),
                   learning rate=1.0, n estimators=5, random state=0)
[0.875
            0.90625
                     0.9375 0.90322581 0.87096774 0.80645161
0.87096774 0.90322581 0.87096774 0.86666667]
cv scores mean: 0.881122311827957
Cross validation error: 0.11887768817204303
n estimators: 10
AdaBoostClassifier(algorithm='SAMME.R',
                   base estimator=DecisionTreeClassifier(class weight=None,
                                                          criterion='gini',
                                                         max_depth=1,
                                                         max_features=None,
                                                         max_leaf_nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min_impurity_split=Non
e,
                                                         min samples leaf=1,
                                                         min samples split=2,
                                                         min weight fraction lea
f=0.0,
                                                          presort=False,
                                                          random state=None,
                                                          splitter='best'),
```

```
In [22]: #plot the data points
         ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
         fig = mp.figure()
         mp.scatter(n_estimators_list,cv_errors,s=10)
         mp.xscale('log')
         mp.plot(n estimators list,cv errors)
         #specify the axes
         mp.xlabel("n estimators in AdaBoost")
         mp.ylabel("CV Error")
         #Labeling the plot
         fig.suptitle('Figure 4.7 - AdaBoost model with DecisionTreeClassifier(max depth=
         1)', fontsize=15)
         #table
         figb t = mp.figure()
         table vals b = []
         np_array = np.array(cv_errors)
         np_round_to_tenths = np.around(np_array, 5)
         round to tenths = list(np round to tenths)
         table vals b.append(round to tenths)
         row_labels = ['cv_error']
         col_labels = n_estimators_list
         the table = mp.table(cellText=table vals b,
                               rowLabels=row labels,
                               colLabels=col_labels,
                               loc='center')
         the table.auto set font size (False)
         the_table.set_fontsize(14)
         the table.scale(2,2)
         # Removing ticks and spines enables you to get the figure only with table
         mp.tick_params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
         mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
         for pos in ['right','top','bottom','left']:
             mp.gca().spines[pos].set visible(False)
         mp.savefig('matplotlib-table.png', bbox inches='tight', pad inches=0.05)
         #display the current graph
         mp.show()
```

Figure 4.7 - AdaBoost model with DecisionTreeClassifier(max_depth=1)



	1	5	10	100	1000	10000
cv_error	0.11858	0.11888	0.11192	0.04175	0.03842	0.04821

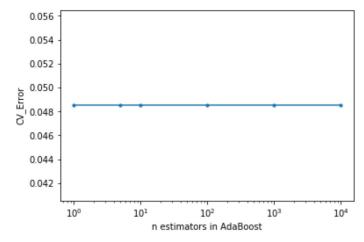
1k) ADABOOST CLASSIFIER - DecisionTreeClassifier(max_depth=10)

```
In [23]: #create a new decision tree model
         n_{estimators_list} = [1, 5, 10, 100, 1000, 10000]
         adaboost_models = []
         cv_scores_list = []
         cv errors = []
         # iterating each number in list
         for num in n estimators list:
             ada clf = AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=1
         0), n estimators=num, random state=0)
             adaboost models.append(ada clf)
          #train model with cv of 10
         for ada in adaboost models:
             cv_scores = cross_val_score(ada, simpleTrain, trainDigits, cv=10)
             cv_scores_list.append(cv_scores)
          #print each cv score (accuracy) and average them
         for i in range(len(n estimators list)):
             print('n estimators:', n_estimators_list[i])
             print(adaboost_models[i])
             print(cv_scores_list[i])
             print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
             error = 1-np.mean(cv_scores_list[i])
             cv_errors.append(error)
             print('Cross validation error:', error)
```

```
n estimators: 1
AdaBoostClassifier(algorithm='SAMME.R',
                   base estimator=DecisionTreeClassifier(class weight=None,
                                                         criterion='gini',
                                                         max_depth=10,
                                                         max features=None,
                                                         max leaf nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min impurity split=Non
                                                         min samples leaf=1,
                                                         min_samples_split=2,
                                                         min_weight_fraction_lea
f=0.0,
                                                         presort=False,
                                                         random state=None,
                                                         splitter='best'),
                   learning_rate=1.0, n_estimators=1, random_state=0)
[0.9375
           1.
                 1. 0.96774194 0.93548387 0.83870968
1.
           0.96774194 0.96774194 0.9
cv_scores mean:0.951491935483871
Cross validation error: 0.04850806451612899
n estimators: 5
AdaBoostClassifier(algorithm='SAMME.R',
                  base estimator=DecisionTreeClassifier(class weight=None,
                                                         criterion='gini',
                                                         max depth=10,
                                                         max features=None,
                                                         max leaf nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min impurity split=Non
e,
                                                         min samples leaf=1,
                                                         min samples split=2,
                                                         min weight fraction lea
f=0.0,
                                                         presort=False,
                                                         random state=None,
                                                         splitter='best'),
                   learning_rate=1.0, n_estimators=5, random state=0)
[0.9375
                  1. 0.96774194 0.93548387 0.83870968
           0.96774194 0.96774194 0.9
                                           ]
cv scores mean: 0.951491935483871
Cross validation error: 0.04850806451612899
n estimators: 10
AdaBoostClassifier(algorithm='SAMME.R',
                  base estimator=DecisionTreeClassifier(class weight=None,
                                                         criterion='gini',
                                                         max_depth=10,
                                                         max_features=None,
                                                         max_leaf_nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min_impurity_split=Non
e,
                                                         min samples leaf=1,
                                                         min samples split=2,
                                                         min_weight_fraction_lea
f=0.0,
                                                         presort=False,
                                                         random state=None,
                                                         splitter='best'),
```

```
In [24]: #plot the data points
         ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
         fig = mp.figure()
         mp.scatter(n_estimators_list,cv_errors,s=10)
         mp.xscale('log')
         mp.plot(n estimators list,cv errors)
         #specify the axes
         mp.xlabel("n estimators in AdaBoost")
         mp.ylabel("CV Error")
         #Labeling the plot
         fig.suptitle('Figure 4.8 - AdaBoost model with DecisionTreeClassifier(max depth=1
         0)', fontsize=15)
         #table
         figb t = mp.figure()
         table vals b = []
         np_array = np.array(cv_errors)
         np_round_to_tenths = np.around(np_array, 5)
         round to tenths = list(np round to tenths)
         table vals b.append(round to tenths)
         row_labels = ['cv_error']
         col_labels = n_estimators_list
         the table = mp.table(cellText=table vals b,
                               rowLabels=row labels,
                               colLabels=col_labels,
                               loc='center')
         the table.auto set font size (False)
         the_table.set_fontsize(14)
         the table.scale(2,2)
         # Removing ticks and spines enables you to get the figure only with table
         mp.tick_params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
         mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
         for pos in ['right','top','bottom','left']:
             mp.gca().spines[pos].set visible(False)
         mp.savefig('matplotlib-table.png', bbox inches='tight', pad inches=0.05)
         #display the current graph
         mp.show()
```

Figure 4.8 - AdaBoost model with DecisionTreeClassifier(max_depth=10)



	1	5	10	100	1000	10000	
cv_error	0.04851	0.04851	0.04851	0.04851	0.04851	0.04851	

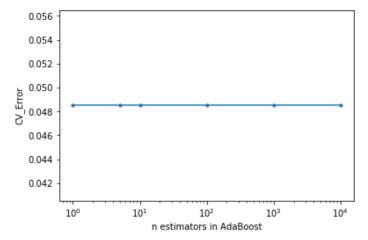
1I) ADABOOST CLASSIFIER - DecisionTreeClassifier(max_depth=1000)

```
In [25]: #create a new decision tree model
         n_{estimators_list} = [1, 5, 10, 100, 1000, 10000]
         adaboost_models = []
         cv_scores_list = []
         cv errors = []
         # iterating each number in list
         for num in n estimators list:
             ada clf = AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=10
         00), n estimators=num, random state=0)
             adaboost models.append(ada clf)
          #train model with cv of 10
         for ada in adaboost models:
             cv_scores = cross_val_score(ada, simpleTrain, trainDigits, cv=10)
             cv_scores_list.append(cv_scores)
          #print each cv score (accuracy) and average them
         for i in range(len(n estimators list)):
             print('n estimators:', n_estimators_list[i])
             print(adaboost_models[i])
             print(cv_scores_list[i])
             print('cv_scores mean:{}'.format(np.mean(cv_scores_list[i])))
             error = 1-np.mean(cv_scores_list[i])
             cv_errors.append(error)
             print('Cross validation error:', error)
```

```
n estimators: 1
AdaBoostClassifier(algorithm='SAMME.R',
                   base estimator=DecisionTreeClassifier(class weight=None,
                                                         criterion='gini',
                                                         max_depth=1000,
                                                         max features=None,
                                                         max leaf nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min impurity split=Non
                                                         min samples leaf=1,
                                                         min_samples_split=2,
                                                         min_weight_fraction_lea
f=0.0,
                                                         presort=False,
                                                         random state=None,
                                                         splitter='best'),
                   learning_rate=1.0, n_estimators=1, random_state=0)
[0.9375
           1.
                 1. 0.96774194 0.93548387 0.83870968
1.
           0.96774194 0.96774194 0.9
cv_scores mean:0.951491935483871
Cross validation error: 0.04850806451612899
n estimators: 5
AdaBoostClassifier(algorithm='SAMME.R',
                  base estimator=DecisionTreeClassifier(class weight=None,
                                                         criterion='gini',
                                                         max depth=1000,
                                                         max features=None,
                                                         max leaf nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min impurity split=Non
e,
                                                         min samples leaf=1,
                                                         min samples split=2,
                                                         min weight fraction lea
f=0.0,
                                                         presort=False,
                                                         random state=None,
                                                         splitter='best'),
                   learning_rate=1.0, n_estimators=5, random state=0)
[0.9375
                  1. 0.96774194 0.93548387 0.83870968
           0.96774194 0.96774194 0.9
                                           ]
cv scores mean: 0.951491935483871
Cross validation error: 0.04850806451612899
n estimators: 10
AdaBoostClassifier(algorithm='SAMME.R',
                  base estimator=DecisionTreeClassifier(class weight=None,
                                                         criterion='gini',
                                                         max_depth=1000,
                                                         max_features=None,
                                                         max_leaf_nodes=None,
                                                         min_impurity_decrease=
0.0,
                                                         min_impurity_split=Non
e,
                                                         min samples leaf=1,
                                                         min samples split=2,
                                                         min weight fraction lea
f=0.0,
                                                         presort=False,
                                                         random state=None,
                                                         splitter='best'),
```

```
In [26]: #plot the data points
         ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
         fig = mp.figure()
         mp.scatter(n_estimators_list,cv_errors,s=10)
         mp.xscale('log')
         mp.plot(n estimators list,cv errors)
         #specify the axes
         mp.xlabel("n estimators in AdaBoost")
         mp.ylabel("CV Error")
         #Labeling the plot
         fig.suptitle('Figure 4.9 - AdaBoost model with DecisionTreeClassifier(max depth=100
         0)', fontsize=15)
         #table
         figb t = mp.figure()
         table vals b = []
         np_array = np.array(cv_errors)
         np_round_to_tenths = np.around(np_array, 5)
         round to tenths = list(np round to tenths)
         table vals b.append(round to tenths)
         row_labels = ['cv_error']
         col_labels = n_estimators_list
         the table = mp.table(cellText=table vals b,
                               rowLabels=row labels,
                               colLabels=col_labels,
                               loc='center')
         the table.auto set font size (False)
         the_table.set_fontsize(14)
         the table.scale(2,2)
         # Removing ticks and spines enables you to get the figure only with table
         mp.tick_params(axis='x', which='both', bottom=False, top=False, labelbottom=False)
         mp.tick_params(axis='y', which='both', right=False, left=False, labelleft=False)
         for pos in ['right','top','bottom','left']:
             mp.gca().spines[pos].set visible(False)
         mp.savefig('matplotlib-table.png', bbox inches='tight', pad inches=0.05)
         #display the current graph
         mp.show()
```

Figure 4.9 - AdaBoost model with DecisionTreeClassifier(max_depth=1000)

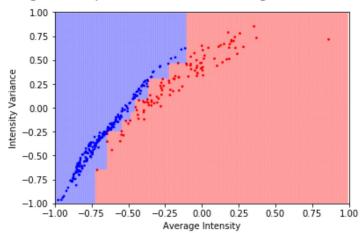


	1	5	10	100	1000	10000	
cv_error	0.04851	0.04851	0.04851	0.04851	0.04851	0.04851	

1m) Optimal model decision region - ADABOOST

```
In [27]: # create the model
          # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCla
         ssifier.html
          # Declare Model
         model = AdaBoostClassifier(base_estimator=DecisionTreeClassifier(max_depth=1), n_es
         timators=1000, random_state=0)
         # Fit model to our data
         model.fit(simpleTrain,trainDigits)
         # Lists to hold inpoints, predictions and assigned colors
         xPred = []
         yPred = []
         cPred = []
          # Use input points to get predictions here
         for xP in range(-100,100):
             xP = xP/100.0
             for yP in range(-100,100):
                 yP = yP/100.0
                 xPred.append(xP)
                  yPred.append(yP)
                  if (model.predict([[xP,yP]]) =="1.0"):
                      cPred.append("b")
                  else:
                      cPred.append("r")
```

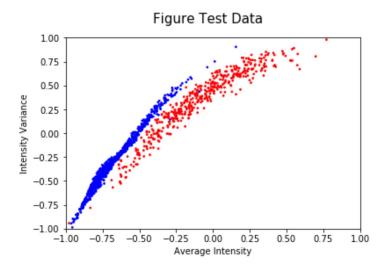
Fig 4.10 - Optimal model decision region - ADABOOST



Answer: Optimal decision region plotted above is for base estimator = DecisionTreeClassifier(max_depth=1) and n_estimators=1000. The graph plotted above also suggests there is no overfitting and underfitting occurence.

2. REPORTING FINAL ERROR

```
In [136]: #Convert the 256D data (testFeatures) to 2D data
          #We need X and Y for plotting and simpleTest for testing the model.
          #They contain the same points in a different arrangement
          X t = []
          Y t = []
          simpleTest = []
          #Colors will be passed to the graphing library to color the points.
          #1's are blue: "b" and 5's are red: "r"
          t colors = []
          \#legends = []
          for index in range(len(testFeatures)):
              #print(index)
              #produce the 2D dataset for graphing/training and scale the data so it is in t
          he [-1,1] square
              xNew = 2*np.average(testFeatures[index])+.75
              yNew = 3*np.var(testFeatures[index])-1.5
              X t.append(xNew)
              Y t.append(yNew)
              simpleTest.append([xNew,yNew])
              #trainDigits will still be the value we try to classify. Here it is the string
          "1.0" or "5.0"
              if(testDigits[index] == "1.0"):
                  t colors.append("b")
                  #legends.append("1")
                  t colors.append("r")
                  #legends.append("5")
          #plot the data points
          ### https://matplotlib.org/api/_as_gen/matplotlib.pyplot.scatter.html
          fig = mp.figure()
          mp.scatter(X t,Y t,s=3,c=t colors)
          #specify the axes
          mp.xlim(-1,1)
          mp.xlabel("Average Intensity")
          mp.ylim(-1,1)
          mp.ylabel("Intensity Variance")
          #Labeling the plot
          #mp.legend(['1'])
          #mp.legend(legends)
          fig.suptitle('Figure Test Data', fontsize=15)
          #display the current graph
          show()
```



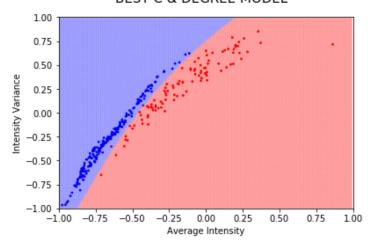
Polynomial SVM

```
In [137]: from sklearn.svm import SVC
```

Optimal Model

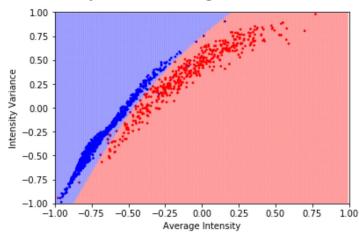
```
In [138]:
          # create the model
          # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCl
          assifier.html
          # Declare Model
          model = SVC(C=2, gamma='scale', kernel='poly', degree=3)
          # Fit model to our data
          model.fit(simpleTrain,trainDigits)
          # Lists to hold inpoints, predictions and assigned colors
          xPred = []
          yPred = []
          cPred = []
          # Use input points to get predictions here
          for xP in range(-100,100):
              xP = xP/100.0
              for yP in range(-100,100):
                  yP = yP/100.0
                  xPred.append(xP)
                  yPred.append(yP)
                  if (model.predict([[xP,yP]]) =="1.0"):
                       cPred.append("b")
                  else:
                      cPred.append("r")
```

BEST C & DEGREE MODEL



Testing data results with the above best model

Polynomial SVM - Regions on test data



```
In [142]: score = model.score(simpleTest, testDigits)
    svm_error = 1-score
    print(score)
    print(svm_error)

    0.9855884707766213
    0.014411529223378738
```

Answer: The optimal model I've choosen for Polynomial SVM is "SVC(C=2, gamma='scale', kernel='poly', degree=3)" with degree 3 and c = 2 which has a cross validation score of 0.9935. I have used this model on the entire test data and obtained a score of 0.9855 which is a very good score for the model above. Since the testing data and the training data in 2D have similar trends as we see from the above graph plots, the testing data performed well on the model too.

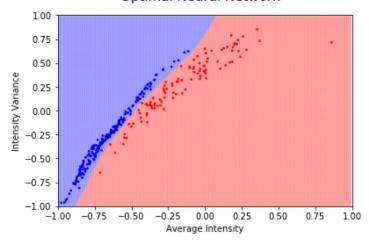
Neural Network

```
In [143]: from sklearn.neural_network import MLPClassifier
```

Optimal Model

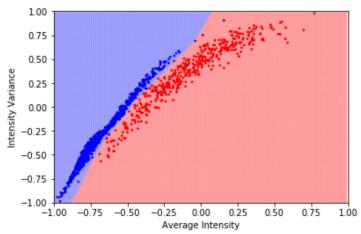
```
In [144]: # create the model
          # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCl
          assifier.html
          # Declare Model
          model = MLPClassifier(hidden layer sizes=(50,50,50,50,50), activation='relu', earl
          y stopping=False, solver='adam', alpha=0, max iter=10000, epsilon=0.001)
          # Fit model to our data
          model.fit(simpleTrain,trainDigits)
          # Lists to hold inpoints, predictions and assigned colors
          xPred = []
          yPred = []
          cPred = []
          # Use input points to get predictions here
          for xP in range(-100,100):
              xP = xP/100.0
              for yP in range(-100,100):
                  yP = yP/100.0
                  xPred.append(xP)
                  yPred.append(yP)
                  if (model.predict([[xP, yP]]) == "1.0"):
                       cPred.append("b")
                  else:
                      cPred.append("r")
```

Optimal Neural Network



Testing data results for the above optimal Neural Network model

Neural Network - Regions on test data



0.9919935948759008 0.008006405124099225

Answer: The optimal model I've choosen for Polynomial SVM is with hidden_layers=5 with 50 nodes and with relu as it's activation function. This model has a cross validation score of 0.99677. I have used this model on the entire test data and obtained a score of 0.99199 which is a very good score for the model above and has better performance than SVM. Since the testing data and the training data in 2D have similar trends as we see from the above graph plots, the testing data performed well on the model too.

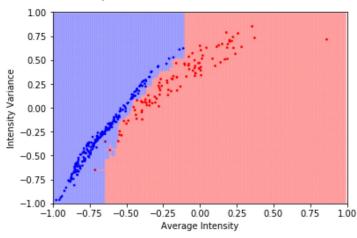
RANDOM FORESTS

```
In [149]: from sklearn.ensemble import RandomForestRegressor
```

Optimal model

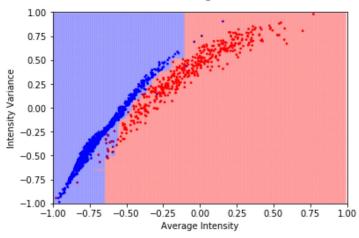
```
In [150]: # create the model
          # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCl
          assifier.html
          # Declare Model
          model = RandomForestClassifier(n estimators=75, max leaf nodes=100, random state=
          # Fit model to our data
          model.fit(simpleTrain, trainDigits)
          # Lists to hold inpoints, predictions and assigned colors
          xPred = []
          yPred = []
          cPred = []
          # Use input points to get predictions here
          for xP in range(-100,100):
              xP = xP/100.0
              for yP in range(-100,100):
                  yP = yP/100.0
                  xPred.append(xP)
                  yPred.append(yP)
                  if (model.predict([[xP,yP]]) == "1.0"):
                       cPred.append("b")
                  else:
                      cPred.append("r")
```

Optimal Random Forests Model



Testing data results for above Random forest optimal model

Random Forests - Regions on test data



Answer: Answer: The optimal model I've choosen for Random Forests is with max_leaf_nodes=100 and n_estimators=75. This model has a cross validation score of 0.96752 on the training data. I have used this model on the entire test data and obtained a score of 0.97598 which is a very good score for the model but has a slightly less performance than polynomial SVM and Neural Networks. Since the testing data and the training data in 2D have almost similar trends but mssed few points in the rectilinear decision boundaries and that's why has a slightly less performance. But overall the model does a weel job.

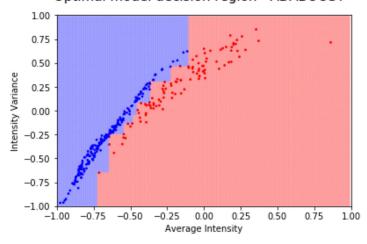
ADABOOST

```
In [155]: from sklearn.ensemble import AdaBoostClassifier
```

Optimal Model

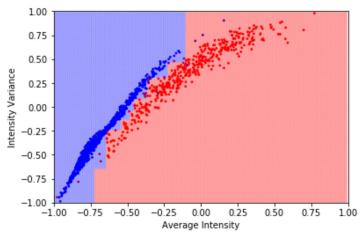
```
In [156]: # create the model
          # https://scikit-learn.org/stable/modules/generated/sklearn.neighbors.KNeighborsCl
          assifier.html
          # Declare Model
          model = AdaBoostClassifier(base estimator=DecisionTreeClassifier(max depth=1), n e
          stimators=1000, random state=0)
          # Fit model to our data
          model.fit(simpleTrain, trainDigits)
          # Lists to hold inpoints, predictions and assigned colors
          xPred = []
          yPred = []
          cPred = []
          # Use input points to get predictions here
          for xP in range(-100,100):
              xP = xP/100.0
              for yP in range(-100,100):
                  yP = yP/100.0
                  xPred.append(xP)
                  yPred.append(yP)
                  if (model.predict([[xP, yP]]) == "1.0"):
                       cPred.append("b")
                  else:
                       cPred.append("r")
```

Optimal model decision region - ADABOOST



Testing data results for above optimal Adaboost model

Adaboost - Regions on test data



```
In [160]: score = model.score(simpleTest, testDigits)
ada_error = 1-score
print(score)
print(ada_error)
```

0.9743795036028823

0.02562049639711772

Answer: The optimal model I've choosen for Adaboost is with DecisionTreeClassifier(max_depth=1) as base estimator and n_estimators=1000. This model has a highest cross validation score of 0.96157 on the training data. I have used this model on the entire test data and obtained a score of 0.97437 which is a very good score for the model. Since the testing data and the training data in 2D have almost similar trends but mssed few points in the rectilinear decision boundaries. But overall the model does a well job. We can also say that the random forests and Adaboost have performed well on test data than training data. This shows how they can reduce variance without decreasing bias.

CONCENTRATION BOUNDS

```
In [177]: import math
```

Markov Bound

```
In [178]: def markov(test_error, confidence_interval):
    prob = (100 - confidence_interval)/100
    a = test_error/prob
    return a
```

Chebyshev Bound

```
In [183]: def chebyshev(test_samples, confidence_interval):
    prob = (100 - confidence_interval)/100
    a = math.sqrt(1/(4*test_samples*prob))
    return a
```

Hoeffding Bound

Markov Bounds for all 4 models with all confidence levels

```
In [166]: confidence_intervals = [75, 95, 99]
           svm_B = []
           nn \bar{B} = []
           rf_B = []
           ada_B = []
           for ci in confidence intervals:
               svm B.append(markov(svm error, ci))
               nn B.append(markov(nn error, ci))
               rf B.append(markov(rf error, ci))
               ada B.append(markov(ada error, ci))
In [169]: markov_bounds_list = pd.DataFrame(
               {'Confidence Intervals': confidence_intervals,
                'SVM Model': svm B,
                'Neural Network Model': nn B,
                'Random Forests Model': rf B,
                'Ada Boost Model': ada B
               })
           markov bounds list
Out[169]:
              Confidence Intervals SVM Model Neural Network Model Random Forests Model Ada Boost Model
```

0.032026

0.160128

0.800641

0.096077

0.480384

2.401922

0.141478

0.141478

0.102482

0.512410

2.562050

Chebyshev Bounds for all 4 models with all confidence levels

75

95

99

0.057646

0.288231

1.441153

0

1

2

2

```
In [184]: confidence intervals = [75, 95, 99]
           svm B = []
           nn B = []
           rf B = []
           ada B = []
           for ci in confidence_intervals:
               svm B.append(chebyshev(test length, ci))
               nn B.append(chebyshev(test length, ci))
               rf B.append(chebyshev(test length, ci))
               ada B.append(chebyshev(test length, ci))
In [185]: chebyshev_bounds_list = pd.DataFrame(
               {'Confidence Intervals': confidence_intervals,
                 'SVM Model': svm_B,
                'Neural Network Model': nn B,
                'Random Forests Model': rf B,
                'Ada Boost Model': ada B
           chebyshev_bounds_list
Out[185]:
              Confidence Intervals SVM Model Neural Network Model Random Forests Model Ada Boost Model
            0
                                 0.028296
                                                  0.028296
                                                                     0.028296
                                                                                   0.028296
            1
                                0.063271
                                                  0.063271
                                                                     0.063271
                                                                                   0.063271
                            95
```

Hoeffding Bounds for all 4 models with all confidence levels

99

0.141478

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0.141478

Out[192]:

	Confidence Intervals	SVM Model	Neural Network Model	Random Forests Model	Ada Boost Model
0	75	0.028852	0.028852	0.028852	0.028852
1	95	0.038428	0.038428	0.038428	0.038428
2	99	0.046055	0.046055	0.046055	0.046055

2a) Answer: With respect to Markov bounds, Random forests and Adaboost changed dramatically followed by Polynomial SVM. The change in Neural Networks bounds is the least with respect to the change in confidence intervals. These(Markov) bounds are mostly effected by the test_error obtained in the models. Since neural networks has least error, the effect is also low. Since random forests and Adaboost has higher test errors, the bounds are largely effected by the confidence intervals. Chebyshev and Hoeffding are dependent on the number of test samples, the bounds are same for all the models since all the models have same number of test samples.

- 2b) I would choose neural networks since the training error and test error are very low i.e., this model has low bias and low variance which is optimal. And also, the concentration bounds are also very low. Since we always want to have low variance in the error and the neural network model has lowest error with low concentration bounds wrt confidence intervals, I would choose this model. i.e., We want to have lowest test error(+/-)councentration bounds, neural networks is optimal here.
- 2c) Yes, the markov bounds given an idea about the range of whole test error that we can have. Even though the bounds cannot give perfect answers, they give any idea about the model upper and lower limits to the test error. Also, chebyshev and hoeffding bounds given an idea about the whole test error with respect to the test data samples. They explain that as test data increases, the overall test error (Eout) value can decrease since the bounds decrease too.

```
In [ ]:
```