## **Team: Mohit Bansal**

```
In [224]: import pandas as pd
  import matplotlib.pyplot as plt
  import seaborn as sns
  import numpy as np
  import math
  import string
```

# **Data Cleaning and splitting method**

```
In [225]: from sklearn.model selection import train test split
          # takes in raw data as given performs feature engineering to
          # add useful columns and clean up existing columns. Does not
          # change the original data
          def clean_and_split(students, size):
              students cleaned = students.copy(deep=True)
              constants = [] # keeping track of constants added for validation data
              # splitting course id to get course and time
              course = []
              time = []
              for course_id in students_cleaned['course_id']:
                  info = course_id.split('/')
                  course.append(info[1])
                  time.append(info[2])
              students_cleaned['course'] = course
              students_cleaned['time'] = time
              # replacing NaN values with 'nan' strings for categorical data
              students cleaned['final cc cname DI'] = students['final cc cname DI'].fill
          na('nan')
              students_cleaned['LoE_DI'] = students_cleaned['LoE_DI'].fillna('nan')
              students cleaned['gender'] = students cleaned['gender'].fillna('nan')
              # one hot encoding for categorical variables
              students_cleaned = pd.get_dummies(students_cleaned, columns=['final_cc_cna
          me_DI', 'LoE_DI', 'gender', 'course', 'time'])
              features = list(students_cleaned.columns)
              features.remove('course id')
              features.remove('userid DI')
              if 'certified' in features:
                  features.remove('certified')
              features.remove('start time DI')
              features.remove('last_event_DI')
              # splitting between train and validation sets
              train_data, validation_data = train_test_split(students_cleaned, test_size
          =size)
              train data = train data.copy(deep=True)
              validation data = validation data.copy(deep=True)
              # converting grade strings to float and replacing NaN values with mean gra
          de
              train data['grade'] = pd.to numeric(train data['grade'], errors='coerce')
              validation data['grade'] = pd.to numeric(validation data['grade'], errors=
           'coerce')
              mean_grade = train_data['grade'].mean()
              mean_grade = round(mean_grade,2) # round off to 2 decimal places
              train data['grade'] = train data['grade'].fillna(mean grade)
              validation_data['grade'] = validation_data['grade'].fillna(mean_grade)
              # replacing NaN values with mean for other numerical features
              mean_yob = round(train_data['YoB'].mean())
              train_data['YoB'] = train_data['YoB'].fillna(mean_yob)
              validation data['YoB'] = validation data['YoB'].fillna(mean yob)
              mean_nevents = round(train_data['nevents'].mean())
```

```
train data['nevents'] = train data['nevents'].fillna(mean nevents)
   validation_data['nevents'] = validation_data['nevents'].fillna(mean_nevent
s)
   mean ndays = round(train data['ndays act'].mean())
   train data['ndays act'] = train data['ndays act'].fillna(mean ndays)
   validation_data['ndays_act'] = validation_data['ndays_act'].fillna(mean_nd
ays)
   mean_nplays = round(train_data['nplay_video'].mean())
   train_data['nplay_video'] = train_data['nplay_video'].fillna(mean_nplays)
   validation data['nplay video'] = validation data['nplay video'].fillna(mea
n nplays)
   mean_chapters = round(train_data['nchapters'].mean())
   train data['nchapters'] = train data['nchapters'].fillna(mean chapters)
   validation_data['nchapters'] = validation_data['nchapters'].fillna(mean_ch
apters)
   mean forums = round(train data['nforum posts'].mean())
   train data['nforum posts'] = train data['nforum posts'].fillna(mean forums
)
   validation data['nforum posts'] = validation data['nforum posts'].fillna(m
ean_forums)
   # these are used to apply same tranformation to test features
   constants = []
   constants.append(mean_grade)
   constants.append(mean_yob)
   constants.append(mean_nevents)
   constants.append(mean_ndays)
   constants.append(mean_nplays)
   constants.append(mean chapters)
   constants.append(mean_forums)
   return constants, features, train data, validation data
```

```
In [226]: def clean_test(constants, students):
              students_cleaned = students.copy(deep=True)
              # splitting course_id to get course and time
              course = []
              time = []
              for course_id in students_cleaned['course_id']:
                  info = course_id.split('/')
                  course.append(info[1])
                  time.append(info[2])
              students_cleaned['course'] = course
              students_cleaned['time'] = time
              # replacing NaN values with 'nan' strings for categorical data
              students_cleaned['final_cc_cname_DI'] = students['final_cc_cname_DI'].fill
          na('nan')
              students_cleaned['LoE_DI'] = students_cleaned['LoE_DI'].fillna('nan')
              students_cleaned['gender'] = students_cleaned['gender'].fillna('nan')
              # one hot encoding for categorical variables
              students_cleaned = pd.get_dummies(students_cleaned, columns=['final_cc_cna
          me_DI', 'LoE_DI', 'gender', 'course', 'time'])
              # converting grade strings to float and replacing NaN values with mean gra
          de
              students_cleaned['grade'] = pd.to_numeric(students_cleaned['grade'], error
          s='coerce')
              mean grade = constants[0]
              students_cleaned['grade'] = students_cleaned['grade'].fillna(mean_grade)
              # replacing NaN values with mean for other numerical features
              mean_yob = constants[1]
              students_cleaned['YoB'] = students_cleaned['YoB'].fillna(mean_yob)
              mean nevents = constants[2]
              students_cleaned['nevents'] = students_cleaned['nevents'].fillna(mean_neve
          nts)
              mean_ndays = constants[3]
              students_cleaned['ndays_act'] = students_cleaned['ndays_act'].fillna(mean_
              mean_nplays = constants[4]
              students_cleaned['nplay_video'] = students_cleaned['nplay_video'].fillna(m
          ean_nplays)
              mean_chapters = constants[5]
              students_cleaned['nchapters'] = students_cleaned['nchapters'].fillna(mean_
          chapters)
              mean forums = constants[6]
              students_cleaned['nforum_posts'] = students_cleaned['nforum_posts'].fillna
          (mean_forums)
              return students cleaned
```

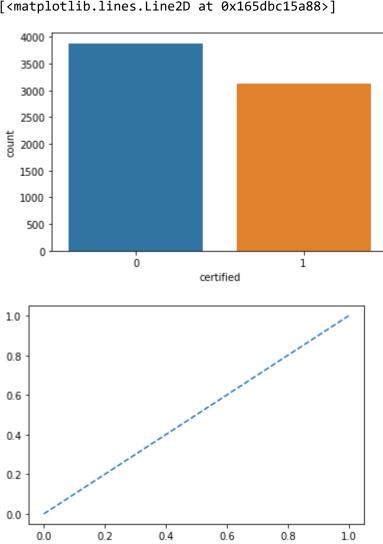
## Training the models

```
In [227]: students_data = pd.read_csv('edx_su2021_train.csv')
          constants, features, train_data, validation_data = clean_and_split(students_da
          ta, 0.2)
In [228]: validation_data['certified']
Out[228]: 6530
          5081
                  1
          7046
                  0
          5910
                  0
          6141
                  1
          2129
          451
                  0
          2397
                  1
          3390
                  0
          7102
          Name: certified, Length: 1752, dtype: int64
```

# **Majority Classifier**

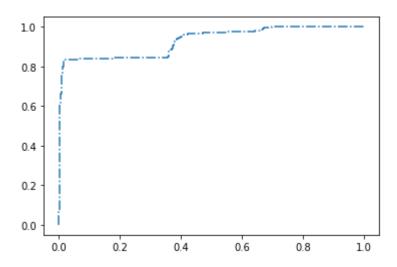
```
In [229]:
          from sklearn.metrics import roc_curve
          import matplotlib.pyplot as plt
          plt.figure()
          sns.countplot(x='certified', data = train data)
          # it is clear that 0 is the majority class
          majority_probs = np.zeros(len(validation_data))
          majority_fpr, majority_tpr, _ = roc_curve(validation_data['certified'], majori
          ty_probs)
          plt.figure()
          plt.plot(majority_fpr, majority_tpr, linestyle='--', label='Majority classifie
          r')
```

Out[229]: [<matplotlib.lines.Line2D at 0x165dbc15a88>]



## **Logistic Regression with L2 penalty**

#### Out[230]: [<matplotlib.lines.Line2D at 0x165dbc1de08>]

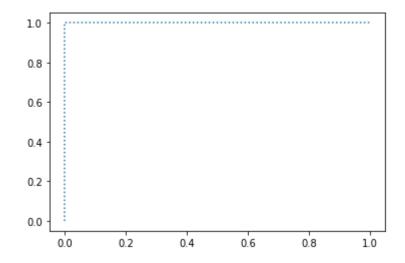


### **Decision Tree**

```
In [231]: from sklearn.tree import DecisionTreeClassifier
    from sklearn.model_selection import GridSearchCV
    hyperparameters = {
        'min_samples_leaf': [1, 10, 50, 100, 200, 300],
        'max_depth': [1, 5, 10, 15, 20]
    }
    estimator = DecisionTreeClassifier()
    search = GridSearchCV(estimator, hyperparameters, cv=6, return_train_score=Tru
    e)
    search.fit(train_data[features], train_data['certified'])
    print(search.best_params_)
```

{'max depth': 10, 'min samples leaf': 1}

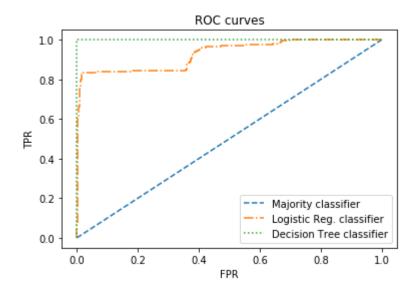
#### Out[232]: [<matplotlib.lines.Line2D at 0x165d8d08288>]



# **ROC** curves of all 3 - Comparing the models

```
In [233]: fig, ax = plt.subplots()
    ax.set_title('A single plot')
    ax.plot(majority_fpr, majority_tpr, linestyle='--', label='Majority classifie
    r')
    ax.plot(logistic_fpr, logistic_tpr, linestyle='-.', label='Logistic Reg. class
    ifier')
    ax.plot(tree_fpr, tree_tpr, linestyle=':', label='Decision Tree classifier')
    ax.legend()
    ax.set_xlabel("FPR")
    ax.set_ylabel("TPR")
    ax.set_title("ROC curves")
```

Out[233]: Text(0.5, 1.0, 'ROC curves')



### **Brief Inference:**

We have plotted the ROC curve which plots the True positive rate against the False positive rate for different threshold values. TPR is an indicator of how accurately the model predicts the positive class. The FPR on the other hand indicates how many times we wrongly predict positive. Based on the nature of these rates, we naturally want the FPR to be low and TPR to be high. According to this criterion, our decision tree classifier has the best model for all the thresholds. Another thing to consider is the Area under the curve. Higher area under an ROC curve generally implies a better model. This holds true since, the decision tree classifier has the highest area under the curve.

Through the ROC curves above we can see that we get the Decision Tree Classifier as the best model. Now, we use this model to predict the test labels

### **Test predictions**

```
In [234]: # Uploading test data
students_test = pd.read_csv('edx_su2021_test.csv')
students_test = clean_test(constants, students_test)
```

```
In [235]: predictions = tree_model.predict(students_test[features])
    to_save = students_test[['userid_DI']].copy()
    to_save.loc[:, 'certified'] = predictions
    to_save.to_csv('submission.csv', index=False)
```

### **Answers to Questions**

Discuss what features you used and whether you did some transformations on them. What features seemed important for your final model?

I used the course\_id feature to extract two columns for course code and time of year. I used the registered, viewed and explored features without any modification. I replaced NaN values with 'nan' strings for categorical data in LoE, final\_cc\_cname\_DI, and gender. I used one hot encoding for 'final\_cc\_cname\_DI', 'LoE\_DI', 'gender', 'course',and 'time' variables. For pother numerical features (YoB, grade, nevent, ndays\_act, nplay\_video, nchapters, nforum\_posts) I replaced the NaN values with mean of those columns in the training data.

The features of course and time of year seemed important for the final model because adding them reduced my validation error.

Explain what did you do to improve your initial predictions. Did it help?

To improve my initial predictions, I removed the features - start time and last event. I also added the course code and time of year features extracted from the course\_id. This seemed to greatly help my predictions. I also used GridSearchCV to perform hyperparameter tuning for 'min\_samples\_leaf' and 'max\_depth' hyperparameters of the decision tree classifier. This helped me in deciding the best values for these hyperparameters.

Consider and discuss the ethical implications of using the model you trained.

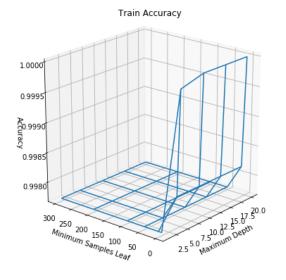
I can think of two ethical implications of using my model. First is the use of gender as feature which may lead to gender discrimination if used incorrectly. The second is the ethical implication of racial discrimination due to the presence of the 'final\_cc\_cname' feature.

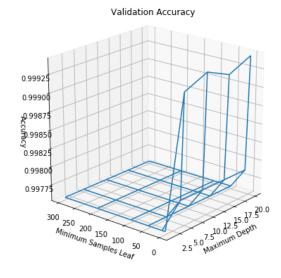
Imagine you were hired to work as a data scientist for an online education platform (exciting, right?). They want to use your model to predict which students they should tailor their course material towards so they can maximize their profits. Their idea is to use the model to help predict which student groups (i.e. from specific countries or certain educational backgrounds) would make them the most money by getting the most participants to complete their paid certificate program. Are there any ethical considerations we should think about before acting on that plan?

Yes, we should consider the ethical considerations of end goal. If our end goal is 'making money', we are bound to discriminate between countries/ continents and even gender if it suits our goal. This is because gender and continents are used as features in the model. This can further widen any cultural or academic disparity between two socio-economic groups by impacting the access to education for those groups.

## **Further Analysis (just for fun)**

```
In [236]:
          from mpl toolkits.mplot3d import Axes3D
          from sklearn import tree
          def plot scores(ax, title, search, hyperparameters, score key):
              # Get results from GridSearch and turn scores into matrix
              cv_results = search.cv_results_
              scores = cv_results[score_key]
              scores = scores.reshape((len(hyperparameters['max depth']), len(hyperparam
          eters['min samples leaf'])))
              max depths = cv results['param max depth'].reshape(scores.shape).data.asty
          pe(int)
              min_samples_leafs = cv_results['param_min_samples_leaf'].reshape(scores.sh
          ape).data.astype(int)
              # Plot result
              ax.plot wireframe(max depths, min samples leafs, scores)
              ax.view init(20, 220)
              ax.set_xlabel('Maximum Depth')
              ax.set ylabel('Minimum Samples Leaf')
              ax.set zlabel('Accuracy')
              ax.set_title(title)
          fig = plt.figure(figsize=(15,7))
          ax1 = fig.add_subplot(121, projection='3d')
          ax2 = fig.add subplot(122, projection='3d')
          plot_scores(ax1, 'Train Accuracy', search, hyperparameters, 'mean_train_score'
          plot scores(ax2, 'Validation Accuracy', search, hyperparameters, 'mean test sc
          ore')
```





```
In [237]: import scipy.stats
          class RandomForest416:
              This class implements the common sklearn model interface (has a fit and pr
          edict function).
              A random forest is a collection of decision trees that are trained on rand
          om subsets of the
              dataset. When predicting the value for an example, takes a majority vote f
          rom the trees.
              def __init__(self, num_trees, max_depth=None):
                  Constructs a RandomForest416 that uses the given numbner of trees, eac
          h with a
                  max depth of max_depth.
                  self. trees = [
                       DecisionTreeClassifier(max depth=max depth, random state=1)
                       for i in range(num_trees)
                   1
              def fit(self, X, y):
                   Takes an input dataset X and a series of targets y and trains the Rand
          omForest416.
                  Each tree will be trained on a random sample of the data that samples
           the examples
                  uniformly at random (with replacement). Each random dataset will have
           the same number
                  of examples as the original dataset, but some examples may be missing
           or appear more
                   than once due to the random sampling with replacement.
                  # TODO implement this method!
                  num examples = len(X)
                  for i,model in enumerate(self._trees):
                       rand_indices = np.random.randint(num_examples, size=num_examples)
                       random X = X.iloc[rand indices]
                       random_y = y.iloc[rand_indices]
                       self._trees[i] = model.fit(random_X, random_y)
              def predict(self, X):
                   Takes an input dataset X and returns the predictions for each example
           in X.
                   .....
                  # Builds up a 2d array with n rows and T columns
                  # where n is the number of points to classify and T is the number of t
          rees
                  predictions = np.zeros((len(X), len(self._trees)))
                  for i, tree in enumerate(self._trees):
```

```
# Make predictions using the current tree
preds = tree.predict(X)

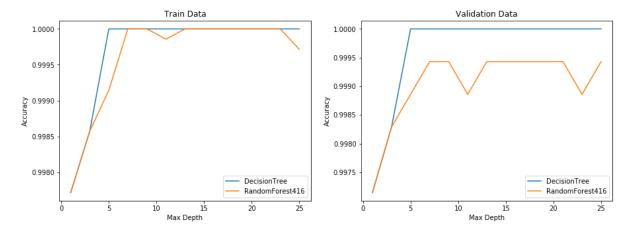
# Store those predictions in ith column of the 2d array
predictions[:, i] = preds

# For each row of predictions, find the most frequent label (axis=1 me
ans across columns)
    return scipy.stats.mode(predictions, axis=1)[0]

In [238]: rf = RandomForest416(2, max_depth=10)
    rf.fit(train_data[features], train_data['certified'])
    rf.predict(validation_data[features])
```

```
In [239]: from sklearn.metrics import accuracy score
          depths = list(range(1, 26, 2))
          dt accuracies = []
          rf accuracies = []
          target = 'certified'
          for i in depths:
              print(f'Depth {i}')
              # Train and evaluate a Decision Tree Classifier with given max depth
              tree = DecisionTreeClassifier(max_depth=i, min_samples_leaf=1)
              tree.fit(train data[features], train data[target])
              dt accuracies.append((
                   accuracy_score(tree.predict(train_data[features]), train_data[target
          ]),
                  accuracy score(tree.predict(validation data[features]), validation dat
          a[target])
              ))
              # Train and evaluate our RandomForest classifier with given max_depth
              rf = RandomForest416(15, max depth=i)
              rf.fit(train_data[features], train_data[target])
              rf_accuracies.append((
                   accuracy score(rf.predict(train data[features]), train data[target]),
                   accuracy_score(rf.predict(validation_data[features]), validation_data[
          target])
              ))
          # Then plot the scores
          fig, axs = plt.subplots(1, 2, figsize=(15, 5))
          # Plot training accuracies
          axs[0].plot(depths, [acc[0] for acc in dt_accuracies], label='DecisionTree')
          axs[0].plot(depths, [acc[0] for acc in rf accuracies], label='RandomForest416'
          )
          # Plot validation accuracies
          axs[1].plot(depths, [acc[1] for acc in dt_accuracies], label='DecisionTree')
          axs[1].plot(depths, [acc[1] for acc in rf_accuracies], label='RandomForest416'
          )
          # Customize plots
          axs[0].set title('Train Data')
          axs[1].set_title('Validation Data')
          for ax in axs:
              ax.legend()
              ax.set_xlabel('Max Depth')
              ax.set_ylabel('Accuracy')
```

Depth 1
Depth 5
Depth 7
Depth 9
Depth 11
Depth 13
Depth 15
Depth 17
Depth 17
Depth 19
Depth 21
Depth 23
Depth 23
Depth 25



#### In [240]:

['registered', 'viewed', 'explored', 'YoB', 'grade', 'nevents', 'ndays\_act', 'nplay\_video', 'nchapters', 'nforum\_posts', 'final\_cc\_cname\_DI\_Australia', inal cc cname DI Bangladesh', 'final cc cname DI Brazil', 'final cc cname DI Canada', 'final\_cc\_cname\_DI\_China', 'final\_cc\_cname\_DI\_Colombia', 'final\_cc\_c name\_DI\_Egypt', 'final\_cc\_cname\_DI\_France', 'final\_cc\_cname\_DI\_Germany', 'fin al cc cname DI Greece', 'final cc cname DI India', 'final cc cname DI Indones ia', 'final\_cc\_cname\_DI\_Japan', 'final\_cc\_cname\_DI\_Mexico', 'final\_cc\_cname\_D I\_Morocco', 'final\_cc\_cname\_DI\_Nigeria', 'final\_cc\_cname\_DI\_Other Africa', 'f inal\_cc\_cname\_DI\_Other East Asia', 'final\_cc\_cname\_DI\_Other Europe', 'final\_c c\_cname\_DI\_Other Middle East/Central Asia', 'final\_cc\_cname\_DI\_Other North & Central Amer., Caribbean', 'final\_cc\_cname\_DI\_Other Oceania', 'final\_cc\_cname \_DI\_Other South America', 'final\_cc\_cname\_DI\_Other South Asia', 'final\_cc\_cna me\_DI\_Pakistan', 'final\_cc\_cname\_DI\_Philippines', 'final\_cc\_cname\_DI\_Poland', 'final\_cc\_cname\_DI\_Portugal', 'final\_cc\_cname\_DI\_Russian Federation', 'final\_ cc\_cname\_DI\_Spain', 'final\_cc\_cname\_DI\_Ukraine', 'final\_cc\_cname\_DI\_United Ki ngdom', 'final\_cc\_cname\_DI\_United States', 'final\_cc\_cname\_DI\_Unknown/Other', "LoE DI Bachelor's", 'LoE DI Doctorate', 'LoE DI Less than Secondary', "LoE D I\_Master's", 'LoE\_DI\_Secondary', 'LoE\_DI\_nan', 'gender\_f', 'gender\_m', 'gende r\_nan', 'course\_CB22x', 'course\_CS50x', 'course\_ER22x', 'course\_PH207x', 'cou rse\_PH278x', 'time\_2012', 'time\_2012\_Fall', 'time\_2013\_Spring']