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)

EDA: https://nycdatascience.com/blog/student-works/amazon-fine-foods-visualization/ (https://nycdatascience.com/blog/student-works/amazon-fine-foods-visualization/ (https://nycdatascience.com/blog/student-works/amazon-fine-foods-visualization/ (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. ProductId 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 re view 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 Score/Rating. A rating of 4 or 5 can be cosnidered as a positive review. A rating of 1 or 2 can be considered as negative one. A review of rating 3 is considered nuetral and such reviews are ignored from our analysis. This is an approximate and proxy way of determining the polarity (positivity/negativity) of a review.

[1] Reading Data

[1.1] Loading the data

The dataset is available in two forms

.csv file SQLite Database

In order to load the data, We have used the SQLITE dataset as it is 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 is above 3, then the recommendation will be set to "positive". Otherwise, it will be set to "negative".

```
In [36]: %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.preprocessing import MaxAbsScaler
         from sklearn.model selection import train_test_split
         from sklearn.cross validation import train test split
         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
         import tqdm
         #from tqdm import tqdm
         import os
         # NB imports
         from sklearn.naive_bayes import MultinomialNB
         from sklearn.model_selection import TimeSeriesSplit, GridSearchCV
         #metrics import
         from sklearn.metrics import confusion_matrix
         from sklearn.metrics import accuracy_score
         from sklearn.metrics import roc_auc_score
         from sklearn.metrics import auc
         from sklearn.metrics import roc_curve
         from sklearn.metrics import recall_score
         from sklearn.metrics import precision_score
         from sklearn.metrics import f1_score
```

```
In [2]: con = sqlite3.connect('database.sqlite')
        #Normal for every dataset contains a score from 1 to 5.
        #The review must classified as positive or a negative reviews. Neutral reviews s
        uch score = 3 are omitted.
        #We consider scores 1,2 as negative and 4,5 as positive. So we will consider revi
        ews which are not equal to 3.
        filtered_data = pd.read_sql_query("SELECT * FROM Reviews WHERE Score != 3 LIMIT
        150000", con)
        filtered data.head(5)
```

Out[2]:		ld	Productid	Userld	ProfileName	HelpfulnessNumerator	HelpfulnessDenominator	Sco
	0	1	B001E4KFG0	A3SGXH7AUHU8GW	delmartian	1	1	
	1	2	B00813GRG4	A1D87F6ZCVE5NK	dll pa	0	0	
	2	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1	1	
	3	4	B000UA0QIQ	A395BORC6FGVXV	Karl	3	3	
	4	5	B006K2ZZ7K	A1UQRSCLF8GW1T	Michael D. Bigham "M. Wassir"	0	0	

```
In [3]: # For Each review the scores will be 1,2 and 4,5.
         # We will mark negative review as 0 and positive reviews as 1 for classification
        purposes
        def partition(x) :
            if x > 3:
                return 1
            return 0
        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 (150000, 10)
Out[3]:
           ld
                 ProductId
                                   UserId ProfileName HelpfulnessNumerator HelpfulnessDenominator Sco
```

 0
 1
 B0001E4KFG0
 A3SGXH7AUHU8GW
 delmartian
 1
 1

 1
 2
 B00813GRG4
 A1D87F6ZCVE5NK
 dll pa
 0
 0

 2
 3
 B000LQOCH0
 ABXLMWJIXXAIN
 Natalia Corres "Natalia Corres"
 1
 1

```
In [4]: #Checking the database statistics
filtered_data.describe()
```

Out[4]:

	ld	HelpfulnessNumerator	HelpfulnessDenominator	Score	Time
count	150000.000000	150000.000000	150000.000000	150000.000000	1.500000e+05
mean	81466.351767	1.683353	2.146233	0.842147	1.296091e+09
std	46958.766654	6.825579	7.418456	0.364605	4.766398e+07
min	1.000000	0.000000	0.000000	0.000000	9.393408e+08
25%	40734.750000	0.000000	0.000000	1.000000	1.270598e+09
50%	81583.500000	0.000000	1.000000	1.000000	1.310947e+09
75%	122047.250000	2.000000	2.000000	1.000000	1.332547e+09
max	162717.000000	559.000000	562.000000	1.000000	1.351210e+09

2. Exploratory Data Analysis

2.1 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 [5]: display= pd.read_sql_query("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND UserId='ABXLMWJIXXAIN'
    ORDER BY ProductID
    """, con)
    display.head()
```

Out[5]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	HelpfulnessDenominator	s
0	320691	B000CQ26E0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	0	0	
1	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1	1	
2	468954	B004DMGQKE	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	0	0	

From the above output we could observe that For a single Userld there are multiple reviews. So we need to sort the data and delete the duplicate records based on the ProductId and UserId

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 [9]: display= pd.read_sql_query("""
         SELECT *
         FROM Reviews
         WHERE Score != 3 AND HelpfulnessNumerator > HelpfulnessDenominator
         ORDER BY ProductID
         """, con)
         display.head()
Out[9]:
               ld
                     ProductId
                                      Userld ProfileName HelpfulnessNumerator HelpfulnessDenominator Sc
                                                  J. E.
          0 64422 B000MIDROQ A161DK06JJMCYF
                                               Stephens
                                                                      3
                                                                                         1
                                               "Jeanne"
          1 44737 B001EQ55RW A2V0I904FH7ABY
                                                  Ram
In [10]: | final=final[final.HelpfulnessNumerator<=final.HelpfulnessDenominator]</pre>
In [11]: #Before starting the next phase of preprocessing lets see the number of entries
         left
         print(final.shape)
         score = final['Score']
         #How many positive and negative reviews are present in our dataset?
         final['Score'].value_counts()
         (126357, 10)
Out[11]: 1
             106326
              20031
         Name: Score, dtype: int64
```

3. Preprocessing

3.1 Preprocessing Review Text

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:-

```
Begin by removing the html tags

Remove any punctuations or limited set of special characters like , or . or # etc.

Check if the word is made up of english letters and is not alpha-numeric

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)

Convert the word to lowercase

Remove Stopwords

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 [12]: sent_100 = final['Text'].values[100]
    print("="*50)
    print(sent_100)
    print("="*50)
```

My only complaint about this product is that sometimes it is difficult to sque eze just a very small amount from the bottle into the mixture to be colored. I don't do massive amounts of baking, and I find that sometimes it is really h ard to get a very light color when adding this product to very small quantities of mixture. However, the colors are very bright and I was very pleased to f ind that the color is evenly distributed throughout the mixture. I thought the is product was a little pricy, but after trying it out, I do believe that it w as worth the extra money. The product seems of high quality and it shows in my cooking whenever I want to add some color to things.

phrase = re.sub(r"\'ve", " have", phrase)
phrase = re.sub(r"\'m", " am", phrase)

return phrase

```
In [13]: def cleanhtml(sentence):
    sent = re.sub(r"http\S+", "", sentence)
    return sent

In [14]: 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"n\'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"\'d", " will", phrase)
    phrase = re.sub(r"\'t", " not", phrase)
    phrase = re.sub(r"\'t", " not", phrase)
```

```
In [15]: | #remove words with numbers python: https://stackoverflow.com/a/18082370/4084039
         def rmwrdswithnumbers(sentence):
             sent = re.sub("\S^*\d\S^*", "", sentence).strip()
             return sent
In [16]: #remove spacial character: https://stackoverflow.com/a/5843547/4084039
         def rmspcchar(sentence):
             sent = re.sub('[^A-Za-z0-9]+', '', sentence)
             return sent
In [17]: nltk.download('stopwords')
         stopwords = set(stopwords.words('english'))
         [nltk\_data] Downloading package stopwords to
         [nltk_data] /home/kirankumar_yeddala/nltk_data...
         [nltk data] Package stopwords is already up-to-date!
In [18]: from tqdm import tqdm
         preprocessed reviews = []
         for sentance in tqdm(final['Text'].values):
            sentance = cleanhtml(sentance)
             sentance = decontracted(sentance)
             sentance = rmwrdswithnumbers(sentance)
             sentance = rmspcchar(sentance)
             sentance = ' '.join(e.lower() for e in sentance.split() if e.lower() not in
            preprocessed reviews.append(sentance.strip())
         100%| 126357/126357 [00:14<00:00, 8779.37it/s]
In [19]: preprocessed reviews[100]
Out[19]: 'complaint product sometimes difficult squeeze small amount bottle mixture col
         ored massive amounts baking find sometimes really hard get light color adding
         product small quantities mixture however colors bright pleased find color even
         ly distributed throughout mixture thought product little pricy trying believe
         worth extra money product seems high quality shows cooking whenever want add c
         olor things'
```

3.2 Preprocessing Review Summary

```
In [20]: from tqdm import tqdm
preprocessed_reviews_summary = []
for sentance in tqdm(final['Summary'].values):
    sentance = cleanhtml(sentance)
    sentance = decontracted(sentance)
    sentance = rmwrdswithnumbers(sentance)
    sentance = rmspcchar(sentance)
    sentance = ' '.join(e.lower() for e in sentance.split() if e.lower() not in stopwords)
    preprocessed_reviews_summary.append(sentance.strip())

100%| 126357/126357 [00:02<00:00, 49708.05it/s]

In [21]: preprocessed_reviews_summary[100]

Out[21]: 'happy'</pre>
```

4. Featurization

4.1 Bag of Words

```
In [23]: #********BOW***********
bow_vect = CountVectorizer()
model = bow_vect.fit(x_train)
bow_train = model.transform(x_train)
bow_cv = model.transform(x_cv)
bow_test = model.transform(x_test)
```

4.2 TFIDF

```
In [68]: #******TFIDF*****************
    tfidf_vect = TfidfVectorizer(ngram_range=(1,2), min_df = 10, max_features=5000)
    model_tf = tfidf_vect.fit(x_train)
    tfidf_train = model_tf.transform(x_train)
    tfidf_cv = model_tf.transform(x_cv)
    tfidf_test = model_tf.transform(x_test)
```

4.3 W2V

4.3.1 AVG W2V

```
avg_w2vs = []
        for sent in tqdm(preprocessed_reviews):
            #initializing number of words
            n_{words} = 0
            #initializing vector of size of 50
            sent vec = np.zeros(50)
            for word in sent.split():
               if word in w2v words list:
                   #creating for each word is an vector
                   vec = w2v.wv[word]
                   sent vec += vec
                   n words += 1
            if n words != 0:
               sent vec /= n words
               avg w2vs.append(sent vec)
        print(len(avg w2vs))
        print(len(avg w2vs[0]))
        100%| 87773/87773 [18:38<00:00, 78.46it/s]
        87773
        50
```

4.4 TFIDF W2V

```
tfidf_vect = TfidfVectorizer(ngram_range=(1,2))
        tfidf = tfidf_vect.fit_transform(preprocessed_reviews)
        features = tfidf_vect.get_feature_names()
        tfidf w2vs = []
        row = 0
        for sent in tqdm(preprocessed_reviews):
            sent vec = np.zeros(50)
            tfidf_sum = 0
            for word in sent.split():
                if(word in w2v_words and word in features):
                   vec = w2v.wv[word]
                   tfidf value = tfidf[row, features.index(word)]
                   sent vec += (vec * tfidf value)
                   tfidf sum += tfidf value
            if(tfidf sum != 0):
                sent vec /= tfidf sum
                tfidf w2vs.append(sent vec)
            row += 1
```

100%| 87773/87773 [2:26:44<00:00, 9.97it/s]

```
In [25]: | # Here we are trying to find AUC score for each alpha using cross validation dat
         a. We are using simple cross
         # validation data
         def Multi_NB_CV(x_cv, x_test, y_cv, y_test):
             cv auc = []
             alpha = [10000, 1000, 100, 10, 1, 0.1, 0.01, 0.001, 0.0001]
             for i in alpha:
                 tbs = TimeSeriesSplit(n_splits=10)
                 nb = MultinomialNB(alpha = i)
                 nb.fit(x_cv, y_cv)
                 pred_cv = nb.predict_proba(x_cv)[:,1]
                 cv_auc.append(roc_auc_score(y_cv,pred_cv))
                 print("for alpha = {0} the roc_auc_score is {1}". format(i, roc_auc_scor
         e(y_cv,pred_cv)))
             optimal_alpha_cv = alpha[cv_auc.index(max(cv_auc))]
             return cv_auc, optimal_alpha_cv
```

```
In [26]: # Here we are trying to find AUC score for each alpha using Train data. We are u
         sing simple cross
         #- validation data
         def Multi NB train(x train, x test, y train, y test):
             train auc = []
             alpha = [10000, 1000, 100, 10, 1, 0.1, 0.01, 0.001, 0.0001]
             for i in alpha:
                 tbs = TimeSeriesSplit(n_splits=10)
                 nb = MultinomialNB(alpha = i)
                 nb.fit(x_train, y_train)
                 pred train = nb.predict proba(x train)[:,1]
                 train auc.append(roc auc score(y train,pred train))
                 print("for alpha = {0} the roc auc score is {1}". format(i, roc auc scor
         e(y_train,pred_train)))
             optimal_alpha_train = alpha[train auc.index(max(train auc))]
             return train auc, optimal alpha train, nb
In [27]: # Plotting the AUC score
         def ploting_train_cv(alpha, train_auc, cv_auc):
             plt.plot(alpha, cv_auc, label = 'cv_auc')
             plt.plot(alpha, train_auc, label = 'train_auc')
             plt.xlabel('hyperparameter')
             plt.ylabel('area under AUC curve')
             plt.title("Train AUC vs Validate AUC")
             plt.grid()
             plt.legend()
             plt.show()
In [87]: # Function for ROC Curve
         from sklearn.metrics import roc_curve
         def roc_cur(x_train, x_test, y_train, y_test, optimal_alpha):
             model = MultinomialNB(alpha = optimal alpha)
             model.fit(x train, y train)
             pred = model.predict(x test)
             pred_score = model.predict_proba(x_test)[:,1]
             print('For alpha {0} the area under AUC curve is = {1}\n'.format(optimal alp
         ha, roc_auc_score(y_test, pred_score)))
             fpr, tpr, threshold = metrics.roc_curve(y_test, pred_score)
             roc_auc = metrics.auc(fpr, tpr)
             plt.title("ROC Curve")
             plt.plot(fpr, tpr, 'b', label = 'AUC %0.2f' % roc_auc)
             plt.legend(loc = 'lower right')
             plt.plot([0, 1], [0, 1], 'r--')
             plt.xlim([0, 1])
             plt.ylim([0, 1])
             plt.ylabel('True Positive Rate')
             plt.xlabel('False Positive Rate')
             plt.show()
             return pred
In [29]: # Function for calculating different metrics
         def metrics(gsearch cv, std test, y test, x train, y train):
             best estimator = gsearch cv.best estimator
             best estimator.fit(x train, y train)
             y pred = best estimator.predict(std test)
             print("Accuracy on test data:", round(accuracy score(y test, y pred) * 100 ,
         2))
             print("Precision on test data:", round(precision score(y test, y pred) * 100
         , 2))
             print("Recall on test data:", round(recall_score(y_test, y_pred) * 100 , 2))
             print("F1_score on test data:", round(f1_score(y_test, y_pred) * 100,2))
             return y pred
```

```
In [30]: # Function for printing the confusion matrix

def confus_mat(y_test, pred):
    co_mt = confusion_matrix(y_test,pred)
    class_label = ['negative', 'positive']
    df_conf_matrix = pd.DataFrame(co_mt, index=class_label, columns=class_label)
    sns.heatmap(df_conf_matrix, annot=True, fmt='d')
    plt.title("Confusion Matrix")
    plt.xlabel("Predicted")
    plt.ylabel("Actual")
    plt.show()
    print('-'*120)
    print("Accuracy on test data:", round(accuracy_score(y_test, pred) * 100 , 2
))
    print("Precision on test data:", round(precision_score(y_test, pred) * 100 , 2))
    print("Recall on test data:", round(frecall_score(y_test, pred) * 100 , 2))
    print("Fl_score on test data:", round(fl_score(y_test, pred) * 100 , 2))
```

5. Applying Naive Bayes

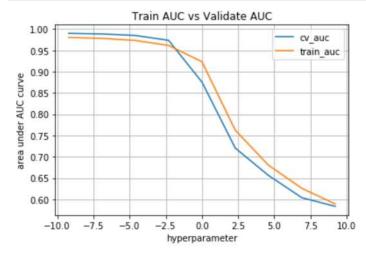
5.1 Applying Multinomial NB on Bow

```
In [31]: | # Normalizing the data
         from sklearn import preprocessing
         bow_train = preprocessing.normalize(bow_train)
         bow_cv = preprocessing.normalize(bow_cv)
         bow test = preprocessing.normalize(bow test)
In [56]: # Calling and Fitting the model
         train_auc, optimal_alpha_train, nb = Multi_NB_train(bow_train, bow_test, y_train
         , y_test)
         optimal_alpha_train
         for alpha = 10000 the roc auc score is 0.5890951916597568
         for alpha = 1000 the roc_auc_score is 0.6258857430229732
         for alpha = 100 the roc auc score is 0.6800231466949458
         for alpha = 10 the roc auc score is 0.762482471412153
         for alpha = 1 the roc_auc_score is 0.9233197889736758
         for alpha = 0.1 the roc_auc_score is 0.9616149385850794
         for alpha = 0.01 the roc_auc_score is 0.9733195484446174
         for alpha = 0.001 the roc auc score is 0.978180940394706
         for alpha = 0.0001 the roc auc score is 0.9803465931079426
Out[56]: 0.0001
In [57]: cv auc, optimal alpha cv = Multi NB CV(bow cv, bow test, y cv, y test)
         optimal_alpha_cv
         for alpha = 10000 the roc_auc_score is 0.5836209590465546
         for alpha = 1000 the roc_auc_score is 0.6038125736443832
         for alpha = 100 the roc_auc_score is 0.6559896115113838
         for alpha = 10 the roc_auc_score is 0.7204371858799408
         for alpha = 1 the roc_auc_score is 0.8756584368423358
         for alpha = 0.1 the roc_auc_score is 0.9735923051605249
         for alpha = 0.01 the roc_auc_score is 0.9850165294659033
         for alpha = 0.001 the roc_auc_score is 0.9885926697815247
         for alpha = 0.0001 the roc_auc_score is 0.9899787655308941
Out[57]: 0.0001
```

Observation

From the above we can see that as the alpha value decreases AUC score increase which means the it is simple model. For alpha = 0.0001 AUC score is higher then we will consider it optimal alpha.

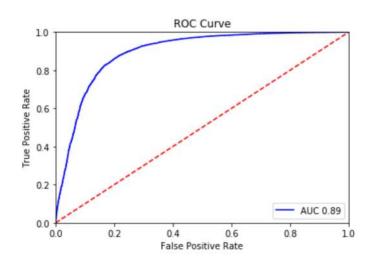
```
In [58]: # Plotting on Train AUC and Validate AUC
    alpha = [10000,1000,100,10,1,0.1,0.01,0.001,0.0001]
    alpha = np.log(np.array(alpha))
    ploting_train_cv(alpha, train_auc, cv_auc)
```



From the above graphy we can see that both train AUC curve and Validate AUC curve decreases with the increase in alpha sharply which means as alpha increases the model becomes dumb model

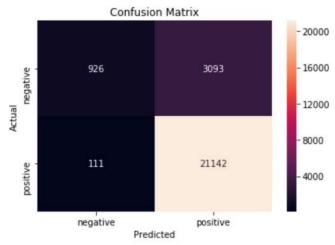
```
In [89]: # ROC Curve
pred = roc_cur(bow_train, bow_test, y_train, y_test, optimal_alpha_cv)
```

For alpha 0.0001 the area under AUC curve is = 0.8942846023804469



From the above we can area under AUC Curve is above the diagnol line then we can say that my model is better than simple linear model

```
In [90]: # Confusion Matrix
confus_mat(y_test, pred)
```



Accuracy on test data: 87.32 Precision on test data: 87.24 Recall on test data: 99.48 F1 score on test data: 92.96

br
like
taste
product
would
one
coffee
good
flavor

Top 10 negative important features

man 10 positive important features

```
Top 10 positive important features
```

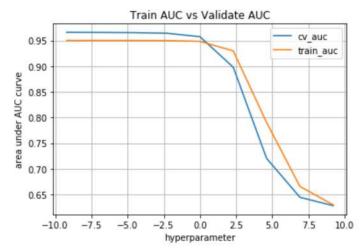
br
great
good
like
love
coffee
one
taste
product
flavor

5.2 Tfidf on Multinomial NB

```
In [69]: # Data Normalisation
         tfidf train = preprocessing.normalize(tfidf train)
         tfidf cv = preprocessing.normalize(tfidf cv)
         tfidf test = preprocessing.normalize(tfidf test)
In [70]: # Calling and fitting the model
         train auc, optimal alpha train, nb = Multi NB train(tfidf train, tfidf test, y t
         rain, y_test)
         optimal_alpha_train
         for alpha = 10000 the roc auc score is 0.6296712542528822
         for alpha = 1000 the roc auc score is 0.6656828421839014
         for alpha = 100 the roc auc score is 0.7903058848603677
         for alpha = 10 the roc auc score is 0.9297995331893694
         for alpha = 1 the roc auc score is 0.9483477156013098
         for alpha = 0.1 the roc auc score is 0.9499045157218561
         for alpha = 0.01 the roc auc score is 0.9500742431282064
         for alpha = 0.001 the roc_auc_score is 0.950111611502867
         for alpha = 0.0001 the roc_auc_score is 0.9501260458636621
Out[70]: 0.0001
In [71]: cv auc, optimal alpha cv = Multi NB CV(tfidf cv, tfidf test, y cv, y test)
         optimal alpha cv
         for alpha = 10000 the roc_auc_score is 0.6281573956131556
         for alpha = 1000 the roc_auc_score is 0.6445344302481425
         for alpha = 100 the roc_auc_score is 0.7204118214473976
         for alpha = 10 the roc_auc_score is 0.897906586115505
         for alpha = 1 the roc_auc_score is 0.9574695673122581
         for alpha = 0.1 the roc auc score is 0.9642367676219629
         for alpha = 0.01 the roc auc score is 0.9653462314366171
         for alpha = 0.001 the roc auc score is 0.9657289931903663
         for alpha = 0.0001 the roc auc score is 0.965919616746104
Out[71]: 0.0001
```

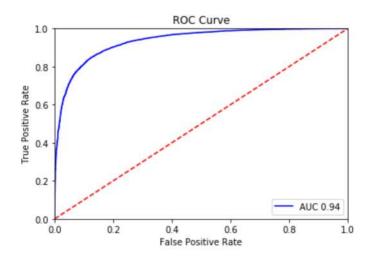
From the above we can see that for alpha > 10 my auc score is constant for decrease in aplha value. For the best alpha value is 0.1 So we will consider alpha = 0.1 as our model.

```
In [72]: # Plotting Train AUC and Validate AUC
    alpha = [10000,1000,100,10,1,0.1,0.01,0.001,0.0001]
    alpha = np.log(np.array(alpha))
    ploting_train_cv(alpha, train_auc, cv_auc)
```



```
In [86]: # ROC curve on Train and Test Data
pred = roc_cur(tfidf_train, tfidf_test, y_train, y_test, optimal_alpha_cv)
```

For alpha 0.0001 the area under AUC curve is = 0.9354631865738856



From the above we can area under AUC Curve is above the diagnol line then we can say that my model is better than simple linear model

```
In [91]: # Confusion Matrix
          confus_mat(y_test, pred)
                         Confusion Matrix
                                                    - 20000
                                                     16000
                      926
                                       3093
                                                    - 12000
                                                     8000
                                      21142
                                                     4000
                    negative
                                      positive
                            Predicted
         Accuracy on test data: 87.32
         Precision on test data: 87.24
         Recall on test data: 99.48
         F1 score on test data: 92.96
In [75]: | #Top positive and Negative features
          features = tfidf_vect.get_feature_names()
         class_labels = nb.classes_
          top_negative = sorted(zip(nb.feature_log_prob_[0], features), reverse=True)[0:10]
          top_positive = sorted(zip(nb.feature_log_prob_[1], features),reverse=True)[0:10]
         print("Top 10 negative important features")
          for coef, feat in top_negative:
```

```
print(feat)
print("----")
print("Top 10 positive important features")
for coef, feat in top_positive:
  print(feat)
```

```
Top 10 negative important features
like
product
taste
would
coffee
br br
one
flavor
_____
Top 10 positive important features
great
good
coffee
like
love
tea
br br
product
one
```

6. Conclusion

30-01-2019, 23:17 18 of 19

Out[92]:

	Vectorizer	Best alpha	ROC_AUC	Accuracy	Precision	Recall	F1_score
0	BOW	0.0001	89.42	87.32	87.24	99.48	92.96
1	TFIDF	0.0001	93.54	87.54	88.43	99.39	93.59

```
In [ ]:
```