Assignment 6: Apply NB

- 1. Minimum data points need to be considered for people having 4GB RAM is 50k and for 8GB RAM is 100k
- 2. When you are using ramdomsearchev or gridsearchev you need not split the data into X_train,X_cv,X_test. As the above methods use kfold. The model will learn better if train data is more so splitting to X_train,X_test will suffice.
- 3. If you are writing for loops to tune your model then you need split the data into X_train,X_cv,X_test.
- 4. While splitting the data explore stratify parameter.
- 5. Apply Multinomial NB on these feature sets
 - Features that need to be considered essav

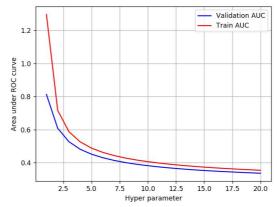
while encoding essay, try to experiment with the max_features and n_grams parameter of vectorizers and see if it increases AUC score.

categorical features

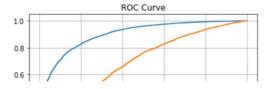
- teacher_prefix
- project_grade_category
- school_state
- clean_categories
- clean subcategories

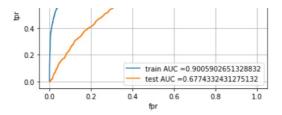
numerical features

- price
- teacher_number_of_previously_posted_projects while encoding the numerical features check this and this
- Set 1: categorical, numerical features + preprocessed_eassay (BOW)
- Set 2: categorical, numerical features + preprocessed_eassay (TFIDF)
- 6. The hyper paramter tuning(find best alpha:smoothing parameter)
 - Consider alpha values in range: 10^-5 to 10^2 like [0.00001,0.0005, 0.0001,0.005,0.001,0.05,0.01,0.1,0.5,1,5,10,50,100]
 - Explore class_prior = [0.5, 0.5] parameter which can be present in MultinomialNB function(go through this) then check how results might change.
 - Find the best hyper parameter which will give the maximum AUC value
 - For hyper parameter tuning using k-fold cross validation(use GridsearchCV or RandomsearchCV)/simple cross validation data (write for loop to iterate over hyper parameter values)
 - You need to plot the performance of model both on train data and cross validation data for each hyper parameter, like shown in the figure



- -while plotting take log(alpha) on your X-axis so that it will be more readable
- Once after you found the best hyper parameter, you need to train your model with it, and find the AUC on test data and plot the ROC curve on both train and test.





 Along with plotting ROC curve, you need to print the <u>confusion matrix</u> with predicted and original labels of test data points

	Predicted: NO	Predicted: YES	
Actual: NO	TN = ??	FP = ??	
Actual: YES	FN = ??	TP = ??	

-plot the confusion matrix in heatmaps, while plotting the confusion matrix go through the link

7. find the top 20 features from either from feature Set 1 or feature Set 2 using values of `feature_log_prob_` parameter of `MultinomialNB` (https://scikit-

learn.org/stable/modules/generated/sklearn.naive_bayes.MultinomialNB.html) and print **BOTH** positive as well as negative corresponding feature names.

- go through the link
- 8. You need to summarize the results at the end of the notebook, summarize it in the table format

Vectorizer	Model	+ Hyper parameter	AUC
BOW	Brute	7	0.78
TFIDF	Brute	12	0.79
W2V	Brute	10	0.78
TFIDFW2V	Brute	6	0.78

In [186]:

2. Naive Bayes

1.1 Loading Data

```
In [187]:
```

```
from google.colab import drive
drive.mount('/content/drive')
```

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount ("/content/drive", force remount=True).

In [188]:

```
#make sure you are loading atleast 50k datapoints
#you can work with features of preprocessed_data.csv for the assignment.
# If you want to add more features, you can add. (This is purely optional, not mandatory)
import pandas
data = pandas.read_csv('/content/drive/MyDrive/6_Donors_choose_NB/preprocessed_data.csv',
nrows = 100000)
# data = pd.read_csv('preprocessed_data.csv', nrows=50000) # you can take less number of
rows like this
```

```
data.head(3)
Out[189]:
  school_state teacher_prefix project_grade_category teacher_number_of_previously_posted_projects project_is_approved
0
                                                                                 53
                                                                                                  1
          ca
                      mrs
                                 grades_prek_2
1
           ut
                      ms
                                    grades_3_5
2
          ca
                      mrs
                                 grades_prek_2
                                                                                 10
                                                                                                   1 lit
In [190]:
data.shape
Out[190]:
(100000, 9)
In [191]:
data.isnull().sum()
Out[191]:
                                                      0
school state
teacher prefix
                                                      0
project_grade_category
                                                      0
{\tt teacher\_number\_of\_previously\_posted\_projects}
                                                      0
project_is_approved
                                                      0
                                                      0
clean_categories
clean_subcategories
                                                      0
                                                      0
essay
                                                      0
price
dtype: int64
In [192]:
data.dtypes
Out[192]:
school_state
                                                       object
teacher_prefix
                                                       object
project grade category
                                                       object
teacher number of previously posted projects
                                                        int64
project is approved
                                                        int64
clean categories
                                                       object
clean_subcategories
                                                       object
                                                       object
essay
price
                                                      float64
dtype: object
In [192]:
```

In [189]:

1.2 Splitting data into Train and cross validation(or test): Stratified Sampling

In [193]:

```
# write your code in following steps for task 1
# 1. Split your data.
# 2. Perform Bag of Words Vectorization of text data.
# 3. Perform tfidf vectorization of text data.
# 4. perform one-hot encoding of categorical features.
# 5. perform normalization of numerical features
# 6. For set 1 stack up all the features using hstack()
 7. For set 2 stack up all the features using hstack()
# 8. Perform hyperparameter tuning and represent the training and cross-validation AUC sc
ores for different 'alpha' values, using a 2D line plot.
# 9. Find the best hyperparameter 'alpha' and fit the model. Plot ROC-AUC curve(by obtain
ing the probabilities using 'predict proba' method)
# 10. Plot confusion matrix based on the best threshold value
# 11. Either for the model in set 1 or in set 2, print the top 20 features (you have to pr
int the names, not the indexes) associated with the positive and negative classes each.
# 12. Summarize your observations and compare both the models(ie., from set 1 and set 2)
in terms of optimal hyperparameter value, train AUC and test AUC scores.
# 13. You can use Prettytable or any other tabular format for comparison.
# please write all the code with proper documentation, and proper titles for each subsect
ion
# go through documentations and blogs before you start coding
# first figure out what to do, and then think about how to do.
# reading and understanding error messages will be very much helpfull in debugging your c
ode
# when you plot any graph make sure you use
    # a. Title, that describes your plot, this will be very helpful to the reader
    # b. Legends if needed
   # c. X-axis label
    # d. Y-axis label
```

In [194]:

```
%matplotlib inline
import warnings
warnings.filterwarnings("ignore")
import pandas as pd
import numpy as np
import nltk
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.feature_extraction.text import TfidfVectorizer
from sklearn.feature extraction.text import CountVectorizer
from sklearn.metrics import confusion matrix
from sklearn import metrics
from sklearn.metrics import roc curve, auc
import re
# Tutorial about Python regular expressions: https://pymotw.com/2/re/
import pickle
from tqdm import tqdm
import os
from collections import Counter
```

```
In [195]:
```

```
features =data.copy().drop(['project_is_approved'],axis=1)
y=data['project_is_approved']
```

```
data.groupby('project_is_approved').count()
```

Out[196]:

school_state teacher_prefix project_grade_category teacher_number_of_previously_posted_projects clear

project_is_approved

	0	15183	15183	15183	15183
	1	84817	84817	84817	84817
4)

In [197]:

```
from sklearn.model_selection import train_test_split

x_train,x_test, y_train_1, y_test = train_test_split(features, y, stratify=y, test_size=
0.25)

x_train, x_cv, y_train, y_cv = train_test_split(x_train, y_train_1, test_size=0.33, stra
tify=y_train_1)
```

In [198]:

```
# Split the dataset
# 1) If you want to apply simple cross-validation, split the dataset into 3 parts (ie., t
rain, CV and test sets)
# 2) If you want to apply K-fold CV (or) GridSearch Cross Validation (or) Randomized Sear
ch Cross Validation, just split the dataset into 2 parts (ie., train and test sets)
```

In [198]:

1.3 Make Data Model Ready: encoding essay, and project_title

```
In [199]:
```

```
from collections import Counter
my_counter = Counter()
for word in x_train['essay'].values:
    my_counter.update(word.split())

cat_dict = dict(my_counter)

sorted_bow_dict = dict(sorted(cat_dict.items(), key=lambda kc: kc[1]))
```

In [200]:

```
# Apply Bag of Words (BOW) vectorization on 'Preprocessed_Essay'
# Apply Bag of Words (BOW) vectorization on 'Preprocessed_Title' (Optional)

from sklearn.feature_extraction.text import CountVectorizer
vectorizer_bow = CountVectorizer(ngram_range=(1,2), vocabulary=list(sorted_bow_dict.keys
()), lowercase=False, binary=True)
#vectorizer.fit(x_train['essay'].values)

x_train_essay_bow = vectorizer_bow.fit_transform(x_train['essay'].values) # fit has to ha
ppen on only train data
x_cv_essay_bow = vectorizer_bow.transform(x_cv['essay'].values)
x_test_essay_bow = vectorizer_bow.transform(x_test['essay'].values)

# Note- apply fit_transform only on training dataset, because if we apply same fit
# transformation on both datset it creates data leakage
# Note- if we apply fit and transform seperately it takes more time than applying fit_tra
nsform
```

```
cat.extend(vectorizer bow.get feature names())
len(cat)
Out[201]:
41733
In [202]:
print(x train.shape, y train.shape)
print(x test.shape, y test.shape)
print('-'*50)
print("After vectorizations")
print(x train essay bow.shape, y train.shape)
print(x test essay bow.shape, y test.shape)
(50250, 8) (50250,)
(25000, 8) (25000,)
After vectorizations
(50250, 41733) (50250,)
(25000, 41733) (25000,)
In [203]:
from collections import Counter
my counter = Counter()
for word in x train['essay'].values:
   my counter.update(word.split())
cat dict = dict(my counter)
sorted tf dict = dict(sorted(cat dict.items(), key=lambda kc: kc[1]))
In [204]:
# Apply TF-IDF vectorization on 'Preprocessed Essay'
# Apply TF-IDF vectorization on 'Preprocessed Title' (Optional)
from sklearn.feature extraction.text import TfidfVectorizer
vectorizer tfidf = TfidfVectorizer(stop words="english", ngram range=(1,2), vocabulary=1
ist(sorted tf dict.keys()), lowercase=False, binary=True)
x train essay tfidf = vectorizer tfidf.fit transform(x train['essay'].values) # fit has
to happen on only train data
x cv essay tfidf = vectorizer tfidf.transform(x cv['essay'].values) # fit has to happen
on only train data
x test essay tfidf = vectorizer tfidf.transform(x test['essay'].values)
1.4 Make Data Model Ready: encoding numerical, categorical features
In [205]:
data.head(2)
Out[205]:
  school_state teacher_prefix project_grade_category teacher_number_of_previously_posted_projects project_is_approved cl
0
                    mrs
                              grades_prek_2
                                                                          53
                                                                                          1
         ca
```

cat = []

```
4
```

In [205]:

```
In [206]:
```

```
# Apply One-Hot Encoding on the categorical features either using OneHotEncoder() (or) Co
untVectorizer(binary=True)
# Apply Normalization on the numerical features using Normalizer().
from collections import Counter
my counter = Counter()
for word in x train['school state'].values:
   my counter.update(word.split())
cat dict = dict(my_counter)
sorted_school_dict = dict(sorted(cat dict.items(), key=lambda kc: kc[1]))
vectorizer school = CountVectorizer(vocabulary=list(sorted school dict.keys()), lowercase
=False, binary=True)
x train school ohe = vectorizer school.fit transform(x train['school state'].values)
x cv school ohe = vectorizer school.transform(x cv['school state'].values)
x test school ohe = vectorizer school.transform(x test['school state'].values)
from collections import Counter
my counter = Counter()
for word in x train['teacher prefix'].values:
   my counter.update(word.split())
cat dict = dict(my counter)
sorted teacher dict = dict(sorted(cat dict.items(), key=lambda kc: kc[1]))
vectorizer teacher = CountVectorizer(vocabulary=list(sorted teacher dict.keys()), lowerca
se=False, binary=True)
x train teacher ohe = vectorizer teacher.fit transform(x train['teacher prefix'].values)
x cv teacher ohe = vectorizer teacher.transform(x cv['teacher prefix'].values)
x test teacher ohe = vectorizer teacher.transform(x test['teacher prefix'].values)
from collections import Counter
my counter = Counter()
for word in x_train['project_grade_category'].values:
   my_counter.update(word.split())
cat dict = dict(my counter)
sorted grade dict = dict(sorted(cat dict.items(), key=lambda kc: kc[1]))
vectorizer grade = CountVectorizer(vocabulary=list(sorted grade dict.keys()), lowercase=F
alse, binary=True)
x train grade ohe = vectorizer grade.fit transform(x train['project grade category'].valu
x cv grade ohe = vectorizer grade.transform(x cv['project grade category'].values)
x test grade ohe = vectorizer grade.transform(x test['project grade category'].values)
from collections import Counter
my counter = Counter()
for word in x_train['clean_categories'].values:
   my_counter.update(word.split())
cat_dict = dict(my_counter)
sorted clean cat dict = dict(sorted(cat dict.items(), key=lambda kc: kc[1]))
vectorizer_clean_cat = CountVectorizer(vocabulary=list(sorted_clean_cat_dict.keys()), low
ercase=False, binary=True)
x train clean category ohe = vectorizer clean cat.fit transform(x train['clean categories
'].values)
x cv clean category ohe = vectorizer clean cat.fit transform(x cv['clean categories'].val
x test clean category ohe = vectorizer clean cat.transform(x test['clean categories'].val
ues)
```

```
from collections import Counter
my_counter = Counter()
for word in x_train['clean_subcategories'].values:
    my_counter.update(word.split())
cat_dict = dict(my_counter)
sorted_clean_sub_cat_dict = dict(sorted(cat_dict.items(), key=lambda kc: kc[1]))

vectorizer_clean_sub_cat = CountVectorizer(vocabulary=list(sorted_clean_sub_cat_dict.keys
()), lowercase=False, binary=True)
x_train_clean_subcategory_ohe = vectorizer_clean_sub_cat.fit_transform(x_train['clean_subcategories'].values)
x_cv_clean_subcategory_ohe = vectorizer_clean_sub_cat.transform(x_cv['clean_subcategories'].values)
x_test_clean_subcategory_ohe = vectorizer_clean_sub_cat.transform(x_test['clean_subcategories'].values)
```

In [207]:

```
from sklearn.preprocessing import Normalizer
# normalizer.fit(X train['price'].values)
# this will rise an error Expected 2D array, got 1D array instead:
# array=[105.22 215.96 96.01 ... 368.98 80.53 709.67].
# Reshape your data either using
# array.reshape(-1, 1) if your data has a single feature
# array.reshape(1, -1) if it contains a single sample.
# -1 means you don't need to specify the dimension number or it will automatically input
remaining dimensions
normalizer_1 = Normalizer()
x train price norm = normalizer 1.fit transform(x train['price'].values.reshape(-1,1))
x cv price norm = normalizer 1.transform(x cv['price'].values.reshape(-1,1))
x test price norm = normalizer 1.transform(x test['price'].values.reshape(-1,1))
normalizer = Normalizer()
x train posts norm = normalizer.fit transform(x train['teacher number of previously poste
d projects'].values.reshape(-1,1))
x cv posts norm = normalizer.transform(x cv['teacher number of previously posted projects
'].values.reshape(-1,1))
x_test_posts_norm = normalizer.transform(x_test['teacher_number_of_previously_posted_proj
ects'].values.reshape(-1,1))
```

In [208]:

```
# merge two sparse matrices: https://stackoverflow.com/a/19710648/4084039
from scipy.sparse import hstack
# set-1, with BOW
x tr = hstack((x train essay bow, x train school ohe, x train teacher ohe, x train grade
ohe, x_train_clean_category_ohe, x_train_clean_subcategory_ohe, x_train_price_norm, x_tra
in posts norm)).tocsr()
x c = hstack((x cv essay bow, x cv school ohe, x cv teacher ohe, x cv grade ohe, x cv cle
an_category_ohe, x_cv_clean_subcategory_ohe, x_cv_price_norm, x_cv_posts_norm)).tocsr()
x te = hstack((x test essay bow, x test school ohe, x test teacher ohe, x test grade ohe
, x_test_clean_category_ohe, x_test_clean_subcategory_ohe, x_test_price_norm, x_test_post
s norm)).tocsr()
\#set - 2 , with tfidf
x tr tf = hstack((x train essay tfidf, x train school ohe, x train teacher ohe, x train g
rade ohe, x train clean category ohe, x train clean subcategory ohe, x train price norm,
x train posts norm)).tocsr()
x c tf = hstack((x cv essay tfidf, x cv school ohe, x cv teacher ohe, x cv grade ohe, x
cv clean category ohe, x cv clean subcategory ohe, x cv price norm, x cv posts norm)).toc
x te tf = hstack((x test essay tfidf, x test school ohe, x test teacher ohe, x test grad
e_ohe, x_test_clean_category_ohe, x_test_clean_subcategory_ohe, x_test_price_norm, x_test
posts norm)).tocsr()
print(x_tr.shape, y_train.shape)
print(x c.shape, y cv.shape)
```

```
print(x_te.shape, y_test.shape)

print('*'*100)

print(x_tr_tf.shape, y_train.shape)
print(x_c_tf.shape, y_cv.shape)
print(x_te_tf.shape, y_test.shape)

(50250, 41834) (50250,)
(24750, 41834) (24750,)
(25000, 41834) (25000,)

*************
(50250, 41834) (50250,)
(24750, 41834) (50250,)
(24750, 41834) (24750,)
(25000, 41834) (25000,)
```

1.5 Appling NB on different kind of featurization as mentioned in the instructions

Apply NB on different kind of featurization as mentioned in the instructions

For Every model that you work on make sure you do the step 2 and step 3 of instrucations

Set 1

```
In [209]:
# Perform Hyperparameter Tuning.
# Plot the training and the CV AUC scores, for different values of 'alpha', using a 2D li
ne plot
```

```
In [210]:
```

```
def batch_predict(clf, data):
    # roc_auc_score(y_true, y_score) the 2nd parameter should be probability estimates of
the positive class
    # not the predicted outputs

y_data_pred = []
    tr_loop = data.shape[0] - data.shape[0]%1000
    # consider you X_tr shape is 49041, then your tr_loop will be 49041 - 49041%1000 = 49

000

# in this for loop we will iterate unti the last 1000 multiplier
for i in range(0, tr_loop, 1000):
    y_data_pred.extend(clf.predict_proba(data[i:i+1000])[:,1])
# we will be predicting for the last data points
if data.shape[0]%1000 !=0:
    y_data_pred.extend(clf.predict_proba(data[tr_loop:])[:,1])

return y_data_pred
```

In [211]:

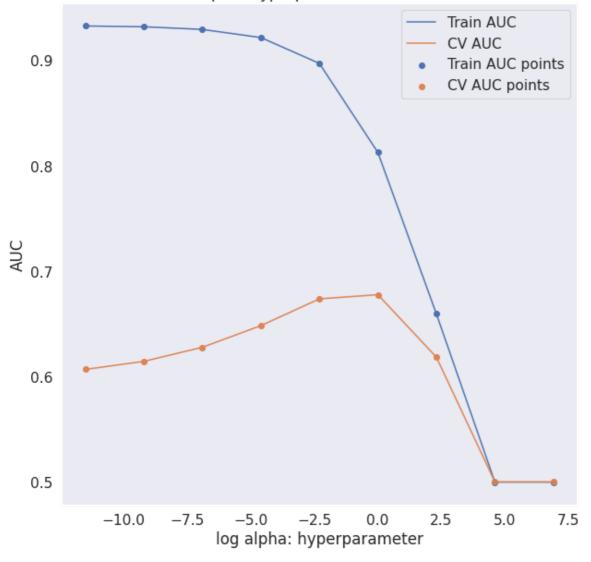
In [212]:

```
plt.figure(figsize=(10,10))
plt.plot(log_alphas, train_auc, label='Train AUC')
plt.plot(log_alphas, cv_auc, label='CV AUC')

plt.scatter(log_alphas, train_auc, label='Train AUC points')
plt.scatter(log_alphas, cv_auc, label='CV AUC points')

plt.legend()
plt.xlabel("log alpha: hyperparameter")
plt.ylabel("AUC")
plt.title("alpha: hyperparameter v/s AUC")
plt.grid()
plt.show()
```

alpha: hyperparameter v/s AUC



```
from sklearn.model selection import GridSearchCV
nb = MultinomialNB(class prior=[0.5,0.5])
parameters = { 'alpha': [0.00001, 0.0001, 0.001, 0.01, 0.1, 0.5, 0.8, 1, 10, 100,
clf = GridSearchCV(nb, parameters, cv= 10, scoring='roc auc', return train score=True, ver
bose=2)
clf.fit(x tr, y train)
train auc= clf.cv results ['mean train score']
train auc std= clf.cv results ['std train score']
cv auc = clf.cv results ['mean test score']
cv auc std= clf.cv results ['std test score']
Fitting 10 folds for each of 11 candidates, totalling 110 fits
[CV] END .....alpha=1e-05; total time=
                                              0.1s
[CV] END ......alpha=0.0001; total time=
                                              0.1s
[CV] END .....alpha=0.0001; total time=
                                              0.1s
[CV] END ......alpha=0.0001; total time=
                                              0.1s
[CV] END .....alpha=0.0001; total time=
                                              0.1s
[CV] END ......alpha=0.0001; total time=
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[CV] END ......alpha=0.0001; total time=
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[CV] END .....alpha=0.0001; total time=
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[CV] END .....alpha=0.01; total time=
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[CV] END .....alpha=0.1; total time=
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[CV] END .....alpha=0.5; total time=
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0.1s

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[CV] END .....alpha=0.5; total time=
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[CV] END .....alpha=0.5; total time=
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[CV] END ......alpha=1; total time=
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[CV] END .....alpha=10; total time=
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[CV] END ......alpha=10; total time=
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[CV] END ......alpha=10; total time=
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[CV] END .....alpha=10; total time=
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[CV] END .....alpha=100; total time=
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[CV] END ......alpha=100; total time=
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[CV] END .....alpha=100; total time=
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[CV] END .....alpha=100; total time=
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[CV] END .....alpha=100; total time=
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[CV] END ......alpha=100; total time=
                                       0.1s
[CV] END ......alpha=100; total time=
                                       0.1s
[CV] END .....alpha=100; total time=
                                       0.1s
[CV] END .....alpha=100; total time=
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[CV] END .....alpha=1000; total time=
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[CV] END ......alpha=1000; total time=
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[CV] END ......alpha=1000; total time=
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[CV] END .....alpha=1000; total time=
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[CV] END ......alpha=1000; total time=
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[CV] END ......alpha=1000; total time=
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[CV] END ......alpha=1000; total time=
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[CV] END .....alpha=1000; total time=
                                       0.1s
In [213]:
```

In [214]:

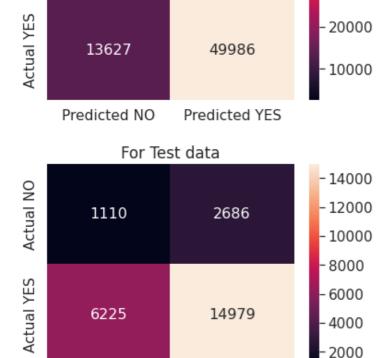
```
# Print params
print(clf.best_estimator_)
print(clf.score(x_tr, y_train))
print(clf.score(x_c, y_cv))
```

MultinomialNB(alpha=0.5, class_prior=[0.5, 0.5]) 0.8479187529562191 0.6818110819885058

In [215]: # Obtain the optimal value for 'alpha' and using the obtained optimal 'alpha' value, fit a multinomial naive bayes model, on the train data, # Note: If you have split the datase into 3 parts (ie., train, cv and test sets) in the b eginning, then the training data for this final model would be (train set + cv set) # Make class label and probability predictions on the train and test data. In [216]: from scipy.sparse import vstack # for vertical stacking $x = vstack((x_tr, x_c)).tocsr()$ print(x.shape) print(y train 1.shape) x tr.shape (75000, 41834) (75000,)Out[216]: (50250, 41834)In [217]: nb 1 = MultinomialNB(alpha = 0.1, class prior=[0.5, 0.5]) nb 1.fit(x,y train 1) Out[217]: MultinomialNB(alpha=0.1, class prior=[0.5, 0.5]) In [218]: # Pick the best threshold among the probability estimates, such that it has to yield maxi mum value for TPR*(1-FPR) # Plot the confusion matrices(each for train and test data) afer encoding the predicted c lass labels, on the basis of the best threshod probability estimate. In [219]: from sklearn.metrics import confusion matrix #function to get heatmap confusion matrix def get confusion matrix(clf, X te, y test): y pred = clf.predict(X te) df cm = pd.DataFrame(confusion matrix(y test, y pred), range(2), range(2)) df cm.columns = ['Predicted NO', 'Predicted YES'] df cm = df cm.rename({0: 'Actual NO', 1: 'Actual YES'}) sns.set(font scale=1.4) #for label size sns.heatmap(df cm, annot=True, annot kws={"size": 16}, fmt='g') In [220]: # Confusion Matrix

```
get confusion_matrix(nb_1,x,y_train_1)
plt.title('For Train data')
plt.show()
get confusion matrix(nb_1,x_te,y_test)
plt.title('For Test data')
plt.show()
```

For Train data - 40000 8759 2628 30000



Predicted YES

In [221]:

Predicted NO

AUC Score - 0.509629299321022

Plot the ROC-AUC curves using the probability predictions made on train and test data.

In [222]:

```
from sklearn.metrics import roc_auc_score

# predict probabilities
pred_prob_1 = nb_1.predict_proba(x_te)

# auc scores
auc_score = roc_auc_score(y_test, pred_prob_1[:,1])
print('AUC Score - ',auc_score)
```

In [223]:

```
# For best threshold
from sklearn.metrics import roc_curve

fpr, tpr, thresholds = roc_curve(y_test, pred_prob_1[:,1])

best_idx = np.argmax(tpr*(1- fpr))
best_threshold = thresholds[best_idx]

best_threshold
```

Out[223]:

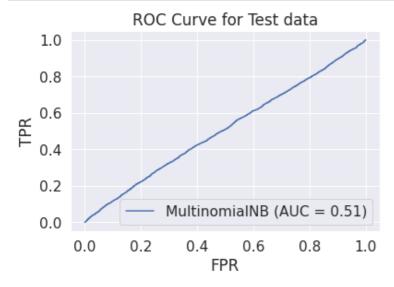
0.6872561813240533

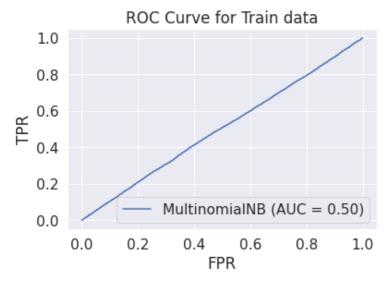
In [224]:

```
from sklearn import metrics

metrics.plot_roc_curve(nb_1, x_te, y_test)
plt.title('ROC Curve for Test data')
plt.xlabel('FPR')
plt.ylabel('TPR')
metrics.plot_roc_curve(nb_1, x_tr, y_train)
plt.title('ROC Curve for Train data')
plt.xlabel('FPR')
```

```
plt.ylabel('TPR')
plt.show()
```





Set 2

In [225]:

```
# Perform Hyperparameter Tuning.
# Plot the training and the CV AUC scores, for different values of 'alpha', using a 2D line plot
```

In [226]:

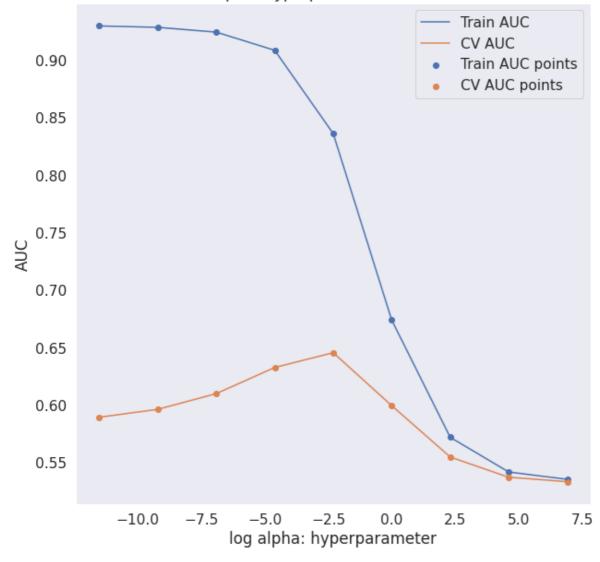
In [227]:

```
plt.figure(figsize=(10,10))
plt.plot(log_alphas, train_auc, label='Train AUC')
plt.plot(log_alphas, cv_auc, label='CV AUC')

plt.scatter(log_alphas, train_auc, label='Train AUC points')
plt.scatter(log_alphas, cv_auc, label='CV AUC points')

plt.legend()
plt.xlabel("log alpha: hyperparameter")
plt.ylabel("AUC")
plt.title("alpha: hyperparameter v/s AUC")
plt.grid()
plt.show()
```

alpha: hyperparameter v/s AUC



In [228]:

```
from sklearn.model_selection import GridSearchCV

nb = MultinomialNB(class_prior=[0.5,0.5])
```

```
parameters = { 'alpha': [0.00001, 0.0001, 0.001, 0.01, 0.1, 0.5, 0.8, 1, 10, 100,
clf = GridSearchCV(nb, parameters, cv= 10, scoring='roc auc', return train score=True, ver
bose=2)
clf.fit(x_tr_tf, y_train)
train auc= clf.cv results ['mean train score']
train auc std= clf.cv results ['std train score']
cv auc = clf.cv results ['mean test score']
cv auc std= clf.cv results ['std test score']
Fitting 10 folds for each of 11 candidates, totalling 110 fits
[CV] END .....alpha=1e-05; total time=
                                             0.1s
[CV] END .....alpha=0.0001; total time=
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[CV] END .....alpha=0.0001; total time=
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[CV] END ......alpha=0.0001; total time=
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[CV] END .....alpha=0.0001; total time=
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[CV] END .....alpha=0.0001; total time=
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[CV] END .....alpha=0.0001; total time=
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[CV] END ......alpha=0.0001; total time=
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[CV] END .....alpha=0.001; total time=
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[CV] END ......alpha=0.01; total time=
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[CV] END ......alpha=0.01; total time=
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[CV] END .....alpha=0.01; total time=
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[CV] END ......alpha=0.01; total time=
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[CV] END .....alpha=0.1; total time=
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[CV] END .....alpha=0.5; total time=
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```

```
[CV] END .....alpha=0.5; total time=
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[CV] END .....alpha=1000; total time=
                                        0.1s
[CV] END .....alpha=1000; total time=
```

0.1s

In [229]:

```
# Print params
print(clf.best estimator )
print(clf.score(x tr tf, y train))
print(clf.score(x_c_tf, y_cv))
```

MultinomialNB(alpha=0.1, class prior=[0.5, 0.5])

0.8360001749193261

0.645452607607835

In [230]:

```
# Obtain the optimal value for 'alpha' and using the obtained optimal 'alpha' value, fit
a multinomial naive bayes model, on the train data,
# Note: If you have split the datase into 3 parts (ie., train, cv and test sets) in the b
eginning, then the training data for this final model would be (train set + cv set)
# Make class label and probability predictions on the train and test data.
```

In [231]:

```
from scipy.sparse import vstack # for vertical stacking
x = vstack((x_tr_tf,x_c_tf)).tocsr()
print(x.shape)
print(y_train_1.shape)
```

(75000, 41834) (75000,)

In [232]:

```
nb_2 = MultinomialNB(alpha = 0.001, class_prior=[0.5,0.5])
nb_2.fit(x,y_train_1)
```

Out[232]:

MultinomialNB(alpha=0.001, class prior=[0.5, 0.5])

In [233]:

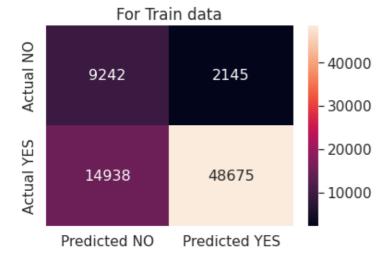
Pick the best threshold among the probability estimates, such that it has to yield maximum value for TPR*(1-FPR)

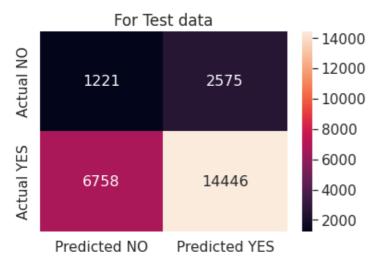
Plot the confusion matrices(each for train and test data) afer encoding the predicted c lass labels, on the basis of the best threshod probability estimate.

In [234]:

```
# Confusion Matrix
get_confusion_matrix(nb_2,x,y_train_1)
plt.title('For Train data')
plt.show()

get_confusion_matrix(nb_2,x_te_tf,y_test)
plt.title('For Test data')
plt.show()
```





```
In [235]:
from sklearn.metrics import roc_auc_score

# predict probabilities
pred_prob_2 = nb_2.predict_proba(x_te_tf)

# auc scores
auc_score = roc_auc_score(y_test, pred_prob_2[:,1])
```

print('AUC Score - ',auc_score)
AUC Score - 0.5057692730103014

In [236]:

```
# For best threshold
from sklearn.metrics import roc_curve

fpr, tpr, thresholds = roc_curve(y_test, pred_prob_2[:,1])

best_idx = np.argmax(tpr*(1- fpr))
best_threshold = thresholds[best_idx]

best_threshold
```

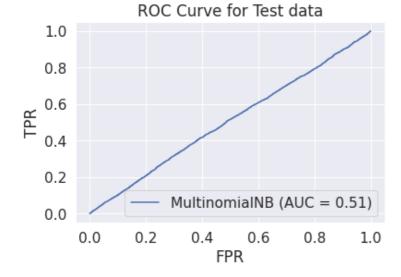
Out[236]:

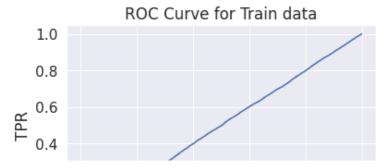
0.538054579331292

In [237]:

```
from sklearn import metrics

metrics.plot_roc_curve(nb_2, x_te_tf, y_test)
plt.title('ROC Curve for Test data')
plt.xlabel('FPR')
plt.ylabel('TPR')
metrics.plot_roc_curve(nb_2, x_tr_tf, y_train)
plt.title('ROC Curve for Train data')
plt.xlabel('FPR')
plt.ylabel('TPR')
plt.show()
```







```
Task -2
In [238]:
# Either from set 1 (or) set 2, print the names of the top 20 features associated with th
e positive and negative classes each. (You have to print the names of the features, but n
ot the indexes)
In [239]:
bow features = []
# for Categorical Features
bow features.extend(vectorizer bow.get feature names())
bow features.extend(vectorizer school.get feature names())
bow features.extend(vectorizer teacher.get feature names())
bow features.extend(vectorizer grade.get feature names())
bow features.extend(vectorizer clean cat.get feature names())
bow features.extend(vectorizer clean sub cat.get feature names())
# For Numerical features
bow features.append("price")
bow features.append("teacher number of previously posted projects")
print(len(bow features))
41834
In [240]:
positive=list(np.argsort((nb 1.feature log prob )[1]))
positive.reverse()
positive featuers=np.array(bow features)[np.array(positive[:10])]
positive featuers
Out[240]:
array(['price', 'students', 'nannan', 'school', 'my',
       'teacher_number_of_previously_posted_projects', 'learning',
       'classroom', 'not', 'learn'], dtype='<U44')
In [241]:
negetive=list(np.argsort((nb 1.feature log prob )[0]))
negetive.reverse()
negetive featuers=np.array(bow features)[np.array(negetive[:10])]
negetive featuers
Out[241]:
array(['price', 'students', 'nannan', 'school', 'my',
```

'teacher_number_of_previously_posted_projects', 'learning',

'classroom', 'not', 'the'], dtype='<U44')

3 Summary

v. varriiriar y

as mentioned in the step 5 of instructions

In [242]:

```
#Summarize your assignment work here in a few points, and also compare the final models (
from set 1 and set 2), in terms of optimal hyperparameter value 'alpha', training AUC and
test AUC scores.

# You can either use a pretty table or any other tabular structure.
```

Reference Link for Pretty table: https://pypi.org/project/prettytable/

In [244]:

```
from prettytable import PrettyTable
table = PrettyTable()

table.field_names = ["Vectorizer", "Model", "Hyper-parameter (Alpha)", "AUC Score", "Best
Threshold"]
table.add_row(["BOW", "Naive Bayes", 0.1, 0.50, 0.68])
table.add_row(["TFIDF", "Naive Bayes", 0.01, 0.50, 0.533])
print(table)
```

Vectorizer	+	Hyper-parameter (Alpha)	+	Best Threshold
BOW TFIDF	Naive Bayes Naive Bayes	0.1 0.01	0.5	0.68