# **Amazon Fine Food Reviews Analysis**

Data Source: <a href="https://www.kaggle.com/snap/amazon-fine-food-reviews">https://www.kaggle.com/snap/amazon-fine-food-reviews</a>

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. ld
- 2. Productld unique identifier for the product
- 3. Userld 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 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

- 1. .csv file
- 2. 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 wil be set to "positive". Otherwise, it will be set to "negative".

```
In [1]: %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 tadm import tadm
import os
```

```
In [2]: # using SQLite Table to read data.
    con = sqlite3.connect('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 50
0000 data 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 Sco
    re != 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 sc
ore<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)</pre>
```

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

#### Out[2]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfulnes
0	1	B001E4KFG0	A3SGXH7AUHU8GW	delmartian	1	1
1	2	B00813GRG4	A1D87F6ZCVE5NK	dll pa	0	0

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfulnes
2	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1	1

```
In [0]: display = pd.read_sql_query("""
    SELECT UserId, ProductId, ProfileName, Time, Score, Text, COUNT(*)
    FROM Reviews
    GROUP BY UserId
    HAVING COUNT(*)>1
    """, con)
```

In [0]: print(display.shape)
display.head()

(80668, 7)

Out[0]:

	Userld	ProductId	ProfileName	Time	Score	Text	COU
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

	Userld	ProductId	ProfileName	Time	Score	Text	COU
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 [0]: display[display['UserId']=='AZY10LLTJ71NX']

Out[0]:

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

In [0]: display['COUNT(\*)'].sum()

Out[0]: 393063

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

#### Out[0]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfuln
0	78445	B000HDL1RQ	AR5J8UI46CURR	Geetha Krishnan	2	2
1	138317	B000HDOPYC	AR5J8UI46CURR	Geetha Krishnan	2	2

		ld	ProductId	UserId	ProfileName	HelpfulnessNumerator	Helpfuln
2	2	138277	B000HDOPYM	AR5J8UI46CURR	Geetha Krishnan	2	2
3	3	73791	B000HDOPZG	AR5J8UI46CURR	Geetha Krishnan	2	2
4	4	155049	B000PAQ75C	AR5J8UI46CURR	Geetha Krishnan	2	2

As it can be seen above that same user has multiple reviews with 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 ProductId 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 delelte 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 [3]: #Sorting data according to ProductId in ascending order
    sorted_data=filtered_data.sort_values('ProductId', axis=0, ascending=Tr
    ue, inplace=False, kind='quicksort', na_position='last')
```

```
In [4]: #Deduplication of entries
    final=sorted_data.drop_duplicates(subset={"UserId","ProfileName","Time"
        ,"Text"}, keep='first', inplace=False)
    final.shape
```

Out[4]: (364173, 10)

```
In [5]: #Checking to see how much % of data still remains
  (final['Id'].size*1.0)/(filtered_data['Id'].size*1.0)*100
```

Out[5]: 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

		ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfuln	
	0	64422	B000MIDROQ	A161DK06JJMCYF	J. E. Stephens "Jeanne"	3	1	
	1	44737	B001EQ55RW	A2V0I904FH7ABY	Ram	3	2	
	4						<b>&gt;</b>	
In [6]:	fi	nal=fi	inal[final.He	elpfulnessNumera	tor<=final.	HelpfulnessDenomina	tor]	
In [7]:	е	ntries		e next phase of	preprocessi	ng lets see the num	ber of	
	<pre>#How many positive and negative reviews are present in our dataset? final['Score'].value_counts()</pre>							
	(3	64171,	10)					
Out[7]:	1 0 Na	57	7061 7110 core, 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:-

- 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 [0]: # 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)
```

Why is this \$[...] when the same product is available for \$[...] here?<br/>br />http://www.amazon.com/VICTOR-FLY-MAGNET-BATT-REFTLL/dn/B00004RBDY<

br /><br/>
br />The Victor M380 and M502 traps are unreal, of course -- total fly genocide. Pretty stinky, but only right nearby.

\_\_\_\_\_

I recently tried this flavor/brand and was surprised at how delicious t hese chips are. The best thing was that there were a lot of "brown" chips in the bsg (my favorite), so I bought some more through amazon and shared with family and friends. I am a little disappointed that there are not, so far, very many brown chips in these bags, but the flavor is still very good. I like them better than the yogurt and green onion flavor because they do not seem to be as salty, and the onion flavor is better. If you haven't eaten Kettle chips before, I recommend that you try a bag before buying bulk. They are thicker and crunchier than Lays but just as fresh out of the bag.

\_\_\_\_\_

Wow. So far, two two-star reviews. One obviously had no idea what the y were ordering; the other wants crispy cookies. Hey, I'm sorry; but t hese reviews do nobody any good beyond reminding us to look before ord ering.<br/>
/>t /><br/>
/>These are chocolate-oatmeal cookies. If you don't li ke that combination, don't order this type of cookie. I find the combo quite nice, really. The oatmeal sort of "calms" the rich chocolate fla vor and gives the cookie sort of a coconut-type consistency. Now let's also remember that tastes differ; so, I've given my opinion.<br /><br / >Then, these are soft, chewy cookies -- as advertised. They are not "c rispy" cookies, or the blurb would say "crispy," rather than "chewy." I happen to like raw cookie dough; however, I don't see where these tas te like raw cookie dough. Both are soft, however, so is this the confu sion? And, yes, they stick together. Soft cookies tend to do that. T hev aren't individually wrapped, which would add to the cost. Oh yeah, chocolate chip cookies tend to be somewhat sweet.<br /><br />So, if you want something hard and crisp, I suggest Nabiso's Ginger Snaps. If you want a cookie that's soft, chewy and tastes like a combination of choco late and oatmeal, give these a try. I'm here to place my second order.

\_\_\_\_\_

love to order my coffee on amazon. easy and shows up quickly.<br/>This k cup is great coffee. dcaf is very good as well

\_\_\_\_\_\_

In [0]: # remove urls from text python: https://stackoverflow.com/a/40823105/40
84039

```
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)
```

Why is this \$[...] when the same product is available for \$[...] here?<br/>br /> <br/>
br /> <br/>
The Victor M380 and M502 traps are unreal, of course -- total fly genocide. Pretty stinky, but only right nearby.

```
In [0]: # https://stackoverflow.com/questions/16206380/python-beautifulsoup-how
        -to-remove-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)
```

Why is this \$[...] when the same product is available for \$[...] here? />The Victor M380 and M502 traps are unreal, of course -- total fly gen ocide. Pretty stinky, but only right nearby.

\_\_\_\_\_\_

I recently tried this flavor/brand and was surprised at how delicious t hese chips are. The best thing was that there were a lot of "brown" ch

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\_\_\_\_\_

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love to order my coffee on amazon. easy and shows up quickly. This k cu p is great coffee. dcaf is very good as well

```
In [0]: # 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"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"\'ll", " will", phrase)
phrase = re.sub(r"\'t", " not", phrase)
phrase = re.sub(r"\'ve", " have", phrase)
phrase = re.sub(r"\'ve", " am", phrase)
return phrase
```

```
In [0]: sent_1500 = decontracted(sent_1500)
    print(sent_1500)
    print("="*50)
```

Wow. So far, two two-star reviews. One obviously had no idea what the y were ordering; the other wants crispy cookies. Hey, I am sorry; but these reviews do nobody any good beyond reminding us to look before or dering.<br /><br />These are chocolate-oatmeal cookies. If you do not like that combination, do not order this type of cookie. I find the co mbo quite nice, really. The oatmeal sort of "calms" the rich chocolate flavor and gives the cookie sort of a coconut-type consistency. Now le t is also remember that tastes differ; so, I have given my opinion.<br/> /><br />Then, these are soft, chewy cookies -- as advertised. They are not "crispy" cookies, or the blurb would say "crispy," rather than "che wy." I happen to like raw cookie dough; however. I do not see where th ese taste like raw cookie dough. Both are soft, however, so is this th e confusion? And, yes, they stick together. Soft cookies tend to do t hat. They are not individually wrapped, which would add to the cost. Oh yeah, chocolate chip cookies tend to be somewhat sweet.<br/>>br/>S o, if you want something hard and crisp, I suggest Nabiso is Ginger Sna ps. If you want a cookie that is soft, chewy and tastes like a combina tion of chocolate and oatmeal, give these a try. I am here to place my second order.

```
In [0]: #remove words with numbers python: https://stackoverflow.com/a/1808237
0/4084039
```

```
sent_0 = re.sub("\S*\d\S*", "", sent_0).strip()
print(sent_0)
```

Why is this \$[...] when the same product is available for \$[...] here?<br/>br /> <br/>br /> The Victor and traps are unreal, of course -- total fly genocide. Pretty stinky, but only right nearby.

```
In [0]: #remove spacial character: https://stackoverflow.com/a/5843547/4084039
sent_1500 = re.sub('[^A-Za-z0-9]+', ' ', sent_1500)
print(sent_1500)
```

Wow So far two two star reviews One obviously had no idea what they wer e ordering the other wants crispy cookies Hey I am sorry but these revi ews do nobody any good beyond reminding us to look before ordering br b r These are chocolate oatmeal cookies If you do not like that combinati on do not order this type of cookie I find the combo quite nice really The oatmeal sort of calms the rich chocolate flavor and gives the cooki e sort of a coconut type consistency Now let is also remember that tast es differ so I have given my opinion br br Then these are soft chewy co okies as advertised They are not crispy cookies or the blurb would say crispy rather than chewy I happen to like raw cookie dough however I do not see where these taste like raw cookie dough Both are soft however s o is this the confusion And yes they stick together Soft cookies tend t o do that They are not individually wrapped which would add to the cost Oh yeah chocolate chip cookies tend to be somewhat sweet br br So if yo u want something hard and crisp I suggest Nabiso is Ginger Snaps If you want a cookie that is soft chewy and tastes like a combination of choco late and oatmeal give these a try I am here to place my second order

```
In [0]: # https://gist.github.com/sebleier/554280
    # we are removing the words from the stop words list: 'no', 'nor', 'no
    t'
    # <br /><br /> ==> after the above steps, we are getting "br br"
    # we are including them into stop words list
    # instead of <br /> if we have <br/> these tags would have revmoved in
    the 1st step

stopwords= set(['br', 'the', 'i', 'me', 'my', 'myself', 'we', 'our', 'o
    urs', 'ourselves', 'you', "you're", "you've",\
```

```
"you'll", "you'd", 'your', 'yours', 'yourself', 'yourselve
s', 'he', 'him', 'his', 'himself', \
            'she', "she's", 'her', 'hers', 'herself', 'it', "it's", 'it
s', 'itself', 'they', 'them', 'their',\
            'theirs', 'themselves', 'what', 'which', 'who', 'whom', 'th
is', 'that', "that'll", 'these', 'those', \
            'am', 'is', 'are', 'was', 'were', 'be', 'been', 'being', 'h
ave', 'has', 'had', 'having', 'do', 'does', \
            'did', 'doing', 'a', 'an', 'the', 'and', 'but', 'if', 'or',
'because', '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', 'h
ow', 'all', 'any', 'both', 'each', 'few', 'more',\
            'most', 'other', 'some', 'such', 'only', 'own', 'same', 's
o', 'than', 'too', 'very', \
            's', 't', 'can', 'will', 'just', 'don', "don't", 'should',
"should'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', "is
n't", 'ma', 'mightn', "mightn't", 'mustn',\
            "mustn't", 'needn', "needn't", 'shan', "shan't", 'shouldn',
"shouldn't", 'wasn', "wasn't", 'weren', "weren't", \
            'won', "won't", 'wouldn', "wouldn't"])
```

```
In [0]: # Combining all the above stundents
    from tqdm import tqdm
    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
```

#### In [0]: preprocessed\_reviews[1500]

Out[0]: 'wow far two two star reviews one obviously no idea ordering wants cris py cookies hey sorry reviews nobody good beyond reminding us look order ing chocolate oatmeal cookies not like combination not order type cookie e find combo quite nice really oatmeal sort calms rich chocolate flavor gives cookie sort coconut type consistency let also remember tastes differ given opinion soft chewy cookies advertised not crispy cookies blur b would say crispy rather chewy happen like raw cookie dough however not see taste like raw cookie dough soft however confusion yes stick toge ther soft cookies tend not individually wrapped would add cost oh yeah chocolate chip cookies tend somewhat sweet want something hard crisp su ggest nabiso ginger snaps want cookie soft chewy tastes like combination chocolate oatmeal give try place second order'

## [3.2] Preprocessing Review Summary

```
In [30]: ## Similartly you can do preprocessing for review summary also.
#set of stopwords in English
from nltk.corpus import stopwords
stop = set(stopwords.words('english'))
import re

words_to_keep = set(('not'))
stop -= words_to_keep

#initialising the snowball stemmer
sno = nltk.stem.SnowballStemmer('english')

#function to clean the word of any html-tags
def cleanhtml(sentence):
```

```
cleanr = re.compile('<.*?>')
    cleantext = re.sub(cleanr, ' ', sentence)
    return cleantext
#function to clean the word of any punctuation or special characters
def cleanpunc(sentence):
    cleaned = re.sub(r'[?|!|\'|"|#]',r'',sentence)
    cleaned = re.sub(r'[.|,|)|(||/|,r'|,cleaned)
    return cleaned
#Code for removing HTML tags , punctuations . Code for removing stopwor
ds . Code for checking if word is not alphanumeric and
# also greater than 2 . Code for stemming and also to convert them to l
owercase letters
i = 0
str1=' '
final string=[]
all positive words=[] # store words from +ve reviews here
all negative words=[] # store words from -ve reviews here.
S=' -
for sent in final['Summary'].values:
    filtered sentence=[]
    #print(sent);
    sent=cleanhtml(sent) # remove HTMl tags
    for w in sent.split():
        for cleaned words in cleanpunc(w).split():
            if((cleaned words.isalpha()) & (len(cleaned words)>2)):
                if(cleaned words.lower() not in stop):
                    s=(sno.stem(cleaned words.lower())).encode('utf8')
                    filtered sentence.append(s)
                    if (final['Score'].values)[i] == 1:
                        all positive words.append(s) #list of all words
used to describe positive reviews
                    if(final['Score'].values)[i] == 0:
                        all negative words.append(s) #list of all words
used to describe negative reviews reviews
                else:
                    continue
            else:
                continue
```

```
strl = b" ".join(filtered_sentence) #final string of cleaned words
final_string.append(strl)
i+=1
```

In [31]: #adding a column of CleanedText which displays the data after pre-proce
 ssing of the review
 final['CleanedSummary']=final\_string
 final['CleanedSummary']=final['CleanedSummary'].str.decode("utf-8")
 #below the processed review can be seen in the CleanedText Column
 print('Shape of final', final.shape)
 final.head()

Shape of final (364171, 13)

#### Out[31]:

	index	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator
O	138706	150524	0006641040	ACITT7DI6IDDL	shari zychinski	0
1	138688	150506	0006641040	A2IW4PEEKO2R0U	Tracy	1

	index	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator
2	138689	150507	0006641040	A1S4A3IQ2MU7V4	sally sue "sally sue"	1
3	138690	150508	0006641040	AZGXZ2UUK6X	Catherine Hallberg " (Kate)"	1
4	138691	150509	0006641040	A3CMRKGE0P909G	Teresa	3

```
In [33]: print(final['CleanedText'][0])
    print("******")
    print(final['CleanedSummary'][0])
    print("****")
    final['CleanedText'][:]=final['CleanedText'][:]+' '+final['CleanedText'][:]+' '+final['CleanedSummary'][:]
    print(final['CleanedText'][0])
```

witti littl book make son laugh loud recit car drive along alway sing r

efrain hes learn whale india droop love new word book introduc silli cl assic book will bet son still abl recit memori colleg \*\*\*\*\*\*\*\* everi book educ \*\*\*\*\*

witti littl book make son laugh loud recit car drive along alway sing r efrain hes learn whale india droop love new word book introduc silli cl assic book will bet son still abl recit memori colleg witti littl book make son laugh loud recit car drive along alway sing refrain hes learn whale india droop love new word book introduc silli classic book will b et son still abl recit memori colleg everi book educ everi book educ

## Note:

preprocessing is done in previous assignments and stored in final.sqlite

```
In [2]: con = sqlite3.connect('final.sqlite')
final = pd.read_sql_query(""" SELECT * FROM Reviews WHERE Score != 3
```

""", con) final.head()

#### Out[2]:

	index	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator
0	138706	150524	0006641040	ACITT7DI6IDDL	shari zychinski	0
1	138688	150506	0006641040	A2IW4PEEKO2R0U	Tracy	1
2	138689	150507	0006641040	A1S4A3IQ2MU7V4	sally sue "sally sue"	1

	index	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator
3	138690	150508	0006641040	AZGXZ2UUK6X	Catherine Hallberg " (Kate)"	1
4	138691	150509	0006641040	A3CMRKGE0P909G	Teresa	3

```
In [3]: final.shape
Out[3]: (364171, 12)

In [4]: from sklearn.model_selection import train_test_split
    ##Sorting data according to Time in ascending order for Time Based Spli
    tting
    time_sorted_data = final.sort_values('Time', axis=0, ascending=True, in
    place=False, kind='quicksort', na_position='last')

X = time_sorted_data['CleanedText'].values
Y = time_sorted_data['Score']

# split the data set into train and test
```

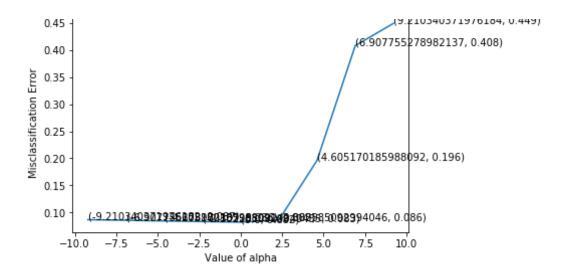
```
X train, X test, Y train, Y test = train test split(X,Y, test size=0.3,
          shuffle=False)
In [5]: print(X train.shape, Y train.shape)
         print(X test.shape, Y test.shape)
         (254919,) (254919,)
         (109252,) (109252,)
         Applying Multinomial Naive Bayes
         [5.1] Applying Naive Bayes on BOW, SET 1
In [6]: #BoW
         count vect = CountVectorizer(min df = 50)
         X train vec = count vect.fit transform(X train)
         X test vec = count vect.transform(X test)
         print("the type of count vectorizer :",type(X train vec))
         print("the shape of out text BOW vectorizer : ",X train vec.get shape
         print("the number of unique words :", X train vec.get shape()[1])
         the type of count vectorizer : <class 'scipy.sparse.csr.csr matrix'>
         the shape of out text BOW vectorizer: (254919, 6050)
         the number of unique words : 6050
In [10]: from sklearn.naive bayes import MultinomialNB
         from sklearn.model selection import cross val score
         # Creating alpha values in the range from 10^-4 to 10^4
         neighbors = [10**-4,10**-3,10**-2,10**-1,10**0,10**1,10**2,10**3.10**4]
         # empty list that will hold cv scores
         cv scores = []
         training scores =[]
```

# perform 10-fold cross validation

```
for k in neighbors:
    nb = MultinomialNB(alpha=k)
    nb.fit(X train vec, Y train)
    #print(nb.predict(X test[2:39]))
    scores = cross val score(nb, X test vec, Y test, cv=10, scoring='ro
c auc')
    scores training = nb.fit(X train vec, Y train).score(X train vec, Y
train)
    cv scores.append(scores.mean())
    training scores.append(scores training)
    #print((nb))
MSE = [1 - x \text{ for } x \text{ in } cv \text{ scores}]
#determining best k
optimal alpha = neighbors[MSE.index(min(MSE))]
print('\nThe optimal value of alpha is %.8f.' % optimal alpha)
plt.plot(np.log(neighbors), MSE)
for xy in zip(np.log(neighbors), np.round(MSE,3)):
    plt.annotate('(%s, %s)' % xy, xy=xy, textcoords='data')
plt.xlabel('Value of alpha')
plt.ylabel('Misclassification Error')
plt.show()
print("the misclassification error for each k value is : ", np.round(MS
E,3))
plt.plot(np.log(neighbors), cv scores)
plt.plot(np.log(neighbors), training scores)
plt.xlabel('alpha')
plt.ylabel('score')
# determining best value of alpha
#best alpha = neighbors[cv scores.index(max(cv scores))]
#print('\nThe optimal value of alpha is %.3f.' % best alpha)
```

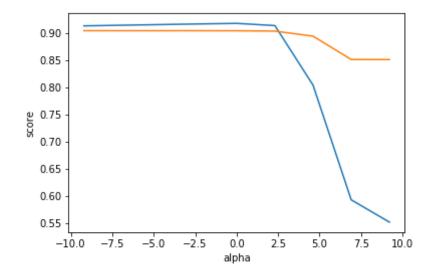
The optimal value of alpha is 1.00000000.

(0.10240271075194.0.440)



the misclassification error for each k value is : [0.087 0.086 0.085 0.083 0.082 0.086 0.196 0.408 0.449]

Out[10]: Text(0,0.5,'score')

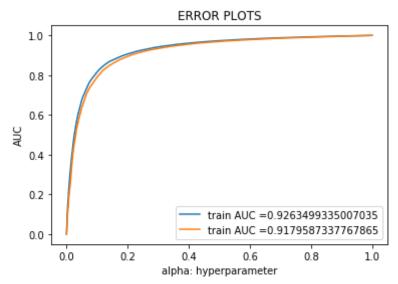


```
from sklearn.metrics import accuracy score
         mnb1 = MultinomialNB(alpha = optimal alpha)
         # fitting the model
         mnb1.fit(X train vec, Y train)
         # predict the response
         predictions= mnb1.predict(X test vec)
         predictions1= mnb1.predict(X train vec)
         # evaluate accuracy
         acc = accuracy score(Y test, predictions) * 100
         print('\nThe Test Accuracy of the Multinomial naive Bayes classifier fo
         r alpha = %.3f is %f%%' % (optimal alpha, acc))
         # Variables that will be used for making table in Conclusion part of t
         his assignment
         bow mnb alpha = optimal alpha
         bow mnb train acc = max(cv scores)*100
         bow mnb test acc = acc
         The Test Accuracy of the Multinomial naive Bayes classifier for alpha =
         1.000 is 89.169992%
In [12]: # https://scikit-learn.org/stable/modules/generated/sklearn.metrics.roc
          curve.html#sklearn.metrics.roc curve
         from sklearn.metrics import roc curve, auc
         # roc auc score(y true, y score) the 2nd parameter should be probabilit
         v estimates of the positive class
         # not the predicted outputs
         train fpr, train tpr, thresholds = roc curve(Y train, mnb1.predict prob
         a(X train vec)[:,1])
         test_fpr, test_tpr, thresholds = roc_curve(Y_test, mnb1.predict proba(X)
         test vec)[:,1])
         plt.plot(train fpr, train tpr, label="train AUC ="+str(auc(train fpr, t
         rain tpr)))
```

```
plt.plot(test_fpr, test_tpr, label="train AUC ="+str(auc(test_fpr, test _tpr)))
plt.legend()
plt.xlabel("alpha: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()

print("="*100)

from sklearn.metrics import confusion_matrix
print("Train confusion matrix")
print(confusion_matrix(Y_train, mnbl.predict(X_train_vec)))
print("Test confusion_matrix")
print(confusion_matrix(Y_test, mnbl.predict(X_test_vec)))
```



\_\_\_\_\_\_

#### [5.1.1] Top 10 important features of negative class from SET 1

```
In [13]: # Please write all the code with proper documentation
         # top 10 features
         import operator
         from nltk.probability import FreqDist, DictionaryProbDist, ELEProbDist,
          sum logs
         from nltk.classify.api import ClassifierI
         from nltk.classify.naivebayes import NaiveBayesClassifier
         nb = MultinomialNB(alpha=optimal alpha).fit(X train vec, Y train)
         pos imp features = nb.feature log prob [1,:]
         neg imp features = nb.feature log prob [0,:]
         imp features = {}
         feature names= count vect.get_feature_names()
         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.itemget
         ter(1), reverse = True)
         print("\n\nNegative top 10 important features are:")
         for i in range(10):
             print(names diff sorted[i])
         Negative top 10 important features are:
         ('tast', -4.174592588303286)
         ('like', -4.255594226628023)
         ('product', -4.424308118931364)
         ('one', -4.698637689183009)
         ('flavor', -4.7400729276772715)
         ('tri', -4.849917472839326)
         ('would', -4.850507866311542)
         ('qood', -5.007094086504942)
         ('coffe', -5.031362516749736)
```

```
('use', -5.0391770386804815)
```

#### [5.1.2] Top 10 important features of positive class from SET 1

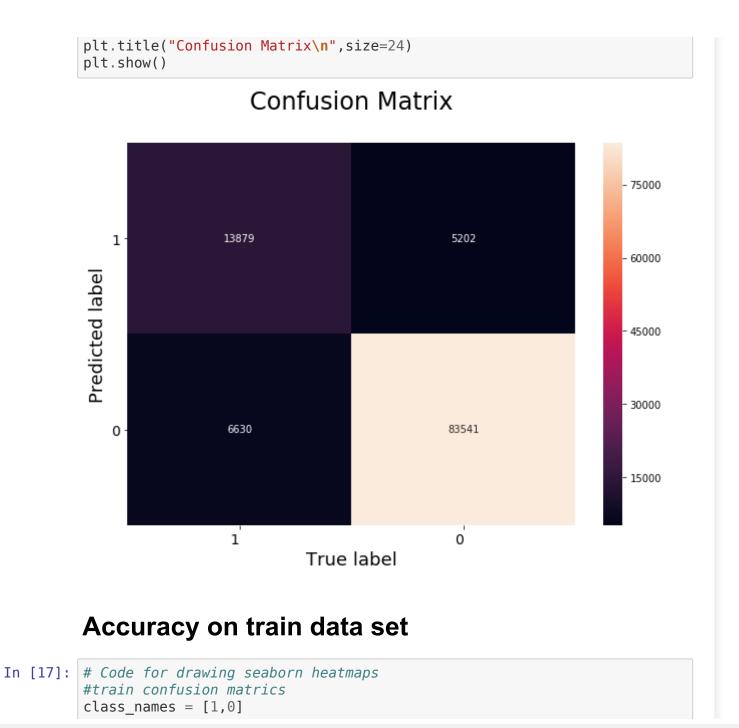
```
In [14]: # Please write all the code with proper documentation
         for i in range(len(feature names)):
             imp features[feature names[i]] = pos imp features[i]
         names diff sorted = sorted(imp features.items(), key = operator.itemget
         ter(1), reverse = True)
         print("Postive top 10 important features are:")
         for i in range(10):
             print(names_diff_sorted[i])
         Postive top 10 important features are:
         ('like', -4.40894771917271)
         ('tast', -4.47930292964824)
         ('qood', -4.614126623248737)
         ('flavor', -4.6357357369625625)
         ('love', -4.66429007687182)
         ('great', -4.684453739591529)
         ('use', -4.706008320452014)
         ('one', -4.762344073425835)
         ('product', -4.848698285348165)
         ('tea', -4.856661564089681)
```

# Accuracy on test data

```
In [15]: # evaluate accuracy
#libraries
from sklearn.metrics import precision_score
from sklearn.metrics import recall_score
from sklearn.metrics import fl_score

acc = accuracy_score(Y_test, predictions) * 100
print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
f%%' % (optimal_alpha, acc))
```

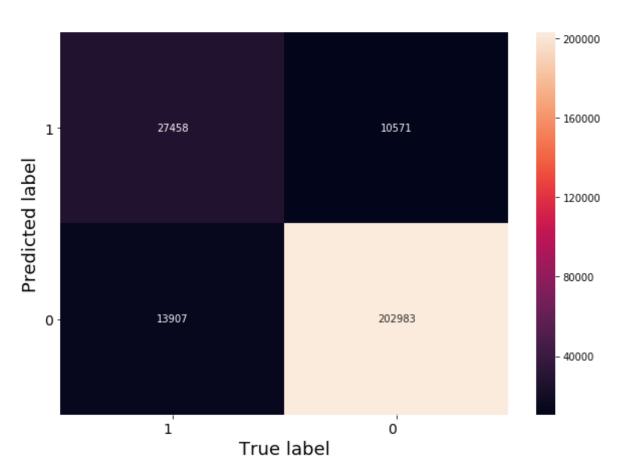
```
# evaluate precision
         acc = precision_score(Y_test, predictions, pos label = 1)
         print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
         f' % (optimal alpha, acc))
         # evaluate recall
         acc = recall score(Y test, predictions, pos label = 1)
         print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
         % (optimal alpha, acc))
         # evaluate f1-score
         acc = f1_score(Y_test, predictions, pos label = 1)
         print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f
         ' % (optimal alpha, acc))
          Accuracy of the Multinomial naive Bayes of alpha = 1.000 is 89.169992%
          Precision of the Multinomial naive Bayes of alpha = 1.000 is 0.941381
          Recall of the Multinomial naive Bayes of alpha = 1.000 is 0.926473
          F1-Score of the Multinomial naive Bayes of alpha = 1.000 is 0.933868
In [16]: # Code for drawing seaborn heatmaps
         #test confusion matrics
         class names = [1,0]
         df heatmap = pd.DataFrame(confusion matrix(Y test, predictions), index=
         class names, columns=class names )
         fig = plt.figure(figsize=(10,7))
         heatmap = sns.heatmap(df heatmap, annot=True, fmt="d")
         # Setting tick labels for heatmap
         heatmap.yaxis.set ticklabels(heatmap.yaxis.get ticklabels(), rotation=0
         , ha='right', fontsize=14)
         heatmap.xaxis.set ticklabels(heatmap.xaxis.get ticklabels(), rotation=0
         , ha='right', fontsize=14)
         plt.ylabel('Predicted label',size=18)
         plt.xlabel('True label', size=18)
```



```
df_heatmap = pd.DataFrame(confusion_matrix(Y_train, mnbl.predict(X_train_vec)), index=class_names, columns=class_names)
fig = plt.figure(figsize=(10,7))
heatmap = sns.heatmap(df_heatmap, annot=True, fmt="d")

# Setting tick labels for heatmap
heatmap.yaxis.set_ticklabels(heatmap.yaxis.get_ticklabels(), rotation=0, ha='right', fontsize=14)
heatmap.xaxis.set_ticklabels(heatmap.xaxis.get_ticklabels(), rotation=0, ha='right', fontsize=14)
plt.ylabel('Predicted label',size=18)
plt.xlabel('True label',size=18)
plt.title("Confusion Matrix\n",size=24)
plt.show()
```

#### Contusion Matrix



```
In [18]: # evaluate accuracy on train data set
    acc = accuracy_score(Y_train,mnbl.predict(X_train_vec) ) * 100
    print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
    f%%' % (optimal_alpha, acc))

# evaluate precision
    acc = precision_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1
    )
    print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
    f' % (optimal_alpha, acc))
```

```
# evaluate recall
acc = recall_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1)
print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
% (optimal_alpha, acc))

# evaluate f1-score
acc = f1_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1)
print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f'
% (optimal_alpha, acc))
```

Accuracy of the Multinomial naive Bayes of alpha = 1.000 is 90.397734%

Precision of the Multinomial naive Bayes of alpha = 1.000 is 0.950500

Recall of the Multinomial naive Bayes of alpha = 1.000 is 0.935880

F1-Score of the Multinomial naive Bayes of alpha = 1.000 is 0.943133

## [5.2] Applying Naive Bayes on TFIDF, SET 2

```
In [19]: # Please write all the code with proper documentation
    tf_idf_vect = TfidfVectorizer(min_df=10)
    X_train_vec = tf_idf_vect.fit_transform(X_train)
    X_test_vec = tf_idf_vect.transform(X_test)
    print("the type of count vectorizer : ",type(X_train_vec))
    print("the shape of out text TFIDF vectorizer : ",X_train_vec.get_shape
    ())
    print("the number of unique words : ", X_train_vec.get_shape()[1])

    the type of count vectorizer : <class 'scipy.sparse.csr.csr_matrix'>
        the shape of out text TFIDF vectorizer : (254919, 12671)
    the number of unique words : 12671

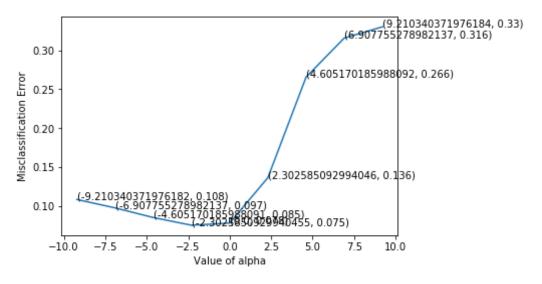
In [20]: from sklearn.naive_bayes import MultinomialNB
    from sklearn.model_selection import cross_val_score

# Creating alpha values in the range from 10^-4 to 10^4
```

```
neighbors = [10**-4,10**-3,10**-2,10**-1,10**0,10**1,10**2,10**3,10**4]
# empty list that will hold cv scores
cv scores = []
training scores =[]
# perform 10-fold cross validation
for k in neighbors:
    nb = MultinomialNB(alpha=k)
    nb.fit(X train vec, Y train)
    #print(nb.predict(X test[2:39]))
    scores = cross val score(nb, X test vec, Y test, cv=10, scoring='ro
c auc')
    scores training = nb.fit(X train vec, Y train).score(X train vec, Y
train)
    cv scores.append(scores.mean())
    training scores.append(scores training)
    #print((nb))
MSE = [1 - x \text{ for } x \text{ in } cv \text{ scores}]
#determining best k
optimal alpha = neighbors[MSE.index(min(MSE))]
print('\nThe optimal value of alpha is %.8f.' % optimal alpha)
plt.plot(np.log(neighbors), MSE)
for xy in zip(np.log(neighbors), np.round(MSE,3)):
    plt.annotate('(%s, %s)' % xy, xy=xy, textcoords='data')
plt.xlabel('Value of alpha')
plt.ylabel('Misclassification Error')
plt.show()
print("the misclassification error for each k value is : ", np.round(MS
E,3))
plt.plot(np.log(neighbors), cv scores, 'r')
plt.plot(np.log(neighbors), training scores, 'b')
plt.xlabel('alpha')
plt.ylabel('score')
# determining best value of alpha
```

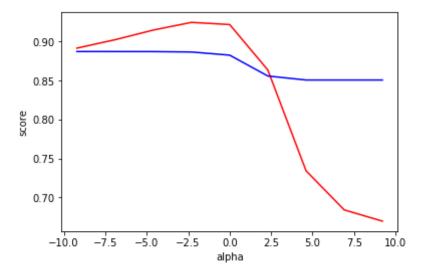
```
#best_alpha = neighbors[cv_scores.index(max(cv_scores))]
#print('\nThe optimal value of alpha is %.3f.' % best_alpha)
```

The optimal value of alpha is 0.10000000.



the misclassification error for each k value is :  $[0.108 \ 0.097 \ 0.085 \ 0.075 \ 0.078 \ 0.136 \ 0.266 \ 0.316 \ 0.33 ]$ 

Out[20]: Text(0,0.5,'score')



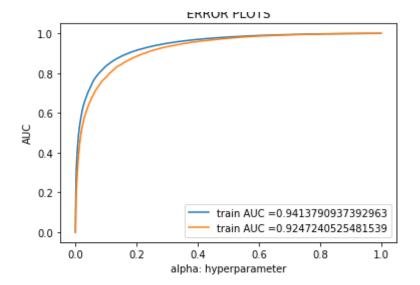
```
ptimal alpha ======
        # instantiate learning model alpha = optimal alpha
        from sklearn.metrics import accuracy score
        mnb1 = MultinomialNB(alpha = optimal alpha)
        # fitting the model
        mnb1.fit(X_train_vec, Y_train)
        # predict the response
        predictions= mnbl.predict(X test vec)
        predictions1= mnb1.predict(X train vec)
        # evaluate accuracy
        acc = accuracy score(Y test, predictions) * 100
        print('\nThe Test Accuracy of the Multinomial naive Bayes classifier fo
        r alpha = %.3f is %f%%' % (optimal alpha, acc))
        # Variables that will be used for making table in Conclusion part of t
        his assignment
        tfidf_mnb_alpha = optimal_alpha
```

```
tfidf_mnb_train_acc = max(cv_scores)*100
tfidf_mnb_test_acc = acc
```

The Test Accuracy of the Multinomial naive Bayes classifier for alpha = 0.100 is 86.599788%

```
In [22]: # https://scikit-learn.org/stable/modules/generated/sklearn.metrics.roc
          curve.html#sklearn.metrics.roc curve
         from sklearn.metrics import roc curve, auc
         # roc auc score(y true, y score) the 2nd parameter should be probabilit
         y estimates of the positive class
         # not the predicted outputs
         train fpr, train tpr, thresholds = roc curve(Y train, mnb1.predict prob
         a(X train vec)[:,1])
         test fpr, test tpr, thresholds = roc curve(Y test, mnb1.predict proba(X
         test vec)[:,1])
         plt.plot(train fpr, train tpr, label="train AUC ="+str(auc(train fpr, t
         rain tpr)))
         plt.plot(test fpr, test tpr, label="train AUC ="+str(auc(test fpr, test
         tpr)))
         plt.legend()
         plt.xlabel("alpha: hyperparameter")
         plt.ylabel("AUC")
         plt.title("ERROR PLOTS")
         plt.show()
         print("="*100)
         from sklearn.metrics import confusion matrix
         print("Train confusion matrix")
         print(confusion matrix(Y train, mnb1.predict(X train vec)))
         print("Test confusion matrix")
         print(confusion matrix(Y test, mnb1.predict(X test vec)))
```

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```
Train confusion matrix
[[ 10192 27837]
  [ 1029 215861]]
Test confusion matrix
[[ 4943 14138]
  [ 502 89669]]
```

#### [5.2.1] Top 10 important features of positive class from SET 2

```
In [23]: # Please write all the code with proper documentation
    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.itemget
        ter(1), reverse = True)
    print("Postive top 10 important features are:")
    for i in range(10):
        print(names_diff_sorted[i])

Postive top 10 important features are:
    ('like', -4.40894771917271)
```

```
('tast', -4.47930292964824)
('good', -4.614126623248737)
('flavor', -4.6357357369625625)
('love', -4.66429007687182)
('great', -4.684453739591529)
('use', -4.706008320452014)
('one', -4.762344073425835)
('product', -4.848698285348165)
('tea', -4.856661564089681)
```

#### [5.2.2] Top 10 important features of negative class from SET 2

```
In [24]: # Please write all the code with proper documentation
         # top 10 features
         nb = MultinomialNB(alpha=optimal alpha).fit(X train vec, Y train)
         pos imp features = nb.feature log prob [1,:]
         neg imp features = nb.feature log prob [0,:]
         imp features = {}
         feature names= count vect.get feature names()
         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.itemget
         ter(1), reverse = True)
         print("\n\nNegative top 10 important features are:")
         for i in range(10):
             print(names diff sorted[i])
         Negative top 10 important features are:
         ('probabl', -5.323152765579801)
         ('fruit', -5.333199555522202)
         ('diabet', -5.462993387162749)
         ('creativ', -5.56281606872518)
         ('lodg', -5.617180870847336)
         ('sever', -5.6207953076083825)
         ('matter'. -5.648986574169769)
```

```
('santa', -5.663096226251608)
('outsid', -5.7395052235046435)
('cal', -5.781468980250179)
```

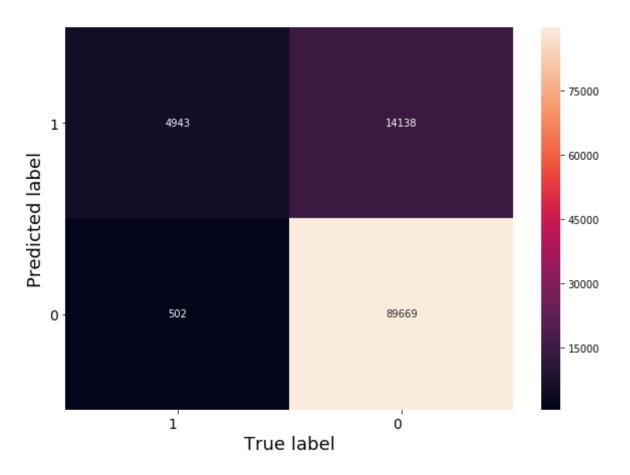
## Accuracy on test data

```
In [25]: # evaluate accuracy
         acc = accuracy score(Y test, predictions) * 100
         print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
         f%%' % (optimal alpha, acc))
         # evaluate precision
         acc = precision score(Y test, predictions, pos label = 1)
         print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
         f' % (optimal alpha, acc))
         # evaluate recall
         acc = recall score(Y test, predictions, pos label = 1)
         print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
         % (optimal alpha, acc))
         # evaluate f1-score
         acc = f1_score(Y_test, predictions, pos label = 1)
         print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f
         ' % (optimal alpha, acc))
          Accuracy of the Multinomial naive Bayes of alpha = 0.100 is 86.599788%
          Precision of the Multinomial naive Bayes of alpha = 0.100 is 0.863805
          Recall of the Multinomial naive Bayes of alpha = 0.100 is 0.994433
          F1-Score of the Multinomial naive Bayes of alpha = 0.100 is 0.924528
In [26]: # Code for drawing seaborn heatmaps
         class names = [1,0]
         df heatmap = pd.DataFrame(confusion matrix(Y test, predictions), index=
```

```
class_names, columns=class_names )
fig = plt.figure(figsize=(10,7))
heatmap = sns.heatmap(df_heatmap, annot=True, fmt="d")

# Setting tick labels for heatmap
heatmap.yaxis.set_ticklabels(heatmap.yaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
heatmap.xaxis.set_ticklabels(heatmap.xaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
plt.ylabel('Predicted label',size=18)
plt.xlabel('True label',size=18)
plt.title("Confusion Matrix\n",size=24)
plt.show()
```





# **Accuracy On train data**

```
In [27]: # evaluate accuracy on train data set
acc = accuracy_score(Y_train,mnbl.predict(X_train_vec)) * 100
print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
f%%' % (optimal_alpha, acc))
```

```
# evaluate precision
acc = precision_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1
)
print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
f' % (optimal_alpha, acc))

# evaluate recall
acc = recall_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1)
print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
% (optimal_alpha, acc))

# evaluate f1-score
acc = f1_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1)
print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f'
' % (optimal_alpha, acc))

Accuracy of the Multinomial naive Bayes of alpha = 0.100 is 88.676403%
```

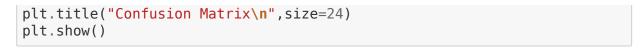
Precision of the Multinomial naive Bayes of alpha = 0.100 is 0.885773

Recall of the Multinomial naive Bayes of alpha = 0.100 is 0.995256

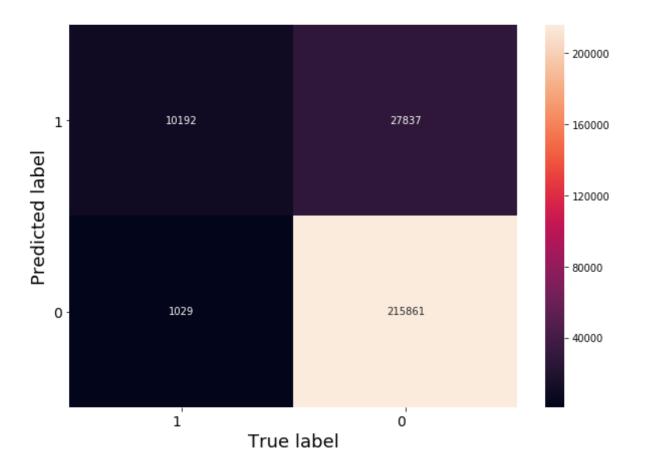
F1-Score of the Multinomial naive Bayes of alpha = 0.100 is 0.937328

```
In [28]: # Code for drawing seaborn heatmaps
    #train confusion matrics
    class_names = [1,0]
    df_heatmap = pd.DataFrame(confusion_matrix(Y_train, mnbl.predict(X_train_vec)), index=class_names, columns=class_names)
    fig = plt.figure(figsize=(10,7))
    heatmap = sns.heatmap(df_heatmap, annot=True, fmt="d")

# Setting tick labels for heatmap
heatmap.yaxis.set_ticklabels(heatmap.yaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
heatmap.xaxis.set_ticklabels(heatmap.xaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
plt.ylabel('Predicted label',size=18)
plt.xlabel('True label',size=18)
```



## **Confusion Matrix**



# (a). Procedure followed:

- STEP 1 :- Text Preprocessing
- STEP 2:- Time-based splitting of whole dataset into train\_data and test\_data

- STEP 3:- Training the vectorizer on train\_data and later applying same vectorizer on both train\_data and test\_data to transform them into vectors
- STEP 4:- Using Bernoulli Naive Bayes as an estimator in 10-Fold Cross-Validation in order to find optimal value of alpha.
- STEP 5:- Draw various plots auc's vs aplha,k
- STEP 6:- Once , we get optimal value of alpha then train BernoulliNB again with this optimal alpha and make predictions on test\_data
- STEP 7:- Find important features per class
- STEP 8 :- Evaluate : Accuracy , F1-Score , Precision , Recall , TPR , FPR , TNR , FNR
- STEP 9:- Draw Seaborn Heatmap for Confusion Matrix .

## [6] Conclusions

```
In [29]: # Creating table using PrettyTable library
from prettytable import PrettyTable

model = ["MultinomialNB for BoW", "MultinomialNB for TFIDF"]

best_alpha = [bow_mnb_alpha, tfidf_mnb_alpha]

train_acc = [bow_mnb_train_acc, tfidf_mnb_train_acc]

test_acc = [bow_mnb_test_acc, tfidf_mnb_test_acc]

sno = [1,2]

Fl_score = [0.941, 0.932]

# Initializing prettytable
ptable = PrettyTable()

# Adding columns
ptable.add_column("S.NO.",sno)
ptable.add_column("MODEL",model)
ptable.add_column("Best Alpha",best_alpha)
```

# feature engineering

# here i combined {cleaned text} and {summary} twice

```
In [30]: conn = sqlite3.connect('featureeng.sqlite')
  final = pd.read_sql_query(""" SELECT * FROM Reviews WHERE Score != 3
    """, conn)
  final.head()
Out[30]:

Level_0 index ld ProductId UserId ProfileName HelpfulnessNumer.
```

	level_0	index	ld	ProductId	Userld	ProfileName	HelpfulnessNur
0	0	138706	150524	0006641040	ACITT7DI6IDDL	shari zychinski	0
1	1	138688	150506	0006641040	A2IW4PEEKO2R0U	Tracy	1
2	2	138689	150507	0006641040	A1S4A3IQ2MU7V4	sally sue "sally sue"	1
3	3	138690	150508	0006641040	AZGXZ2UUK6X	Catherine Hallberg " (Kate)"	1

	level_0	index	ld	ProductId	Userld	ProfileName	HelpfulnessNur
4	4	138691	150509	0006641040	A3CMRKGE0P909G	Teresa	3

In [31]: print(final['CleanedText'][0])

witti littl book make son laugh loud recit car drive along alway sing r efrain hes learn whale india droop love new word book introduc silli cl assic book will bet son still abl recit memori colleg witti littl book make son laugh loud recit car drive along alway sing refrain hes learn whale india droop love new word book introduc silli classic book will b et son still abl recit memori colleg everi book educ everi book educ

In [32]: from sklearn.model\_selection import train\_test\_split
 ##Sorting data according to Time in ascending order for Time Based Spli
 tting
 time\_sorted\_data = final.sort\_values('Time', axis=0, ascending=True, in
 place=False, kind='quicksort', na\_position='last')

X = time\_sorted\_data['CleanedText'].values
Y = time\_sorted\_data['Score']

# 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, shuffle=False)

In [33]: print(X\_train.shape, Y\_train.shape)
print(X\_test.shape, Y\_test.shape)

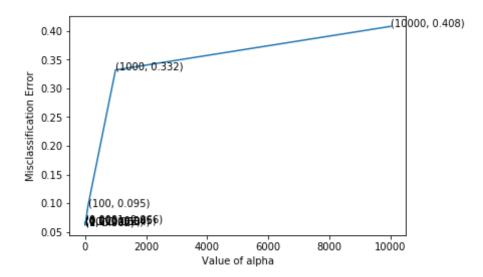
```
(254919,) (254919,)
(109252,) (109252,)
```

## Naive bayes bow

```
In [55]: #BoW
         count vect = CountVectorizer(min df = 50)
         X train vec = count vect.fit transform(X train)
         X test vec = count vect.transform(X test)
         print("the type of count vectorizer :",type(X train vec))
         print("the shape of out text BOW vectorizer : ",X train vec.get shape
         print("the number of unique words :", X train vec.get shape()[1])
         the type of count vectorizer : <class 'scipy.sparse.csr.csr matrix'>
         the shape of out text BOW vectorizer: (254919, 6219)
         the number of unique words : 6219
In [56]: from sklearn.naive bayes import MultinomialNB
         from sklearn.model selection import cross val score
         # Creating alpha values in the range from 10^-4 to 10^4
         neighbors = [10**-4,10**-3,10**-2,10**-1,10**0,10**1,10**2,10**3,10**4]
         # empty list that will hold cv scores
         cv scores = []
         training scores =[]
         # perform 10-fold cross validation
         for k in neighbors:
             nb = MultinomialNB(alpha=k)
             nb.fit(X train vec, Y train)
             #print(nb.predict(X test[2:39]))
             scores = cross val score(nb, X test vec, Y_test, cv=10, scoring='ro
         c auc')
             scores training = nb.fit(X train vec, Y train).score(X train vec, Y
         train)
             cv scores.append(scores.mean())
             training scores.append(scores training)
```

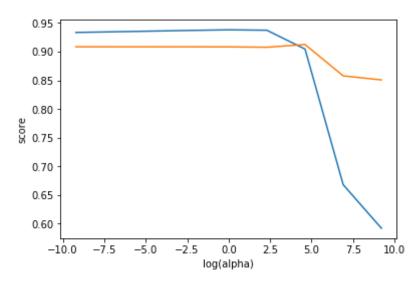
```
#print((nb))
MSE = [1 - x \text{ for } x \text{ in } cv \text{ scores}]
#determining bst k
optimal alpha = neighbors[MSE.index(min(MSE))]
print('\nThe optimal value of alpha is %.8f.' % optimal alpha)
plt.plot(neighbors, MSE)
for xy in zip(neighbors, np.round(MSE,3)):
    plt.annotate('(%s, %s)' % xy, xy=xy, textcoords='data')
plt.xlabel('Value of alpha')
plt.ylabel('Misclassification Error')
plt.show()
print("the misclassification error for each k value is : ", np.round(MS
E,3))
plt.plot(np.log(neighbors), cv scores)
plt.plot(np.log(neighbors), training scores)
plt.xlabel('log(alpha)')
plt.ylabel('score')
# determining best value of alpha
#best alpha = neighbors[cv scores.index(max(cv scores))]
#print('\nThe optimal value of alpha is %.3f.' % best alpha)
```

The optimal value of alpha is 1.00000000.



the misclassification error for each k value is :  $[0.066\ 0.065\ 0.064\ 0.063\ 0.062\ 0.063\ 0.095\ 0.332\ 0.408]$ 

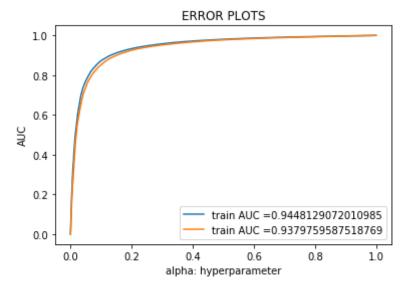
Out[56]: Text(0,0.5,'score')



```
ptimal alpha ======
         # instantiate learning model alpha = optimal alpha
         from sklearn.metrics import accuracy score
         mnb1 = MultinomialNB(alpha = optimal alpha)
         # fitting the model
         mnb1.fit(X train vec, Y train)
         # predict the response
         predictions= mnb1.predict(X test vec)
         predictions1= mnb1.predict(X train vec)
         # evaluate accuracy
         acc = accuracy score(Y test, predictions) * 100
         print('\nThe Test Accuracy of the Multinomial naive Bayes classifier fo
         r alpha = %.3f is %f%%' % (optimal alpha, acc))
         # Variables that will be used for making table in Conclusion part of t
         his assignment
         bow mnb alpha = optimal alpha
         bow mnb train acc = max(cv scores)*100
         bow mnb test acc = acc
         The Test Accuracy of the Multinomial naive Bayes classifier for alpha =
         1.000 is 89.687145%
In [38]: # https://scikit-learn.org/stable/modules/generated/sklearn.metrics.roc
         curve.html#sklearn.metrics.roc curve
         from sklearn.metrics import roc curve, auc
         # roc auc score(y true, y score) the 2nd parameter should be probabilit
         y estimates of the positive class
         # not the predicted outputs
         train fpr, train tpr, thresholds = roc curve(Y train, mnb1.predict prob
         a(X train vec)[:,1])
         test fpr, test tpr, thresholds = roc curve(Y test, mnb1.predict proba(X
         test vec)[:,1])
```

```
plt.plot(train_fpr, train_tpr, label="train AUC ="+str(auc(train_fpr, train_tpr)))
plt.plot(test_fpr, test_tpr, label="train AUC ="+str(auc(test_fpr, test_tpr)))
plt.plot(test_fpr, test_tpr, label="train AUC ="+str(auc(test_fpr, test_tpr)))
plt.legend()
plt.xlabel("alpha: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()

from sklearn.metrics import confusion_matrix
print("Train confusion matrix")
print(confusion_matrix(Y_train, mnb1.predict(X_train_vec)))
print("Test confusion_matrix")
print(confusion_matrix(Y_test, mnb1.predict(X_test_vec)))
```



-----

Train confusion matrix [[ 31438 6591] [ 16758 200132]]

```
Test confusion matrix [[15845 3236] [ 8031 82140]]
```

#### top feaatures

```
In [39]: # Please write all the code with proper documentation
         # top 10 features
         import operator
         from nltk.probability import FreqDist, DictionaryProbDist, ELEProbDist,
          sum logs
         from nltk.classify.api import ClassifierI
         from nltk.classify.naivebayes import NaiveBayesClassifier
         nb = MultinomialNB(alpha=optimal alpha).fit(X train vec, Y train)
         pos imp features = nb.feature log prob [1,:]
         neg imp features = nb.feature log prob [0,:]
         imp features = {}
         feature names= count vect.get feature names()
         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.itemget
         ter(1), reverse = True)
         print("\n\nNegative top 10 important features are:")
         for i in range(10):
             print(names diff sorted[i])
         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.itemget
         ter(1), reverse = True)
         print("Postive top 10 important features are:")
         for i in range(10):
             print(names diff sorted[i])
```

```
Negative top 10 important features are:
('tast', -4.121339830375106)
('like', -4.241897689477554)
('product', -4.420751321071874)
('flavor', -4.71711501591672)
('one', -4.7313051044323835)
('would', -4.892526612966876)
('tri', -4.896172774039712)
('good', -4.905849596162241)
('coffe', -5.005919819152609)
('use', -5.0662578511971805)
Postive top 10 important features are:
('great', -4.384114458401665)
('like', -4.436313730832627)
('tast', -4.446972529290646)
('qood', -4.452495879427524)
('love', -4.547745136136161)
('flavor', -4.633749554154672)
('use', -4.758270705270817)
('tea', -4.779857906273282)
('product', -4.796793016587792)
('one', -4.8027358293294)
```

## Accuracy on test data

```
In [40]: # evaluate accuracy
#libraries
from sklearn.metrics import precision_score
from sklearn.metrics import recall_score
from sklearn.metrics import f1_score

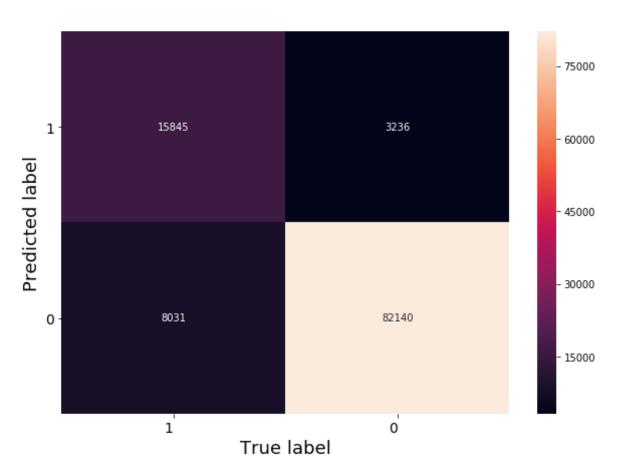
acc = accuracy_score(Y_test, predictions) * 100
print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
f%%' % (optimal_alpha, acc))

# evaluate precision
acc = precision_score(Y_test, predictions, pos_label = 1)
```

```
print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
         f' % (optimal alpha, acc))
         # evaluate recall
         acc = recall score(Y test, predictions, pos label = 1)
         print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
         % (optimal alpha, acc))
         # evaluate f1-score
         acc = f1 score(Y test, predictions, pos label = 1)
         print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f
         ' % (optimal alpha, acc))
          Accuracy of the Multinomial naive Bayes of alpha = 1.000 is 89.687145%
          Precision of the Multinomial naive Bayes of alpha = 1.000 is 0.962097
          Recall of the Multinomial naive Bayes of alpha = 1.000 is 0.910936
          F1-Score of the Multinomial naive Bayes of alpha = 1.000 is 0.935818
In [41]: # Code for drawing seaborn heatmaps
         #test confusion matrics
         class names = [1,0]
         df heatmap = pd.DataFrame(confusion matrix(Y_test, predictions), index=
         class names, columns=class names )
         fig = plt.figure(figsize=(10,7))
         heatmap = sns.heatmap(df heatmap, annot=True, fmt="d")
         # Setting tick labels for heatmap
         heatmap.yaxis.set ticklabels(heatmap.yaxis.get ticklabels(), rotation=0
         , ha='right', fontsize=14)
         heatmap.xaxis.set ticklabels(heatmap.xaxis.get ticklabels(), rotation=0
         , ha='right', fontsize=14)
         plt.ylabel('Predicted label', size=18)
         plt.xlabel('True label', size=18)
         plt.title("Confusion Matrix\n", size=24)
         plt.show()
```

Confincion Matrix





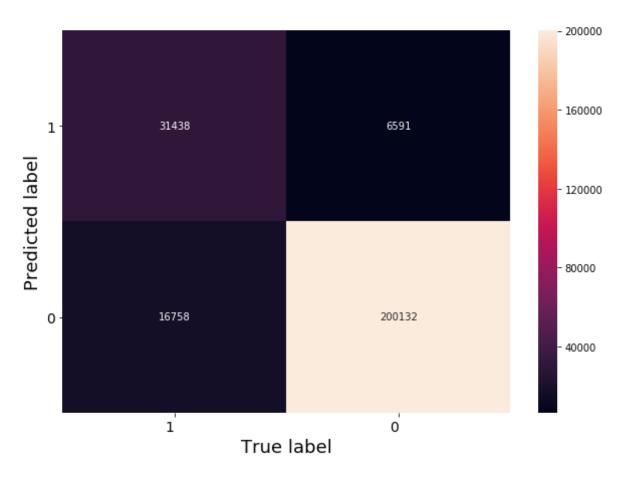
# Accuracy on train data

```
In [42]: # Code for drawing seaborn heatmaps
#train confusion matrics
class_names = [1,0]
df_heatmap = pd.DataFrame(confusion_matrix(Y_train, mnb1.predict(X_train_vec)), index=class_names, columns=class_names)
fig = plt.figure(figsize=(10,7))
```

```
heatmap = sns.heatmap(df_heatmap, annot=True, fmt="d")

# Setting tick labels for heatmap
heatmap.yaxis.set_ticklabels(heatmap.yaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
heatmap.xaxis.set_ticklabels(heatmap.xaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
plt.ylabel('Predicted label',size=18)
plt.xlabel('True label',size=18)
plt.title("Confusion Matrix\n",size=24)
plt.show()
```





```
In [43]: # evaluate accuracy on train data set
acc = accuracy_score(Y_train,mnbl.predict(X_train_vec)) * 100
print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
f%%' % (optimal_alpha, acc))

# evaluate precision
acc = precision_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1
)
print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
```

```
f' % (optimal_alpha, acc))

# evaluate recall
acc = recall_score(Y_train, mnb1.predict(X_train_vec), pos_label = 1)
print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
% (optimal_alpha, acc))

# evaluate f1-score
acc = f1_score(Y_train, mnb1.predict(X_train_vec), pos_label = 1)
print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f'
% (optimal_alpha, acc))
```

Accuracy of the Multinomial naive Bayes of alpha = 1.000 is 90.840620%

Precision of the Multinomial naive Bayes of alpha = 1.000 is 0.968117

Recall of the Multinomial naive Bayes of alpha = 1.000 is 0.922735

F1-Score of the Multinomial naive Bayes of alpha = 1.000 is 0.944881

#### on tfidf

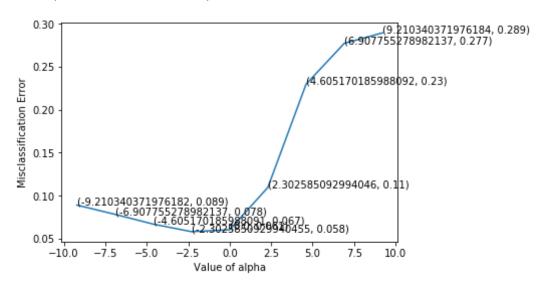
```
In [44]: # Please write all the code with proper documentation
    tf_idf_vect = TfidfVectorizer(min_df=10)
    X_train_vec = tf_idf_vect.fit_transform(X_train)
    X_test_vec = tf_idf_vect.transform(X_test)
    print("the type of count vectorizer : ",type(X_train_vec))
    print("the shape of out text TFIDF vectorizer : ",X_train_vec.get_shape
    ())
    print("the number of unique words : ", X_train_vec.get_shape()[1])

    the type of count vectorizer : <class 'scipy.sparse.csr.csr_matrix'>
        the shape of out text TFIDF vectorizer : (254919, 13053)
    the number of unique words : 13053
In [46]: from sklearn.naive_bayes import MultinomialNB
    from sklearn.model_selection import cross_val_score
```

```
# Creating alpha values in the range from 10^-4 to 10^4
neighbors = [10**-4,10**-3,10**-2,10**-1,10**0,10**1,10**2,10**3,10**4]
# empty list that will hold cv scores
cv scores = []
training scores =[]
# perform 10-fold cross validation
for k in neighbors:
    nb = MultinomialNB(alpha=k)
    nb.fit(X train vec, Y train)
    #print(nb.predict(X test[2:39]))
    scores = cross val score(nb, X test vec, Y test, cv=10, scoring='ro
c auc')
    scores training = nb.fit(X train vec, Y_train).score(X_train_vec, Y
train)
    cv scores.append(scores.mean())
    training scores.append(scores training)
    #print((nb))
MSE = [1 - x \text{ for } x \text{ in } cv \text{ scores}]
#determining best k
optimal alpha = neighbors[MSE.index(min(MSE))]
print('\nThe optimal value of alpha is %.8f.' % optimal alpha)
plt.plot(np.log(neighbors), MSE)
for xy in zip(np.log(neighbors), np.round(MSE,3)):
    plt.annotate('(%s, %s)' % xy, xy=xy, textcoords='data')
plt.xlabel('Value of alpha')
plt.ylabel('Misclassification Error')
plt.show()
print("the misclassification error for each k value is : ", np.round(MS
E,3))
plt.plot(np.log(neighbors), cv scores, 'r')
plt.plot(np.log(neighbors), training scores, 'b')
plt.xlabel('alpha')
plt.vlabel('score')
```

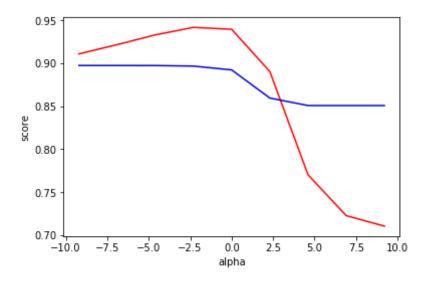
```
# determining best value of alpha
#best_alpha = neighbors[cv_scores.index(max(cv_scores))]
#print('\nThe optimal value of alpha is %.3f.' % best_alpha)
```

The optimal value of alpha is 0.10000000.



the misclassification error for each k value is : [0.089 0.078 0.067 0.058 0.061 0.11 0.23 0.277 0.289]

Out[46]: Text(0,0.5,'score')



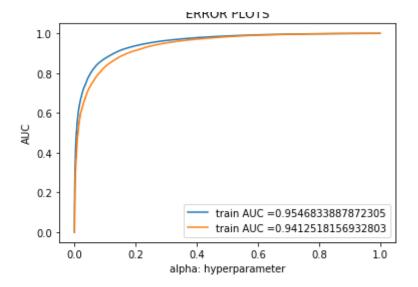
```
ptimal alpha ======
        # instantiate learning model alpha = optimal alpha
        from sklearn.metrics import accuracy score
        mnb1 = MultinomialNB(alpha = optimal alpha)
        # fitting the model
        mnb1.fit(X_train_vec, Y_train)
        # predict the response
        predictions= mnbl.predict(X test vec)
        predictions1= mnb1.predict(X train vec)
        # evaluate accuracy
        acc = accuracy score(Y test, predictions) * 100
        print('\nThe Test Accuracy of the Multinomial naive Bayes classifier fo
        r alpha = %.3f is %f%%' % (optimal alpha, acc))
        # Variables that will be used for making table in Conclusion part of t
        his assignment
        tfidf_mnb_alpha = optimal_alpha
```

```
tfidf_mnb_train_acc = max(cv_scores)*100
tfidf_mnb_test_acc = acc
```

The Test Accuracy of the Multinomial naive Bayes classifier for alpha = 0.100 is 87.818072%

```
In [48]: # https://scikit-learn.org/stable/modules/generated/sklearn.metrics.roc
          curve.html#sklearn.metrics.roc curve
         from sklearn.metrics import roc curve, auc
         # roc auc score(y true, y score) the 2nd parameter should be probabilit
         y estimates of the positive class
         # not the predicted outputs
         train fpr, train tpr, thresholds = roc curve(Y train, mnb1.predict prob
         a(X train vec)[:,1])
         test fpr, test tpr, thresholds = roc curve(Y test, mnb1.predict proba(X
         test vec)[:,1])
         plt.plot(train fpr, train tpr, label="train AUC ="+str(auc(train fpr, t
         rain tpr)))
         plt.plot(test fpr, test tpr, label="train AUC ="+str(auc(test fpr, test
         tpr)))
         plt.legend()
         plt.xlabel("alpha: hyperparameter")
         plt.ylabel("AUC")
         plt.title("ERROR PLOTS")
         plt.show()
         print("="*100)
         from sklearn.metrics import confusion matrix
         print("Train confusion matrix")
         print(confusion matrix(Y train, mnb1.predict(X train vec)))
         print("Test confusion matrix")
         print(confusion matrix(Y test, mnb1.predict(X test vec)))
```

EDDAD DLATE



\_\_\_\_\_\_\_

\_\_\_\_\_\_

```
Train confusion matrix
[[ 12814 25215]
  [ 1124 215766]]
Test confusion matrix
[[ 6300 12781]
  [ 528 89643]]
```

## top features

```
In [49]: # Please write all the code with proper documentation
# top 10 features
import operator
from nltk.probability import FreqDist, DictionaryProbDist, ELEProbDist,
    sum_logs
from nltk.classify.api import ClassifierI
from nltk.classify.naivebayes import NaiveBayesClassifier
nb = MultinomialNB(alpha=optimal_alpha).fit(X_train_vec, Y_train)
pos_imp_features = nb.feature_log_prob_[1,:]
neg_imp_features = nb.feature_log_prob_[0,:]
```

```
imp features = {}
feature names= count vect.get feature names()
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.itemget
ter(1), reverse = True)
print("\n\nNegative top 10 important features are:")
for i in range(10):
    print(names diff sorted[i])
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.itemget
ter(1), reverse = True)
print("Postive top 10 important features are:")
for i in range (10):
    print(names diff sorted[i])
Negative top 10 important features are:
('procedur', -5.312587448981749)
('frys', -5.318103599134482)
('logic', -5.469848179132611)
('diamet', -5.470594991668931)
('shallow', -5.568710118651474)
('matter', -5.607535304607177)
('creativ', -5.609405435847388)
('cake', -5.653858336588055)
('satiat', -5.7055347498175975)
('outing', -5.789471974356955)
Postive top 10 important features are:
('skippi', -4.851330484981341)
('shallow', -5.0082087311264285)
('frys', -5.134762873626148)
('procedur', -5.206547838510216)
('citi', -5.394596512032123)
            5 640901903409752)
('catiat'
```

```
('quit', -5.650236142479857)
('marshmallow', -5.682865187640305)
('diamet', -5.699873409450561)
('back', -5.727390200429296)
```

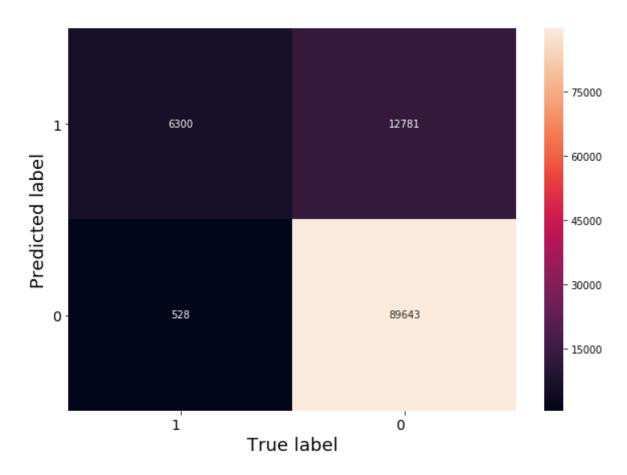
#### Accuracy on test data

```
In [50]: # evaluate accuracy
         acc = accuracy score(Y test, predictions) * 100
         print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
         f%%' % (optimal alpha, acc))
         # evaluate precision
         acc = precision score(Y test, predictions, pos label = 1)
         print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
         f' % (optimal alpha, acc))
         # evaluate recall
         acc = recall score(Y test, predictions, pos label = 1)
         print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
         % (optimal alpha, acc))
         # evaluate f1-score
         acc = f1 score(Y test, predictions, pos label = 1)
         print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f
         ' % (optimal alpha, acc))
          Accuracy of the Multinomial naive Bayes of alpha = 0.100 is 87.818072%
          Precision of the Multinomial naive Bayes of alpha = 0.100 is 0.875215
          Recall of the Multinomial naive Bayes of alpha = 0.100 is 0.994144
          F1-Score of the Multinomial naive Bayes of alpha = 0.100 is 0.930896
In [51]: # Code for drawing seaborn heatmaps
         class names = [1,0]
```

```
df_heatmap = pd.DataFrame(confusion_matrix(Y_test, predictions), index=
class_names, columns=class_names )
fig = plt.figure(figsize=(10,7))
heatmap = sns.heatmap(df_heatmap, annot=True, fmt="d")

# Setting tick labels for heatmap
heatmap.yaxis.set_ticklabels(heatmap.yaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
heatmap.xaxis.set_ticklabels(heatmap.xaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
plt.ylabel('Predicted label',size=18)
plt.xlabel('True label',size=18)
plt.title("Confusion Matrix\n",size=24)
plt.show()
```





# Accuracy on train data

```
In [52]: # evaluate accuracy on train data set
acc = accuracy_score(Y_train,mnbl.predict(X_train_vec) ) * 100
print('\n Accuracy of the Multinomial naive Bayes of alpha = %.3f is %
f%%' % (optimal_alpha, acc))
```

```
# evaluate precision
acc = precision_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1
)
print('\n Precision of the Multinomial naive Bayes of alpha = %.3f is %
f' % (optimal_alpha, acc))

# evaluate recall
acc = recall_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1)
print('\n Recall of the Multinomial naive Bayes of alpha = %.3f is %f'
% (optimal_alpha, acc))

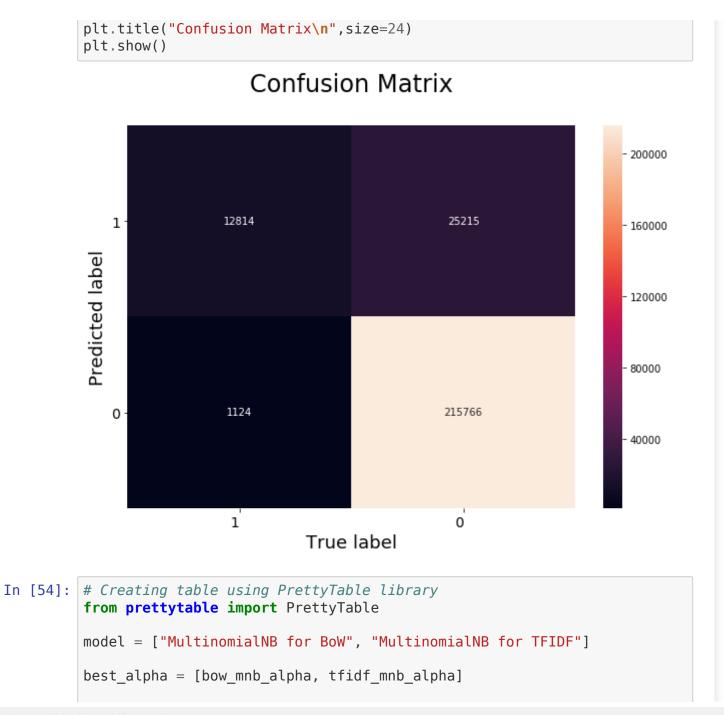
# evaluate f1-score
acc = f1_score(Y_train, mnbl.predict(X_train_vec), pos_label = 1)
print('\n F1-Score of the Multinomial naive Bayes of alpha = %.3f is %f'
% (optimal_alpha, acc))
```

Accuracy of the Multinomial naive Bayes of alpha = 0.100 is 89.667698% Precision of the Multinomial naive Bayes of alpha = 0.100 is 0.895365 Recall of the Multinomial naive Bayes of alpha = 0.100 is 0.994818

F1-Score of the Multinomial naive Bayes of alpha = 0.100 is 0.942475

```
In [53]: # Code for drawing seaborn heatmaps
    #train confusion matrics
    class_names = [1,0]
    df_heatmap = pd.DataFrame(confusion_matrix(Y_train, mnbl.predict(X_train_vec)), index=class_names, columns=class_names)
    fig = plt.figure(figsize=(10,7))
    heatmap = sns.heatmap(df_heatmap, annot=True, fmt="d")

# Setting tick labels for heatmap
heatmap.yaxis.set_ticklabels(heatmap.yaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
heatmap.xaxis.set_ticklabels(heatmap.xaxis.get_ticklabels(), rotation=0
, ha='right', fontsize=14)
plt.ylabel('Predicted label',size=18)
plt.xlabel('True label',size=18)
```



```
train acc = [bow mnb train acc, tfidf mnb train acc]
test acc = [bow mnb test acc, tfidf mnb test acc]
sno = [1,2]
F1 \text{ score} = [0.941, 0.932]
# Initializing prettytable
ptable = PrettvTable()
# Adding columns
ptable.add column("S.NO.",sno)
ptable.add column("MODEL", model)
ptable.add column("Best Alpha", best alpha)
ptable.add column("Training Accuracy", train acc)
ptable.add column("Test Accuracy", test acc)
ptable.add column("F1 Score",F1 score)
# Printing the Table
print(ptable)
Test Accuracy | F1 Score |
   1 | MultinomialNB for BoW | 1 | 93.81366874026789 | 8
9.68714531541757 | 0.941 |
| 2 | MultinomialNB for TFIDF | 0.1 | 94.172472790216 | 8
7.81807198037565 | 0.932 |
-----+
```

# procedure followed

- STEP 1 :- Text Preprocessing(where i combined cleaned text, summary twice)
- STEP 2:- Time-based splitting of whole dataset into train\_data and test\_data
- STEP 3:- Training the vectorizer on train\_data and later applying same vectorizer on both train\_data and test\_data to transform them into vectors
- STEP 4:- Using Bernoulli Naive Bayes as an estimator in 10-Fold Cross-Validation in order to find optimal value of alpha.
- STEP 5:- Draw various plots auc's vs aplha,k
- STEP 6:- Once , we get optimal value of alpha then train BernoulliNB again with this optimal alpha and make predictions on test data
- STEP 7:- Find important features per class
- STEP 8:- Evaluate: Accuracy, F1-Score, Precision, Recall, TPR, FPR, TNR, FNR
- STEP 9:- Draw Seaborn Heatmap for Confusion Matrix .

#### conclusion

- by using feature engineering our performance of the models got increased slighlty
- observe the pretty table before and after feature engineering where accuracy got increased