Amazon Fine Food Reviews Analysis

Data Source: https://www.kaggle.com/snap/amazon-fine-food-reviews (https://www.kaggle.com/snap/amazon-fine-food-reviews (https://www.kaggle.com/snap/amazon-fine-food-reviews (https://www.kaggle.com/snap/amazon-fine-food-reviews (https://www.kaggle.com/snap/amazon-fine-food-reviews)

EDA: https://nycdatascience.com/blog/student-works/amazon-fine-foods-visualization/)

The Amazon Fine Food Reviews dataset consists of reviews of fine foods from Amazon.

Number of reviews: 568,454 Number of users: 256,059 Number of products: 74,258 Timespan: Oct 1999 - Oct 2012 Number of Attributes/Columns in data: 10

Attribute Information:

- 1. Id
- 2. Productld unique identifier for the product
- 3. UserId unqiue identifier for the user
- 4. ProfileName
- 5. HelpfulnessNumerator number of users who found the review helpful
- 6. HelpfulnessDenominator number of users who indicated whether they found the review helpful or not
- 7. Score rating between 1 and 5
- 8. Time timestamp for the review
- 9. Summary brief summary of the review
- 10. Text text of the review

Objective:

Given a review, determine whether the review is positive (Rating of 4 or 5) or negative (rating of 1 or 2).

[Q] How to determine if a review is positive or negative?

[Ans] We could use the Score/Rating. A rating of 4 or 5 could be cosnidered a positive review. A review of 1 or 2 could be considered negative. A review of 3 is nuetral and ignored. This is an approximate and proxy way of determining the polarity (positivity/negativity) of a review.

Loading the data

The dataset is available in two forms

- 1. .csv file
- 2. SQLite Database

In order to load the data, We have used the SQLITE dataset as it easier to query the data and visualise the data efficiently.

Here as we only want to get the global sentiment of the recommendations (positive or negative), we will purposefully ignore all Scores equal to 3. If the score id above 3, then the recommendation will be set to "positive". Otherwise, it will be set to "negative".

```
In [74]: %matplotlib inline
         import warnings
         warnings.filterwarnings("ignore")
         import sqlite3
         import pandas as pd
         import numpy as np
         import nltk
         import string
         import matplotlib.pyplot as plt
         import seaborn as sns
         from sklearn.feature extraction.text import TfidfTransformer
         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
         from nltk.stem.porter import PorterStemmer
         import re
         # Tutorial about Python regular expressions: https://pymotw.com/2/re/
         import string
         from nltk.corpus import stopwords
         from nltk.stem import PorterStemmer
         from nltk.stem.wordnet import WordNetLemmatizer
         from gensim.models import Word2Vec
         from gensim.models import KeyedVectors
         import pickle
         from tqdm import tqdm
         import os
         from sklearn.model selection import train test split
         from sklearn.metrics import roc auc score
         from sklearn.neighbors import KNeighborsClassifier
         from sklearn.model selection import GridSearchCV
         from sklearn.feature extraction.text import CountVectorizer
         from sklearn.metrics import accuracy_score
         from sklearn.cross validation import cross val score
         from collections import Counter
         from sklearn import cross validation
```

[1]. Reading Data

```
In [75]: # using SQLite Table to read data.
         con = sqlite3.connect('D:\\TGM\\ML\\AmazonFineFoodReviews\\database.sqlite')
         # filtering only positive and negative reviews i.e.
         # not taking into consideration those reviews with Score=3
         # SELECT * FROM Reviews WHERE Score != 3 LIMIT 500000, will give top 500000 da
         ta points
         # you can change the number to any other number based on your computing power
         # filtered_data = pd.read_sql_query(""" SELECT * FROM Reviews WHERE Score != 3
         LIMIT 500000""", con)
         # for tsne assignment you can take 5k data points
         filtered data = pd.read sql query(""" SELECT * FROM Reviews WHERE Score !=
          3""", con)
         # Give reviews with Score>3 a positive rating(1), and reviews with a score<3 a
         negative rating(0).
         def partition(x):
             if x < 3:
                 return 0
             return 1
         #changing reviews with score less than 3 to be positive and vice-versa
         actualScore = filtered data['Score']
         positiveNegative = actualScore.map(partition)
         filtered data['Score'] = positiveNegative
         print("Number of data points in our data", filtered data.shape)
         filtered data.head(3)
```

Number of data points in our data (525814, 10)

Out[75]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfulne
0	1	B001E4KFG0	A3SGXH7AUHU8GW	delmartian	1	1
1	2	B00813GRG4	A1D87F6ZCVE5NK	dli pa	0	0
2	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1	1

```
In [76]: display = pd.read_sql_query("""
    SELECT UserId, ProductId, ProfileName, Time, Score, Text, COUNT(*)
    FROM Reviews
    GROUP BY UserId
    HAVING COUNT(*)>1
    """, con)
```

In [77]: print(display.shape)
 display.head()

(80668, 7)

Out[77]:

	UserId	ProductId	ProfileName	Time	Score	Text	COL
0	#oc- R115TNMSPFT9I7	B007Y59HVM	Breyton	1331510400	2	Overall its just OK when considering the price	2
1	#oc- R11D9D7SHXIJB9	B005HG9ET0	Louis E. Emory "hoppy"	1342396800	5	My wife has recurring extreme muscle spasms, u	3
2	#oc- R11DNU2NBKQ23Z	B007Y59HVM	Kim Cieszykowski	1348531200	1	This coffee is horrible and unfortunately not	2
3	#oc- R11O5J5ZVQE25C	B005HG9ET0	Penguin Chick	1346889600	5	This will be the bottle that you grab from the	3
4	#oc- R12KPBODL2B5ZD	B007OSBE1U	Christopher P. Presta	1348617600	1	I didnt like this coffee. Instead of telling y	2

In [78]: display[display['UserId']=='AZY10LLTJ71NX']

Out[78]:

	UserId	ProductId	ProfileName	Time	Score	Text
80638	AZY10LLTJ71NX	B006P7E5ZI	undertheshrine "undertheshrine"	1334707200	5	I was recommended to try green tea extract to

```
In [79]: display['COUNT(*)'].sum()
Out[79]: 393063
```

Exploratory Data Analysis

[2] Data Cleaning: Deduplication

It is observed (as shown in the table below) that the reviews data had many duplicate entries. Hence it was necessary to remove duplicates in order to get unbiased results for the analysis of the data. Following is an example:

In [80]: display= pd.read_sql_query("""
 SELECT *
 FROM Reviews
 WHERE Score != 3 AND UserId="AR5J8UI46CURR"
 ORDER BY ProductID
 """, con)
 display.head()

Out[80]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpful
0	78445	B000HDL1RQ	AR5J8UI46CURR	Geetha Krishnan	2	2
1	138317	B000HDOPYC	AR5J8UI46CURR	Geetha Krishnan	2	2
2	138277	B000HDOPYM	AR5J8UI46CURR	Geetha Krishnan	2	2
3	73791	B000HDOPZG	AR5J8UI46CURR	Geetha Krishnan	2	2
4	155049	B000PAQ75C	AR5J8UI46CURR	Geetha Krishnan	2	2

As can be seen above the same user has multiple reviews of the with the same values for HelpfulnessNumerator, HelpfulnessDenominator, Score, Time, Summary and Text and on doing analysis it was found that

ProductId=B000HDOPZG was Loacker Quadratini Vanilla Wafer Cookies, 8.82-Ounce Packages (Pack of 8)

ProductId=B000HDL1RQ was Loacker Quadratini Lemon Wafer Cookies, 8.82-Ounce Packages (Pack of 8) and so on

It was inferred after analysis that reviews with same parameters other than Productld belonged to the same product just having different flavour or quantity. Hence in order to reduce redundancy it was decided to eliminate the rows having same parameters.

The method used for the same was that we first sort the data according to ProductId and then just keep the first similar product review and delette the others. for eg. in the above just the review for ProductId=B000HDL1RQ remains. This method ensures that there is only one representative for each product and deduplication without sorting would lead to possibility of different representatives still existing for the same product.

```
In [81]: #Sorting data according to ProductId in ascending order
    sorted_data=filtered_data.sort_values('ProductId', axis=0, ascending=True, inp
    lace=False, kind='quicksort', na_position='last')

In [82]: #Deduplication of entries
    final=sorted_data.drop_duplicates(subset={"UserId","ProfileName","Time","Text"
    }, keep='first', inplace=False)
    final.shape

Out[82]: (364173, 10)

In [83]: #Checking to see how much % of data still remains
    (final['Id'].size*1.0)/(filtered_data['Id'].size*1.0)*100

Out[83]: 69.25890143662969
```

Observation:- It was also seen that in two rows given below the value of HelpfulnessNumerator is greater than HelpfulnessDenominator which is not practically possible hence these two rows too are removed from calcualtions

```
In [84]: display= pd.read_sql_query("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND Id=44737 OR Id=64422
    ORDER BY ProductID
    """, con)
display.head()
```

Out[84]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfulr
0	64422	B000MIDROQ	A161DK06JJMCYF	J. E. Stephens "Jeanne"	3	1
1	44737	B001EQ55RW	A2V0I904FH7ABY	Ram	3	2

In [85]: final=final[final.HelpfulnessNumerator<=final.HelpfulnessDenominator]</pre>

#How many positive and negative reviews are present in our dataset?
final['Score'].value_counts()

(364171, 10)

Out[86]: 1 307061

0 57110

Name: Score, dtype: int64

[3]. Text Preprocessing.

Now that we have finished deduplication our data requires some preprocessing before we go on further with analysis and making the prediction model.

Hence in the Preprocessing phase we do the following in the order below:-

- 1. Begin by removing the html tags
- 2. Remove any punctuations or limited set of special characters like, or . or # etc.
- 3. Check if the word is made up of english letters and is not alpha-numeric
- 4. Check to see if the length of the word is greater than 2 (as it was researched that there is no adjective in 2-letters)
- 5. Convert the word to lowercase
- 6. Remove Stopwords
- 7. Finally Snowball Stemming the word (it was observed to be better than Porter Stemming)

After which we collect the words used to describe positive and negative reviews

```
In [87]: # printing some random reviews
    sent_0 = final['Text'].values[0]
    print(sent_0)
    print("="*50)

sent_1000 = final['Text'].values[1000]
    print(sent_1000)
    print("="*50)

sent_1500 = final['Text'].values[1500]
    print(sent_1500)
    print("="*50)

sent_4900 = final['Text'].values[4900]
    print(sent_4900)
    print("="*50)
```

this witty little book makes my son laugh at loud. i recite it in the car as we're driving along and he always can sing the refrain. he's learned about wh ales, India, drooping roses: i love all the new words this book introduces and the silliness of it all. this is a classic book i am willing to bet my son will STILL be able to recite from memory when he is in college

I was really looking forward to these pods based on the reviews. Starbucks i s good, but I prefer bolder taste... imagine my surprise when I ordered 2 bo xes - both were expired! One expired back in 2005 for gosh sakes. I admit th at Amazon agreed to credit me for cost plus part of shipping, but geez, 2 years expired!!! I'm hoping to find local San Diego area shoppe that carries po ds so that I can try something different than starbucks.

Great ingredients although, chicken should have been 1st rather than chicken broth, the only thing I do not think belongs in it is Canola oil. Canola or r apeseed is not someting a dog would ever find in nature and if it did find ra peseed in nature and eat it, it would poison them. Today's Food industries ha ve convinced the masses that Canola oil is a safe and even better oil than ol ive or virgin coconut, facts though say otherwise. Until the late 70's it was poisonous until they figured out a way to fix that. I still like it but it could be better.

Can't do sugar. Have tried scores of SF Syrups. NONE of them can touch the excellence of this product.

/>cbr />Thick, delicious. Perfect. 3 ingredictions: Water, Maltitol, Natural Maple Flavor. PERIOD. No chemicals. No garb age.

/>cbr />Have numerous friends & family members hooked on this stuff. My husband & son, who do NOT like "sugar free" prefer this over major label regular syrup.

/>cbr />I use this as my SWEETENER in baking: cheesecakes, white brownies, muffins, pumpkin pies, etc... Unbelievably delicious...

/>cbr />Can you tell I like it?:)

```
In [88]: # remove urls from text python: https://stackoverflow.com/a/40823105/4084039
    sent_0 = re.sub(r"http\S+", "", sent_0)
    sent_1000 = re.sub(r"http\S+", "", sent_1000)
    sent_150 = re.sub(r"http\S+", "", sent_1500)
    sent_4900 = re.sub(r"http\S+", "", sent_4900)

print(sent_0)
```

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```
In [89]: # https://stackoverflow.com/questions/16206380/python-beautifulsoup-how-to-rem
         ove-all-tags-from-an-element
         from bs4 import BeautifulSoup
         soup = BeautifulSoup(sent 0, 'lxml')
         text = soup.get_text()
         print(text)
         print("="*50)
         soup = BeautifulSoup(sent_1000, 'lxml')
         text = soup.get text()
         print(text)
         print("="*50)
         soup = BeautifulSoup(sent 1500, 'lxml')
         text = soup.get_text()
         print(text)
         print("="*50)
          soup = BeautifulSoup(sent 4900, 'lxml')
         text = soup.get text()
         print(text)
```

this witty little book makes my son laugh at loud. i recite it in the car as we're driving along and he always can sing the refrain. he's learned about wh ales, India, drooping roses: i love all the new words this book introduces and the silliness of it all. this is a classic book i am willing to bet my son will STILL be able to recite from memory when he is in college

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Can't do sugar. Have tried scores of SF Syrups. NONE of them can touch the excellence of this product. Thick, delicious. Perfect. 3 ingredients: Water, Maltitol, Natural Maple Flavor. PERIOD. No chemicals. No garbage. Have nume rous friends & family members hooked on this stuff. My husband & son, who do NOT like "sugar free" prefer this over major label regular syrup. I use this a s my SWEETENER in baking: cheesecakes, white brownies, muffins, pumpkin pies, etc... Unbelievably delicious... Can you tell I like it?:)

```
In [90]: # https://stackoverflow.com/a/47091490/4084039
import re

def decontracted(phrase):
    # specific
    phrase = re.sub(r"won't", "will not", phrase)
    phrase = re.sub(r"can\'t", "can not", phrase)

# general
    phrase = re.sub(r"\'t", " not", phrase)
    phrase = re.sub(r"\'re", " are", phrase)
    phrase = re.sub(r"\'s", " is", phrase)
    phrase = re.sub(r"\'d", " would", phrase)
    phrase = re.sub(r"\'ll", " will", phrase)
    phrase = re.sub(r"\'t", " not", phrase)
    phrase = re.sub(r"\'t", " have", phrase)
    phrase = re.sub(r"\'ve", " have", phrase)
    phrase = re.sub(r"\'re", " am", phrase)
    return phrase
```

```
In [91]: sent_1500 = decontracted(sent_1500)
    print(sent_1500)
    print("="*50)
```

Great ingredients although, chicken should have been 1st rather than chicken broth, the only thing I do not think belongs in it is Canola oil. Canola or r apeseed is not someting a dog would ever find in nature and if it did find ra peseed in nature and eat it, it would poison them. Today is Food industries h ave convinced the masses that Canola oil is a safe and even better oil than o live or virgin coconut, facts though say otherwise. Until the late 70 is it w as poisonous until they figured out a way to fix that. I still like it but it could be better.

```
In [92]: #remove words with numbers python: https://stackoverflow.com/a/18082370/408403
9
sent_0 = re.sub("\S*\d\S*", "", sent_0).strip()
print(sent_0)
```

this witty little book makes my son laugh at loud. i recite it in the car as we're driving along and he always can sing the refrain. he's learned about wh ales, India, drooping roses: i love all the new words this book introduces and the silliness of it all. this is a classic book i am willing to bet my son will STILL be able to recite from memory when he is in college

Great ingredients although chicken should have been 1st rather than chicken b roth the only thing I do not think belongs in it is Canola oil Canola or rape seed is not someting a dog would ever find in nature and if it did find rapes eed in nature and eat it it would poison them Today is Food industries have c onvinced the masses that Canola oil is a safe and even better oil than olive or virgin coconut facts though say otherwise Until the late 70 is it was pois onous until they figured out a way to fix that I still like it but it could be better

In [94]: # https://gist.github.com/sebleier/554280 # we are removing the words from the stop words list: 'no', 'nor', 'not' #

 ==> after the above steps, we are getting "br br" # we are including them into stop words list # instead of
 if we have
 these tags would have revmoved in the 1st step stopwords= set(['br', 'the', 'i', 'me', 'my', 'myself', 'we', 'our', 'ours', , 'him', 'his', 'himself', \ 'she', "she's", 'her', 'hers', 'herself', 'it', "it's", 'its', 'it self', 'they', 'them', 'their', $\$ 'theirs', 'themselves', 'what', 'which', 'who', 'whom', 'this', 't hat', "that'll", 'these', 'those', \ 'am', 'is', 'are', 'was', 'were', 'be', 'been', 'being', 'have', 'has', 'had', 'having', 'do', 'does', \ 'did', 'doing', 'a', 'an', 'the', 'and', 'but', 'if', 'or', 'becau se', 'as', 'until', 'while', 'of', \ 'at', 'by', 'for', 'with', 'about', 'against', 'between', 'into', 'through', 'during', 'before', 'after',\ 'above', 'below', 'to', 'from', 'up', 'down', 'in', 'out', 'on', 'off', 'over', 'under', 'again', 'further',\ 'then', 'once', 'here', 'there', 'when', 'where', 'why', 'how', 'a 'both', 'each', 'few', 'more',\ ll', 'any', 'most', 'other', 'some', 'such', 'only', 'own', 'same', 'so', 'tha n', 'too', 'very', \ 's', 't', 'can', 'will', 'just', 'don', "don't", 'should', "shoul d've", 'now', 'd', 'll', 'm', 'o', 're', \ 've', 'y', 'ain', 'aren', "aren't", 'couldn', "couldn't", 'didn', "didn't", 'doesn', "doesn't", 'hadn',\ "hadn't", 'hasn', "hasn't", 'haven', "haven't", 'isn', "isn't", 'm a', 'mightn', "mightn't", 'mustn',\ "mustn't", 'needn', "needn't", 'shan', "shan't", 'shouldn', "shoul dn't", 'wasn', "wasn't", 'weren', "weren't", \ 'won', "won't", 'wouldn', "wouldn't"])

```
In [95]: # Combining all the above stundents
           from tadm import tadm
           preprocessed reviews = []
           # tqdm is for printing the status bar
           for sentance in tqdm(final['Text'].values):
               sentance = re.sub(r"http\S+", "", sentance)
               sentance = BeautifulSoup(sentance, 'lxml').get_text()
               sentance = decontracted(sentance)
               sentance = re.sub("\S*\d\S*", "", sentance).strip()
sentance = re.sub('[^A-Za-z]+', ' ', sentance)
               # https://gist.github.com/sebleier/554280
               sentance = ' '.join(e.lower() for e in sentance.split() if e.lower() not i
           n stopwords)
               preprocessed reviews.append(sentance.strip())
          100%|
           364171/364171 [05:04<00:00, 1195.86it/s]
 In [96]: preprocessed reviews[1500]
 Out[96]: 'great ingredients although chicken rather chicken broth thing not think belo
          ngs canola oil canola rapeseed not someting dog would ever find nature find r
          apeseed nature eat would poison today food industries convinced masses canola
          oil safe even better oil olive virgin coconut facts though say otherwise late
          poisonous figured way fix still like could better'
 In [97]: | final['cleaned_text']=preprocessed_reviews
 In [98]: final.shape
 Out[98]: (364171, 11)
 In [99]: | final["Score"].value_counts()
 Out[99]: 1
                307061
                 57110
          Name: Score, dtype: int64
In [100]:
          #Getting positive and negetive data
           data_pos = final[final["Score"] == 1].sample(n = 50000)
           data neg = final[final["Score"] == 0].sample(n = 50000)
           final1 = pd.concat([data pos, data neg])
           final1.shape
Out[100]: (100000, 11)
```

```
In [101]: #Sorted the data based on time and took 100k data points
    final1["Time"] = pd.to_datetime(final1["Time"], unit = "s")
    final1 = final1.sort_values(by = "Time")
    final1.head()
```

Out[101]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	ŀ
346041	374343	B00004Cl84	A1B2IZU1JLZA6	Wes	19	2
417901	451923	B00004CXX9	ANIMV3SPDD8SH	Guy De Federicis	1	
346037	374339	B00004Cl84	AZRJH4JFB59VC	Lynwood E. Hines	21	2
346031	374333	B00004Cl84	A1CZICCYP2M5PX	Christian Pelchat	0	(
138691	150509	0006641040	A3CMRKGE0P909G	Teresa	3	2

```
In [102]: Y = final1['Score'].values
    X = final1['cleaned_text'].values
    print(Y.shape)
    print(type(Y))
    print(X.shape)
    print(type(X))

(100000,)
    <class 'numpy.ndarray'>
        (100000,)
        <class 'numpy.ndarray'>
```

```
In [103]: # split the data set into train and test
X_Train, X_Test, Y_Train, Y_Test = train_test_split(X,Y,test_size=0.3, random_state=12, shuffle = False)

# split the train data set into cross validation train and cross validation te
st
X_tr, X_cv, Y_tr, Y_cv = train_test_split(X,Y, test_size=0.3, random_state=12,
shuffle = False)

print('='*100)
print("After splitting")
print("X_Train Shape:",X_Train.shape, "Y_Train Shape:",Y_Train.shape)
print("X_cv Shape:",X_cv.shape, "Y_cv Shape",Y_cv.shape)
print("X_Test Shape",X_Test.shape, "Y_Test Shape",Y_Test.shape)
```

After splitting
X_Train Shape: (70000,) Y_Train Shape: (70000,)
X_cv Shape: (30000,) Y_cv Shape (30000,)
X_Test Shape (30000,) Y_Test Shape (30000,)

[4] Featurization

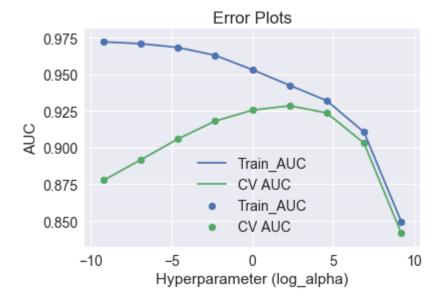
[4.1] Bag Of Words

```
In [104]:
         #BoW
         count vect = CountVectorizer() #in scikit-learn
         count vect.fit(X Train)
         print("some feature names ", count vect.get feature names()[:10])
         X Train Bow = count vect.transform(X Train)
         X_Test_Bow = count_vect.transform(X_Test)
         X CV Bow = count vect.transform(X cv)
         print('='*50)
         #final counts = count vect.transform(X Test)
         print("the type of X Train : ",type(X_Train_Bow))
         print("the shape of Train BOW vectorizer ",X_Train_Bow.get_shape())
         print("the shape of Test BOW vectorizer ",X Test Bow.get shape())
         print("the shape of CV BOW vectorizer ",X_CV_Bow.get_shape())
         #print("the number of unique words ", final_counts.get_shape()[1])
         aaaaaaaaaaaaaaaaaaa', 'aaaaaaarrrrrggghhh', 'aaaaaaahhhhhyaaaaaaa', 'aaaallll',
         'aaaand', 'aaah']
         _____
         the type of X Train : <class 'scipy.sparse.csr.csr matrix'>
         the shape of Train BOW vectorizer (70000, 51472)
         the shape of Test BOW vectorizer (30000, 51472)
         the shape of CV BOW vectorizer (30000, 51472)
```

[4.1.1] AUC Curve Plot

```
In [105]:
          import math
          from sklearn.naive bayes import MultinomialNB
          train AUC = []
          CV AUC = []
          alpha = [10**-4, 10**-3, 10**-2, 10**-1, 1, 10**1, 10**2, 10**3, 10**4]
          log alpha=[]
          for i in tqdm(alpha):
              MNB = MultinomialNB(alpha = i, class_prior=[0.5,0.5], fit_prior=True)
              #fit a model on train BOW vectorizer
              MNB.fit(X_Train_Bow, Y_Train)
              #predict probabilities on train BOW vectorizer
              Y Train Pred = MNB.predict proba(X Train Bow)[:,1]
              #predict probabilities on Cross validation BOW vectorizer
              Y_CV_Pred = MNB.predict_proba(X_CV_Bow)[:,1]
              #calculate AUC score
              train_AUC.append(roc_auc_score(Y_Train,Y_Train_Pred))
              CV_AUC.append(roc_auc_score(Y_cv, Y_CV_Pred))
              log alpha.append(math.log(i))
          plt.plot(log_alpha, train_AUC, label='Train_AUC')
          plt.scatter(log alpha, train AUC, label='Train AUC')
          plt.plot(log_alpha, CV_AUC, label='CV AUC')
          plt.scatter(log_alpha, CV_AUC, label='CV AUC')
          plt.legend()
          plt.xlabel('Hyperparameter (log alpha)')
          plt.ylabel('AUC')
          plt.title('Error Plots')
          plt.show()
```

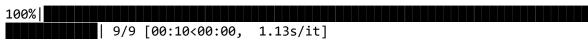




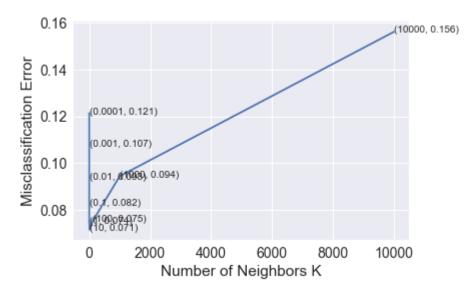
```
In [106]:
           #Finding the best hyper paramter using 10-fold cross validation data
          def Optimal_Alpha(X_Train, Y_Train):
                   #Considered a wide range of alpha values for hyperparameter tuning, s
          tart as low 10^-4 to 10^4.
                  # empty list that will hold cv scores
                  cv scores = []
                  # perform 10-fold cross validation
                  for i in tqdm(alpha):
                      MNB = MultinomialNB(alpha=i, class prior=[0.5,0.5], fit prior=True
                      scores = cross_val_score(MNB, X_Train, Y_Train, cv=10, scoring='ro
          c_auc')
                      cv scores.append(scores.mean())
                  # changing to misclassification error
                  MSE = [1 - x \text{ for } x \text{ in } cv \text{ scores}]
                  # determining best k
                  bestAlpha = alpha[MSE.index(min(MSE))]
                  print('\nThe optimal number of neighbors is %d.' % bestAlpha)
                  # plot misclassification error vs k
                  plt.plot(alpha, MSE)
                  for xy in zip(alpha, np.round(MSE,3)):
                      plt.annotate('(%s, %s)' % xy, xy=xy, textcoords='data')
                  plt.xlabel('Number of Neighbors K')
                  plt.ylabel('Misclassification Error')
                  plt.show()
                  print("the misclassification error for each k value is : ", np.round(M
          SE,3))
                  return bestAlpha
```

[4.1.2] 10-fold cross validation, determining best Alpha

```
In [107]: optimal_alpha_bow = Optimal_Alpha(X_Train_Bow, Y_Train)
    print("_"*100)
    print("optimal_alpha:", optimal_alpha_bow)
    print("_"*100)
```



The optimal number of neighbors is 10.



the misclassification error for each k value is : [0.121 0.107 0.093 0.082 0.074 0.071 0.075 0.094 0.156]

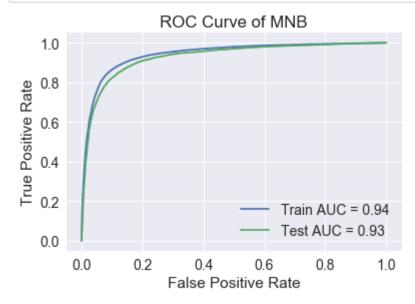
```
optimal alpha: 10
```

```
In [108]: optimal_model = MultinomialNB(alpha=optimal_alpha_bow, class_prior=[0.5,0.5],
    fit_prior=True)
    optimal_model.fit(X_Train_Bow, Y_Train)
    prediction = optimal_model.predict(X_Test_Bow)
    optimal_model
```

Out[108]: MultinomialNB(alpha=10, class_prior=[0.5, 0.5], fit_prior=True)

[4.1.3] ROC Curve of Naive Bayes

```
In [109]:
          #with the reference of below link:
          #https://stackoverflow.com/questions/52910061/implementing-roc-curves-for-k-nn
          -machine-learning-algorithm-using-python-and-sci
          #predict probabilities on X Train Bow and X Test Bow and pass as param to roc
          curve to find roc curve
          Train_FPR, Train_TPR, threshold = roc_curve(Y_Train, optimal_model.predict_pro
          ba(X Train_Bow)[:,1])
          Test FPR, Test TPR, threshold = roc curve(Y Test, optimal model.predict proba(
          X Test Bow)[:,1])
          roc_auc = auc(Train_FPR, Train_TPR)
          roc auc1 = auc(Test FPR, Test TPR)
          plt.plot(Train_FPR, Train_TPR, label = 'Train AUC = %0.2f' % roc_auc)
          plt.plot(Test_FPR, Test_TPR, label = 'Test AUC = %0.2f' % roc_auc1)
          plt.legend()
          plt.xlabel('False Positive Rate')
          plt.ylabel('True Positive Rate')
          plt.title('ROC Curve of MNB')
          plt.show()
```



[4.1.4]Train and Test Accuracy

```
In [110]: Training_Accuracy_Bow = optimal_model.score(X_Train_Bow, Y_Train)
    print('Training_Accuracy=%0.3f'%Training_Accuracy_Bow)
    Training_Error_Bow = 1 - Training_Accuracy_Bow
    print('Training_Error=%0.3f'%Training_Error_Bow)

Test_Accuracy_Bow = accuracy_score(Y_Test, prediction)
    print('Test_Accuracy=%0.3f'%Test_Accuracy_Bow)
    Test_Error_Bow = 1 - Test_Accuracy_Bow
    print('Test_Error=%0.3f'%Test_Error_Bow)
    print('\nThe accuracy of the MNB classifier for k = %d is %f%%' % (optimal_alp ha_bow, Test_Accuracy_Bow))
```

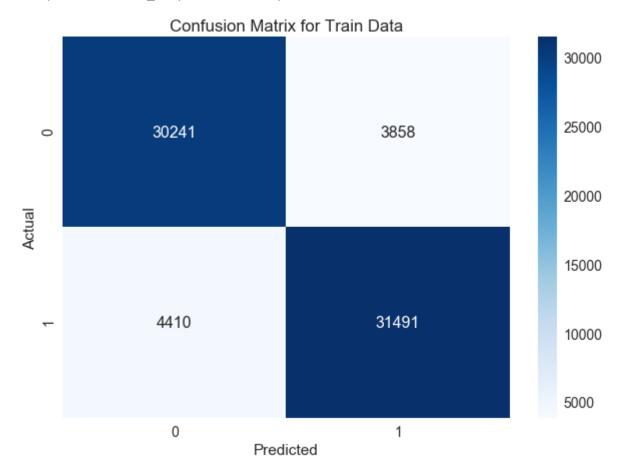
Training_Accuracy=0.882 Training_Error=0.118 Test_Accuracy=0.865 Test_Error=0.135

The accuracy of the MNB classifier for k = 10 is 0.864900%

[4.1.5] Confusion Matrix

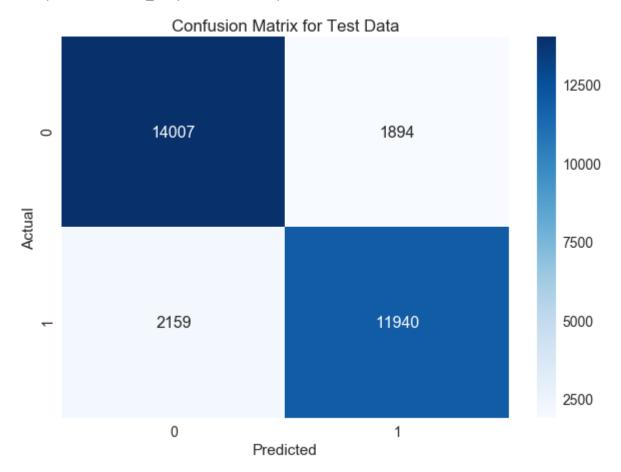
In [111]: from sklearn.metrics import confusion_matrix
 conf_matrix = confusion_matrix(Y_Train, optimal_model.predict(X_Train_Bow))
 df_conf_matrix = pd.DataFrame(conf_matrix, columns=np.unique(Y_Train), index=n
 p.unique(Y_Train))
 df_conf_matrix.index.name = 'Actual'
 df_conf_matrix.columns.name = 'Predicted'
 plt.figure(figsize=(10,7))
 plt.title("Confusion Matrix for Train Data")
 sns.set(font_scale=1.4)
 sns.heatmap(df_conf_matrix, cmap='Blues', annot=True, annot_kws={'size':16}, f
 mt='d')

Out[111]: <matplotlib.axes._subplots.AxesSubplot at 0x1c98daea6a0>



```
In [112]: #With the reference of below link:
    #https://www.kaggle.com/agungor2/various-confusion-matrix-plots
    from sklearn.metrics import confusion_matrix
    conf_matrix = confusion_matrix(Y_Test, optimal_model.predict(X_Test_Bow))
    df_conf_matrix = pd.DataFrame(conf_matrix, columns=np.unique(Y_Test), index=np
    .unique(Y_Test))
    df_conf_matrix.index.name = 'Actual'
    df_conf_matrix.columns.name = 'Predicted'
    plt.figure(figsize=(10,7))
    plt.title("Confusion Matrix for Test Data")
    sns.set(font_scale=1.4)
    sns.heatmap(df_conf_matrix, cmap='Blues', annot=True, annot_kws={'size':16}, f
    mt='d')
```

Out[112]: <matplotlib.axes._subplots.AxesSubplot at 0x1c98daea2b0>



[4.1.6] Classification Report

```
In [113]: from sklearn.metrics import classification_report
    print(classification_report(Y_Test, prediction))
```

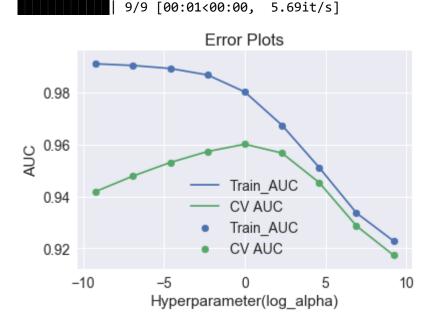
```
precision
                           recall f1-score
                                               support
          0
                  0.87
                             0.88
                                        0.87
                                                  15901
          1
                   0.86
                             0.85
                                        0.85
                                                  14099
                             0.86
                                        0.86
                                                  30000
avg / total
                  0.86
```

[4.2] TF-IDF

```
In [114]:
          #TF-IDF
          tf idf vect = TfidfVectorizer(ngram range=(1,2), min df=5)
          tf idf vect.fit transform(X Train)
          print("some sample features(unique words in the corpus)", tf idf vect.get featu
          re names()[0:10])
          print('='*50)
          X Train TfIdf = tf idf vect.transform(X Train)
          X_Test_TfIdf = tf_idf_vect.transform(X_Test)
          X_CV_TfIdf = tf_idf_vect.transform(X_cv)
          #final tf idf = tf idf vect.transform(X Test)
          print("the type of count vectorizer ",type(X_Train_TfIdf))
          print("the shape of out text TFIDF vectorizer ",X_Train_TfIdf.get_shape())
          print("the shape of out text TFIDF vectorizer ",X_Test_TfIdf.get_shape())
          print("the shape of out text TFIDF vectorizer ",X_CV_TfIdf.get_shape())
          #print("the number of unique words including both unigrams and bigrams ", fina
          l tf idf.get shape()[1])
          some sample features(unique words in the corpus) ['aa', 'aaa', 'aafco', 'abac
          k', 'abandon', 'abandoned', 'abbey', 'abc', 'abdomen', 'abdominal']
          _____
          the type of count vectorizer <class 'scipy.sparse.csr.csr_matrix'>
          the shape of out text TFIDF vectorizer (70000, 92735)
          the shape of out text TFIDF vectorizer (30000, 92735)
          the shape of out text TFIDF vectorizer (30000, 92735)
```

[4.2.1] AUC Curve Plot

```
In [115]:
          import math
          from sklearn.naive bayes import MultinomialNB
          train AUC = []
          CV AUC = []
          alpha = [10**-4, 10**-3, 10**-2, 10**-1, 1,10**1, 10**2, 10**3, 10**4]
          log alpha tfidf=[]
          for i in tqdm(alpha):
              MNB = MultinomialNB(alpha = i, class_prior=[0.5,0.5], fit_prior=True)
              #fit a model on train BOW vectorizer
              MNB.fit(X_Train_TfIdf, Y_Train)
              #predict probabilities on train BOW vectorizer
              Y Train Pred = MNB.predict proba(X Train TfIdf)[:,1]
              #predict probabilities on Cross validation BOW vectorizer
              Y_CV_Pred = MNB.predict_proba(X_CV_TfIdf)[:,1]
              #calculate AUC score
              train_AUC.append(roc_auc_score(Y_Train,Y_Train_Pred))
              CV_AUC.append(roc_auc_score(Y_cv, Y_CV_Pred))
              log alpha tfidf.append(math.log(i))
          plt.plot(log_alpha_tfidf, train_AUC, label='Train_AUC')
          plt.scatter(log alpha tfidf, train AUC, label='Train AUC')
          plt.plot(log_alpha_tfidf, CV_AUC, label='CV AUC')
          plt.scatter(log_alpha_tfidf, CV_AUC, label='CV AUC')
          plt.legend()
          plt.xlabel('Hyperparameter(log alpha)')
          plt.ylabel('AUC')
          plt.title('Error Plots')
          plt.show()
```



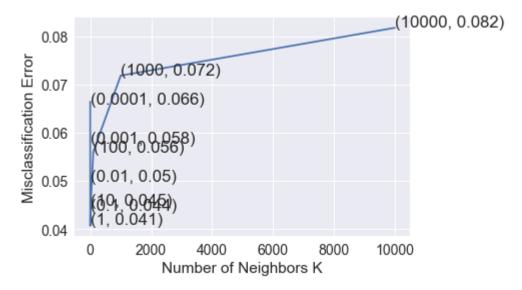
[4.2.2] 10-fold cross validation, determining best Alpha

100%

```
In [116]: optimal_alpha_tfidf = Optimal_Alpha(X_Train_TfIdf, Y_Train)
    print("optimal_alpha:", optimal_alpha_tfidf)
100%
```

The optimal number of neighbors is 1.

| 9/9 [00:15<00:00,

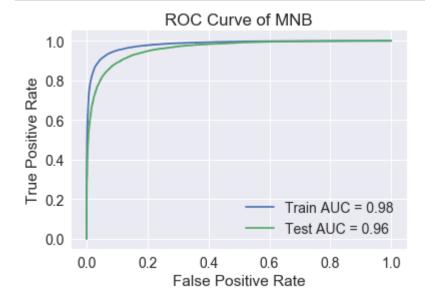


the misclassification error for each k value is : [0.066 0.058 0.05 0.044 0.041 0.045 0.056 0.072 0.082] optimal_alpha: 1

```
In [117]: optimal_model1 = MultinomialNB(alpha=optimal_alpha_tfidf, class_prior=[0.5,0.5
    ], fit_prior=True)
    optimal_model1.fit(X_Train_TfIdf, Y_Train)
    prediction = optimal_model1.predict(X_Test_TfIdf)
```

[4.2.3] ROC Curve of Naive Bayes

```
In [118]:
          #with the reference of below link:
          #https://stackoverflow.com/questions/52910061/implementing-roc-curves-for-k-nn
          -machine-learning-algorithm-using-python-and-sci
          #predict probabilities on X Train Bow and X Test Bow and pass as param to roc
          curve to find roc curve
          Train_FPR, Train_TPR, threshold = roc_curve(Y_Train, optimal_model1.predict_pr
          oba(X_Train_TfIdf)[:,1])
          Test FPR, Test TPR, threshold = roc curve(Y Test, optimal model1.predict proba
          (X Test TfIdf)[:,1])
          roc_auc2 = auc(Train_FPR, Train_TPR)
          roc auc3 = auc(Test FPR, Test TPR)
          plt.plot(Train_FPR, Train_TPR, label = 'Train AUC = %0.2f' % roc_auc2)
          plt.plot(Test_FPR, Test_TPR, label = 'Test AUC = %0.2f' % roc_auc3)
          plt.legend()
          plt.xlabel('False Positive Rate')
          plt.ylabel('True Positive Rate')
          plt.title('ROC Curve of MNB')
          plt.show()
```



[4.2.4]Train and Test Accuracy

```
In [119]: Training_Accuracy_tfidf = optimal_model1.score(X_Train_TfIdf, Y_Train)
    print('Training_Accuracy=%0.3f'%Training_Accuracy_tfidf)
    Training_Error_tfidf = 1 - Training_Accuracy_tfidf
    print('Training_Error=%0.3f'%Training_Error_tfidf)

Test_Accuracy_tfidf = accuracy_score(Y_Test, prediction)
    print('Test_Accuracy=%0.3f'%Test_Accuracy_tfidf)
    Test_Error_tfidf = 1 - Test_Accuracy_tfidf
    print('Test_Error=%0.3f'%Test_Error_tfidf)
    print('NoThe accuracy of the MNB classifier for k = %d is %f%%' % (optimal_alp ha_tfidf, Test_Accuracy_tfidf))
```

Training_Accuracy=0.931 Training_Error=0.069 Test_Accuracy=0.895 Test_Error=0.105

The accuracy of the MNB classifier for k = 1 is 0.895233%

[4.2.5] Confusion Matrix

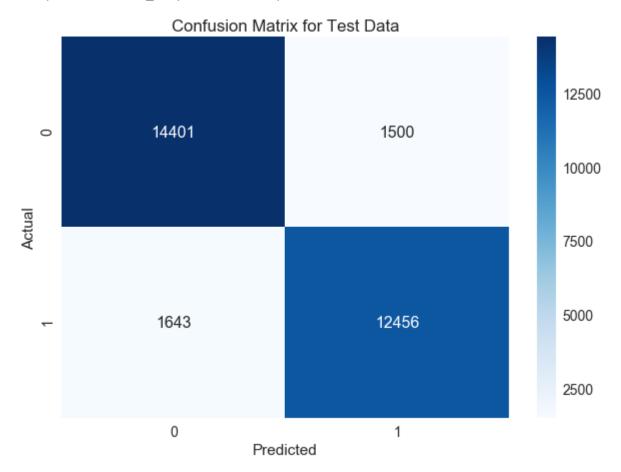
In [120]: from sklearn.metrics import confusion_matrix
 conf_matrix = confusion_matrix(Y_Train, optimal_model1.predict(X_Train_TfIdf))
 df_conf_matrix = pd.DataFrame(conf_matrix, columns=np.unique(Y_Train), index=n
 p.unique(Y_Train))
 df_conf_matrix = pd.DataFrame(conf_matrix, columns=np.unique(Y_Test), index=np
 .unique(Y_Test))
 df_conf_matrix.index.name = 'Actual'
 df_conf_matrix.columns.name = 'Predicted'
 plt.figure(figsize=(10,7))
 plt.title("Confusion Matrix for Train Data")
 sns.set(font_scale=1.4)
 sns.heatmap(df_conf_matrix, cmap='Blues', annot=True, annot_kws={'size':16}, f
 mt='d')

Out[120]: <matplotlib.axes._subplots.AxesSubplot at 0x1c9ad96be48>



```
In [121]: #With the reference of below link:
    #https://www.kaggle.com/agungor2/various-confusion-matrix-plots
    from sklearn.metrics import confusion_matrix
    conf_matrix = confusion_matrix(Y_Test, optimal_model1.predict(X_Test_TfIdf))
    df_conf_matrix = pd.DataFrame(conf_matrix, columns=np.unique(Y_Test), index=np
    .unique(Y_Test))
    df_conf_matrix.index.name = 'Actual'
    df_conf_matrix.columns.name = 'Predicted'
    plt.figure(figsize=(10,7))
    plt.title("Confusion Matrix for Test Data")
    sns.set(font_scale=1.4)
    sns.heatmap(df_conf_matrix, cmap='Blues', annot=True, annot_kws={'size':16}, f
    mt='d')
```

Out[121]: <matplotlib.axes._subplots.AxesSubplot at 0x1c98ddf93c8>



[4.2.6]Classification Report

In [122]: from sklearn.metrics import classification_report
 print(classification_report(Y_Test, prediction))

	precision	recall	f1-score	support
0	0.90	0.91	0.90	15901
1	0.89	0.88	0.89	14099
avg / total	0.90	0.90	0.90	30000

[5] Feature Importance

[5.1]BoW

[5.1.1]Feature Importance for Positive Class

```
In [123]:
          #https://stackoverflow.com/questions/35353150/sklearn-multinomialnb-how-to-fin
          d-most-distinguish-word-in-class
          import operator
          pos imp features = MNB.feature log prob [1,:]
          neg imp features = MNB.feature log prob [0,:]
          print("pos_imp_features",len(pos_imp_features))
          imp features = {}
          feature names= count vect.get feature names()
          print("feature_names",len(feature_names))
          for i in range(len(feature_names)):
              imp features[feature names[i]] = pos imp features[i]
          names_diff_sorted = sorted(imp_features.items(), key = operator.itemgetter(1),
          reverse = True)
          print("Postive top 25 important features are:")
          for i in range(25):
              print(names_diff_sorted[i])
          pos imp features 92735
          feature names 51472
          Postive top 25 important features are:
          ('pupils', -11.36733577032401)
          ('portugal', -11.376557440094475)
           ('sueggestion', -11.38457326712038)
          ('touches', -11.384717570341618)
          ('eventhe', -11.387648758776214)
          ('mucus', -11.395026781036709)
          ('caseits', -11.398145828536254)
          ('aromaticum', -11.401794080610495)
          ('minnesotans', -11.40278108425811)
          ('patronize', -11.404991654726802)
          ('terible', -11.407241317216132)
          ('notgrab', -11.407434579080615)
          ('andexpected', -11.409008854163199)
          ('unfortunatedly', -11.40912985959799)
          ('eic', -11.410060700688014)
          ('havesomething', -11.410297156549989)
          ('curative', -11.410541993801118)
          ('cement', -11.412547064243359)
          ('ivf', -11.413818900871217)
          ('oligofructose', -11.414011561257972)
          ('ordinairy', -11.414444626592093)
          ('truthaboutpetfood', -11.414916458198851)
          ('medifast', -11.415195577824997)
          ('implemented', -11.41590031352609)
```

[5.1.2]Feature Importance for Negetive Class

('leek', -11.416330419652333)

```
In [124]: for i in range(len(feature_names)):
    imp_features[feature_names[i]] = neg_imp_features[i]
    names_diff_sorted = sorted(imp_features.items(), key = operator.itemgetter(1),
    reverse = True)
    print("\n\nNegative top 25 important features are:")
    for i in range(25):
        print(names_diff_sorted[i])
```

```
Negative top 25 important features are:
('sueggestion', -11.370150161443856)
('eventhe', -11.39072028377526)
('mucus', -11.396886574131914)
('portugal', -11.399387104745102)
('curative', -11.4027792525322)
('leek', -11.404987007226316)
('patronize', -11.405735198438393)
('aromaticum', -11.405991007765396)
('complainingsince', -11.407246406855636)
('notgrab', -11.407902666580782)
('colgin', -11.409226508302964)
('boson', -11.409860317452424)
('fudge', -11.409926980310622)
('eic', -11.41088133161944)
('hunting', -11.41329207617512)
('cement', -11.413351001752769)
('xylan', -11.413439178898727)
('moneys', -11.414438331464124)
('ivf', -11.415050398946581)
('procuct', -11.415477110020493)
('uji', -11.415754640459634)
('brainwashing', -11.41625069673083)
('soulful', -11.417110369095674)
('implemented', -11.417996791859633)
('touches', -11.418000951237648)
```

[5.2]TF-IDF

[5.2.1]Feature Importance for Positive Class

```
In [125]:
          #https://stackoverflow.com/questions/35353150/sklearn-multinomialnb-how-to-fin
          d-most-distinguish-word-in-class
          pos imp features = MNB.feature log prob [1,:]
          print("length of pos_imp_features:",len(pos_imp_features))
          imp_features = {}
          feature names= tf idf vect.get feature names()
          print("length of feature names:",len(feature names))
          for i in range(len(feature names)):
               imp_features[feature_names[i]] = pos_imp_features[i]
          #Introduced exception handling to avoid the index error.
            # try:
            #
                    imp_features[feature_names[i]] = pos_imp_features[i]
              except IndexError as error:
                   pass
          names_diff_sorted = sorted(imp_features.items(), key = operator.itemgetter(1),
          reverse = True)
          print("Postive top 25 important features are:\n")
          for i in range(25):
              print(names diff sorted[i])
          length of pos imp features: 92735
          length of feature names: 92735
          Postive top 25 important features are:
          ('not', -11.35577541247079)
          ('great', -11.36733577032401)
          ('good', -11.376557440094475)
          ('tea', -11.38302202715892)
          ('like', -11.38457326712038)
          ('love', -11.384717570341618)
          ('coffee', -11.387648758776214)
          ('one', -11.393973886250503)
          ('product', -11.394467023374107)
           ('taste', -11.394784644046778)
          ('flavor', -11.395026781036709)
          ('best', -11.398145828536254)
          ('amazon', -11.401794080610495)
          ('find', -11.40278108425811)
          ('price', -11.403080957515396)
          ('would', -11.403876351494166)
           ('use', -11.404032250369402)
          ('really', -11.40434932692268)
          ('get', -11.404991654726802)
          ('little', -11.407241317216132)
          ('food', -11.407434579080615)
          ('time', -11.407806419298154)
          ('no', -11.408174829534856)
          ('much', -11.408314533103715)
          ('also', -11.409008854163199)
```

[5.2.2]Feature Importance for Negetive Class

```
In [126]:
          neg imp features = MNB.feature log prob [0,:]
          print("length of neg_imp_features:",len(neg_imp_features))
          print("length of feature names:",len(feature names))
          for i in range(len(feature names)):
              imp features[feature names[i]] = neg imp features[i]
             # try:
              #
                    imp features[feature names[i]] = neg imp features[i]
             # except IndexError as error:
                    pass
          names_diff_sorted = sorted(imp_features.items(), key = operator.itemgetter(1),
          reverse = True)
          print("\n\nNegative top 25 important features are:\n")
          for i in range(25):
              print(names diff sorted[i])
          length of neg_imp_features: 92735
          length of feature names: 92735
          Negative top 25 important features are:
          ('not', -11.300193735430538)
          ('like', -11.370150161443856)
          ('taste', -11.378288708636068)
          ('product', -11.378325867831162)
          ('would', -11.380567943280552)
          ('coffee', -11.39072028377526)
          ('one', -11.391367231022137)
          ('flavor', -11.396886574131914)
          ('no', -11.397739097994315)
          ('good', -11.399387104745102)
          ('tea', -11.400843728957796)
           ('buy', -11.4027792525322)
          ('even', -11.404987007226316)
          ('get', -11.405735198438393)
          ('amazon', -11.405991007765396)
          ('box', -11.407246406855636)
          ('food', -11.407902666580782)
          ('much', -11.408213972903438)
          ('really', -11.408786770340077)
          ('bought', -11.409226508302964)
          ('bad', -11.409860317452424)
          ('could', -11.409926980310622)
          ('chocolate', -11.41088133161944)
          ('tried', -11.411338134884247)
          ('disappointed', -11.41329207617512)
```

```
In [127]: from prettytable import PrettyTable
    comparision = PrettyTable()
    comparision.field_names = ["S.NO","Vectorizer", "Hyperparameter", "Training Er
    ror", "Test Error","AUC"]
    comparision.add_row(["1","BoW", optimal_alpha_bow, Training_Error_Bow, Test_Er
    ror_Bow, np.round(float(roc_auc1),3)])
    comparision.add_row(["2","TF-IDF", optimal_alpha_tfidf,Training_Error_tfidf, T
    est_Error_tfidf, np.round(float(roc_auc3),3)])
    print(comparision)
```

S.NO Vectorizer AUC	Hyperparameter	+ Training Error	Test Error
+ 1 BoW 0.929	10	0.11811428571428573	
2 TF-IDF 667 0.96	1	0.06875714285714285	•
+	•	•	•

Conclusion

- 1. Applied Multinomial NB on two feature sets 1. Bow and 2. TF-IDF vectorizer.
- 2. Sorted the data based on Time and Considered 100 K data points for Training set 70K, Test set: 30K
- 3. Used AUC as a metric for hyperparameter tuning. And took the range of alpha values from (10^-4 to 10^4).
- 4. Found the top 25 features of positive class and top 25 features of negative class for Bow and TF-IDF feature sets.
- 5. With reference to the pretty table, here is my understanding: a. Naive Bayes by using Bow and TF-IDF with best alpha = 10 and AUC score is 0.92 in both.
- 6. Since it is an imbalanced data, not considered accuracy as a metric and AUC Score is high
- 7. Plotted Confusion matrix for both Training and Test set.

Note: I do not have idea on feature engineering concept, One after completing the Module 5 and able to check the accuracy of this model with feature engineering.