Amazon Fine Food Reviews Analysis

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

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

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

Number of reviews: 568,454 Number of users: 256,059 Number of products: 74,258 Timespan: Oct 1999 - Oct 2012

Number of Attributes/Columns in data: 10

Attribute Information:

- 1 Id
- 2. ProductId 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 considered 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 will 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 tqdm import tqdm
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 500000 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 Score != 3 LIMIT 500000""", co
n)
# for tsne assignment you can take 5k data points
```

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SELECT * FROM Reviews WHERE Score != 3 LIMIT 500000, will give top 500000 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 Score != 3 LIMIT 500000""", co
n)
for tsne assignment you can take 5k data points

filtered_data = pd.read_sql_query(""" SELECT * FROM Reviews WHERE Score != 3 """, con)

Give reviews with Score>3 a positive rating(1), and reviews with a score<3 a negative rating(0).

def partition(x):
 if x < 3:
 return 0
 return 1

#changing reviews with score less than 3 to be positive and vice-versa
actualScore = filtered_data['Score']
positiveNegative = actualScore.map(partition)
filtered_data['Score'] = positiveNegative
print("Number of data points in our data", filtered_data.shape)
filtered_data.head(3)</pre>

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

Out[2]:

	ld	ProductId	UserId	ProfileName	HelpfulnessNumerator	HelpfulnessDenominator	Score	Time	Summary
0	1	B001E4KFG0	A3SGXH7AUHU8GW	delmartian	1	1	1	1303862400	Good Quality Dog Food

1 2 B00813GRG4 A1D87F6ZCVE5NK dll pa 0 0 1346976000 Not as Advertised

Natalia
Corres "Deligh

```
ABXLMWJIXXAIN Userld ProfileNtantie
       B000LQOCH0
Productid
                                                                                                           1219017600
Time
                                                      HelpfulnessNumerator HelpfulnessDenominator Score
                                                                                                                        Sanyanitany
4
In [3]:
display = pd.read_sql_query("""
SELECT UserId, ProductId, ProfileName, Time, Score, Text, COUNT(*)
FROM Reviews
GROUP BY UserId
HAVING COUNT(*)>1
""", con)
In [4]:
print(display.shape)
display.head()
(80668, 7)
Out[4]:
                  Userld
                             ProductId
                                               ProfileName
                                                                                                                  Text COUNT(*)
                                                                  Time
                                                                       Score
                                                                                     Overall its just OK when considering the
   #oc-R115TNMSPFT9I7 B007Y59HVM
                                                   Breyton 1331510400
                                                                                                                               2
                                             Louis E. Emory
                                                                                       My wife has recurring extreme muscle
    #oc-R11D9D7SHXIJB9 B005HG9ET0
                                                            1342396800
                                                                            5
                                                                                                                               3
                                                    "hoppy
                                                                                                            spasms, u...
                         B007Y59HVM
 2
                                           Kim Cieszykowski
                                                           1348531200
                                                                                This coffee is horrible and unfortunately not ...
                                                                                                                               2
       R11DNU2NBKQ23Z
                    #oc-
 3
                          B005HG9ET0
                                              Penguin Chick 1346889600
                                                                                This will be the bottle that you grab from the...
                                                                                                                               3
       R11O5J5ZVQE25C
                                                                                                                               2
                         B007OSBE1U
                                        Christopher P. Presta 1348617600
                                                                                   I didnt like this coffee. Instead of telling y...
       R12KPBODL2B5ZD
```

In [5]:

```
display[display['UserId'] == 'AZY10LLTJ71NX']
```

Out[5]:

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

In [6]:

```
display['COUNT(*)'].sum()
```

Out[6]:

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 [7]:
```

```
display= pd.read_sql_query("""
splace *
```

```
FROM Reviews
WHERE Score != 3 AND UserId="AR5J8UI46CURR"
ORDER BY ProductID
""", con)
display.head()
```

Out[7]:

	ld	ProductId	UserId	ProfileName	HelpfulnessNumerator	HelpfulnessDenominator	Score	Time	Summ
0	78445	B000HDL1RQ	AR5J8UI46CURR	Geetha Krishnan	2	2	5	1199577600	LOACH QUADRA VANII WAFE
1	138317	B000HDOPYC	AR5J8UI46CURR	Geetha Krishnan	2	2	5	1199577600	LOACH QUADRAT VANII WAFE
2	138277	B000HDOPYM	AR5J8UI46CURR	Geetha Krishnan	2	2	5	1199577600	LOACH QUADRA VANII WAFE
3	73791	B000HDOPZG	AR5J8UI46CURR	Geetha Krishnan	2	2	5	1199577600	LOACH QUADRA VANII WAFE
4	155049	B000PAQ75C	AR5J8UI46CURR	Geetha Krishnan	2	2	5	1199577600	LOACH QUADRA VANII WAFE
4	<u> </u>								

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 [8]:

```
#Sorting data according to ProductId in ascending order
sorted_data=filtered_data.sort_values('ProductId', axis=0, ascending=True, inplace=False, kind='qui
cksort', na_position='last')
```

In [9]:

```
#Deduplication of entries
final=sorted_data.drop_duplicates(subset={"UserId","ProfileName","Time","Text"}, keep='first', inpl
ace=False)
final.shape
```

Out[9]:

(364173, 10)

In [10]:

```
#Checking to see how much % of data still remains
```

```
(IInat['Ia'].Slze'I.U)/(IILterea_aata['Ia'].Slze'I.U)'LUU
Out[10]:
69.25890143662969
Observation:- It was also seen that in two rows given below the value of HelpfulnessNumerator is greater than
HelpfulnessDenominator which is not practically possible hence these two rows too are removed from calcualtions
In [11]:
display= pd.read_sql_query("""
SELECT 3
FROM Reviews
WHERE Score != 3 AND Id=44737 OR Id=64422
ORDER BY ProductID
""", con)
display.head()
Out[11]:
      ld
            ProductId
                               UserId ProfileName HelpfulnessNumerator HelpfulnessDenominator Score
                                                                                                     Time Summary
                                            J. E.
                                                                                                             This for
 0 64422 B000MIDROQ A161DK06JJMCYF
                                         Stephens
                                                                                             5 1224892800
                                                                                                           My Son at
                                         "Jeanne'
                                                                                                             College
                                                                                                               Pure
                                                                                                           taste with
                                                                                             4 1212883200
 1 44737 B001EQ55RW A2V0I904FH7ABY
                                            Ram
                                                                                                            crunchy
                                                                                                            almonds
                                                                                                              inside
4
In [12]:
final=final(final.HelpfulnessNumerator<=final.HelpfulnessDenominator)</pre>
In [13]:
#Before starting the next phase of preprocessing lets see the number of entries left
print(final.shape)
#How many positive and negative reviews are present in our dataset?
final['Score'].value counts()
(364171, 10)
Out[13]:
     307061
```

[3] Preprocessing

Name: Score, dtype: int64

57110

[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 [14]:

```
# 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(sent_1500)
print("="*50)

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

this witty little book makes my son laugh at loud. i recite it in the car as we're driving along a nd he always can sing the refrain. he's learned about whales, India, drooping roses: i love all t he new words this book introduces and the silliness of it all. this is a classic book i am willing to bet my son will STILL be able to recite from memory when he is in college

I was really looking forward to these pods based on the reviews. Starbucks is good, but I prefer bolder taste... imagine my surprise when I ordered 2 boxes - both were expired! One expired back in 2005 for gosh sakes. I admit that Amazon agreed to credit me for cost plus part of shipping, b ut geez, 2 years expired!!! I'm hoping to find local San Diego area shoppe that carries pods so t hat I can try something different than starbucks.

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

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

// Thick, delicious. Perfect. 3 ingredients: Water, Maltitol, Natural Maple Flavor. PERIOD. No chemicals. No garbage.

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

// Tuse this as my SWEETENER in baking: cheesecakes, white brownies, muffins, pumpkin pies, etc... Unbelievably delicious...

// Can you tell I like it? :)

In [15]:

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

this witty little book makes my son laugh at loud. i recite it in the car as we're driving along a nd he always can sing the refrain. he's learned about whales, India, drooping roses: i love all t he new words this book introduces and the silliness of it all. this is a classic book i am willing to bet my son will STILL be able to recite from memory when he is in college

```
In [16]:
```

```
from bs4 import BeautifulSoup
soup = BeautifulSoup(sent 0, 'lxml')
text = soup.get text()
print(text)
print("="*50)
soup = BeautifulSoup(sent 1000, 'lxml')
text = soup.get text()
print(text)
print("="*50)
soup = BeautifulSoup(sent 1500, 'lxml')
text = soup.get text()
print(text)
print("="*50)
soup = BeautifulSoup(sent 4900, 'lxml')
text = soup.get_text()
print(text)
```

this witty little book makes my son laugh at loud. i recite it in the car as we're driving along a nd he always can sing the refrain. he's learned about whales, India, drooping roses: i love all t he new words this book introduces and the silliness of it all. this is a classic book i am willing to bet my son will STILL be able to recite from memory when he is in college

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In [17]:

```
# 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"\'m", " am", phrase)
   return phrase
```

In [18]:

```
sent_1500 = decontracted(sent_1500)
print(sent_1500)
print("="*50)
```

Creat ingradiants although shigher should have been 1st rather than shigher broth the only thing

I do not think belongs in it is Canola oil. Canola or rapeseed is not someting a dog would ever fi nd in nature and if it did find rapeseed in nature and eat it, it would poison them. Today is Food industries have convinced the masses that Canola oil is a safe and even better oil than olive or v irgin coconut, facts though say otherwise. Until the late 70 is it was poisonous until they figured out a way to fix that. I still like it but it could be better.

In [19]:

```
#remove words with numbers python: https://stackoverflow.com/a/18082370/4084039
sent_0 = re.sub("\S*\d\S*", "", sent_0).strip()
print(sent_0)
```

this witty little book makes my son laugh at loud. i recite it in the car as we're driving along a nd he always can sing the refrain. he's learned about whales, India, drooping roses: i love all t he new words this book introduces and the silliness of it all. this is a classic book i am willing to bet my son will STILL be able to recite from memory when he is in college

In [20]:

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

Great ingredients although chicken should have been 1st rather than chicken broth the only thing I do not think belongs in it is Canola oil Canola or rapeseed is not someting a dog would ever find in nature and if it did find rapeseed in nature and eat it it would poison them Today is Food indu stries have convinced the masses that Canola oil is a safe and even better oil than olive or virgi n coconut facts though say otherwise Until the late 70 is it was poisonous until they figured out a way to fix that I still like it but it could be better

In [21]:

```
# https://gist.github.com/sebleier/554280
# we are removing the words from the stop words list: 'no', 'nor', 'not'
# <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', 'ours', 'ourselves', 'you', "y
ou're", "you've", \
            "you'll", "you'd", 'your', 'yours', 'yourself', 'yourselves', 'he', 'him', 'his',
'himself', \
            'she', "she's", 'her', 'hers', 'herself', 'it', "it's", 'its', 'itself', 'they', 'them',
'their',\
            'theirs', 'themselves', 'what', 'which', 'who', 'whom', 'this', 'that', "that'll",
'these', 'those',
            'am', 'is', 'are', 'was', 'were', 'be', 'been', 'being', 'have', 'has', 'had', 'having',
'do', 'does', \
            'did', 'doing', 'a', 'an', 'the', 'and', 'but', 'if', 'or', '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', 'how', 'all', 'any', 'both', '\epsilon
ach', 'few', 'more', \
            'most', 'other', 'some', 'such', 'only', 'own', 'same', 'so', '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', "de
esn't", 'hadn',\
            "hadn't", 'hasn', "hasn't", 'haven', "haven't", 'isn', "isn'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 [22]:

```
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
    sentance = ' '.join(e.lower() for e in sentance.split() if e.lower() not in stopwords)
    preprocessed_reviews.append(sentance.strip())
```

In [23]:

```
preprocessed_reviews[1500]
```

Out[23]:

'great ingredients although chicken rather chicken broth thing not think belongs canola oil canola rapeseed not someting dog would ever find nature find rapeseed nature eat would poison today food industries convinced masses canola oil safe even better oil olive virgin coconut facts though say otherwise late poisonous figured way fix still like could better'

[3.2] Preprocessing Review Summary

In [24]:

```
## Summary preprocessing
from tqdm import tqdm
preprocessed_summary = []
# tqdm is for printing the status bar
for sentance in tqdm(final['Summary'].values):
   sentance = re.sub(r"http\S+", "", sentance)
    sentance = BeautifulSoup(sentance, 'lxml').get text()
   sentance = decontracted(sentance)
    sentance = re.sub("\S*\d\S*", "", sentance).strip()
    sentance = re.sub('[^A-Za-z]+', ' ', sentance)
    # https://gist.github.com/sebleier/554280
    sentance = ' '.join(e.lower() for e in sentance.split() if e.lower() not in stopwords)
    preprocessed_summary.append(sentance.strip())
              | 22243/364171 [00:03<00:55, 6113.33it/s]/home/lab12/anaconda3/lib/python3.7/site-pa
ckages/bs4/__init__.py:273: UserWarning: "b'.'" looks like a filename, not markup. You should prob
ably open this file and pass the filehandle into Beautiful Soup.
 ' Beautiful Soup.' % markup)
              | 97851/364171 [00:16<00:43, 6121.19it/s]/home/lab12/anaconda3/lib/python3.7/site-pa
ckages/bs4/__init__.py:273: UserWarning: "b'.'" looks like a filename, not markup. You should prob
ably open this file and pass the filehandle into Beautiful Soup.
 ' Beautiful Soup.' % markup)
27%|
              | 98464/364171 [00:16<00:43, 6077.36it/s]/home/lab12/anaconda3/lib/python3.7/site-pa
ckages/bs4/ init .py:273: UserWarning: "b'.'" looks like a filename, not markup. You should prob
ably open this file and pass the filehandle into Beautiful Soup.
   Beautiful Soup.' % markup)
              | 216616/364171 [00:36<00:24, 5989.05it/s]/home/lab12/anaconda3/lib/python3.7/site-r
ackages/bs4/
             _init__.py:273: UserWarning: "b'.'" looks like a filename, not markup. You should pro
bably open this file and pass the filehandle into Beautiful Soup.
  ' Beautiful Soup.' % markup)
           | 354825/364171 [00:59<00:01, 5985.96it/s]/home/lab12/anaconda3/lib/python3.7/site-
              init .py:273: UserWarning: "b'.'" looks like a filename, not markup. You should pr
packages/bs4/_
obably open this file and pass the filehandle into Beautiful Soup.
  ' Beautiful Soup.' % markup)
           | 364171/364171 [01:00<00:00, 5993.20it/s]
4
```

In [23]:

```
final['Cleaned_text'] = preprocessed_reviews
```

```
In [24]:
### Sort data according to time series
final.sort values('Time', inplace=True)
In [25]:
### Taking 100k samples
final 100k = final.sample(n=100000)
In [26]:
x = final 100k['Cleaned text']
Out[26]:
100000
In [27]:
y = final 100k['Score']
y.size
Out [27]:
100000
In [28]:
from sklearn.model_selection import train_test_split
x_train,x_test,y_train,y_test = train_test_split(x,y,test_size=0.3,random_state=42)
```

[4] Featurization

[4.1] BAG OF WORDS

```
In [29]:
#BoW
count vect = CountVectorizer() #in scikit-learn
x_train_bow = count_vect.fit_transform(x_train)
print("some feature names ", count vect.get feature names()[:10])
print('='*50)
x_test_bow = count_vect.transform(x_test)
print("the type of count vectorizer ",type(x_test_bow))
print("the shape of out text BOW vectorizer ",x_test_bow.get_shape())
print("the number of unique words ", x_test_bow.get_shape()[1])
aaaaaahhhh', 'aaaaaaand', 'aaaaaawwwwwwwwwww']
_____
the type of count vectorizer <class 'scipy.sparse.csr.csr_matrix'>
the shape of out text BOW vectorizer (30000, 50937)
the number of unique words 50937
In [30]:
from sklearn import preprocessing
x_train_bow = preprocessing.normalize(x_train_bow)
x_test_bow = preprocessing.normalize(x_test_bow)
```

[4.2] Bi-Grams and n-Grams.

In [32]: #bi-gram, tri-gram and n-gram #removing stop words like "not" should be avoided before building n-grams # count vect = CountVectorizer(ngram range=(1,2)) # please do read the CountVectorizer documentation http://scikitlearn.org/stable/modules/generated/sklearn.feature extraction.text.CountVectorizer.html # you can choose these numebrs min df=10, max features=5000, of your choice count vect = CountVectorizer(ngram range=(1,2), min df=10, max features=5000) final_bigram_counts = count_vect.fit_transform(x_train) print("the type of count vectorizer ",type(final_bigram_counts)) print("the shape of out text BOW vectorizer ",final_bigram_counts.get_shape()) print("the number of unique words including both unigrams and bigrams ", final bigram counts.get s hape()[1]) the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>

the shape of out text BOW vectorizer (70000, 5000) the number of unique words including both unigrams and bigrams 5000

[4.3] TF-IDF

```
In [31]:
```

```
tf idf vect = TfidfVectorizer(ngram range=(1,2), min df=10)
x_train_tfidf = tf_idf_vect.fit_transform(x_train)
print("some sample features(unique words in the corpus)",tf_idf_vect.get_feature_names()[0:10])
print('='*50)
x test tfidf = tf idf vect.transform(x test)
print("the type of count vectorizer ", type (x test tfidf))
print("the shape of out text TFIDF vectorizer ",x test tfidf.get shape())
print ("the number of unique words including both unigrams and bigrams ", x test tfidf.get shape()[
1])
some sample features (unique words in the corpus) ['abandon', 'abandoned', 'abdominal', 'ability',
'able', 'able buy', 'able chew', 'able cut', 'able drink', 'able eat']
_____
the type of count vectorizer <class 'scipy.sparse.csr.csr_matrix'>
the shape of out text TFIDF vectorizer (30000, 40848)
the number of unique words including both unigrams and bigrams 40848
In [32]:
from sklearn import preprocessing
x_train_tfidf = preprocessing.normalize(x_train_tfidf)
x test tfidf = preprocessing.normalize(x test tfidf)
```

[4.4] Word2Vec

```
In [33]:
```

```
# Train your own Word2Vec model using your own text corpus
list of sentance train=[]
for sentance in x train:
   list of sentance train.append(sentance.split())
```

In [34]:

```
# Test your own Word2Vec model using your own text corpus
i=0
list_of_sentance_test=[]
for sentance in x test:
    list of sentance test.append(sentance.split())
```

```
In [35]:
# Using Google News Word2Vectors
# in this project we are using a pretrained model by google
# its 3.3G file, once you load this into your memory
# it occupies ~9Gb, so please do this step only if you have >12G of ram
# we will provide a pickle file wich contains a dict ,
 # and it contains all our courpus words as keys and model[word] as values
# To use this code-snippet, download "GoogleNews-vectors-negative300.bin"
# from https://drive.google.com/file/d/0B7XkCwpI5KDYN1NUTT1SS21pQmM/edit
# it's 1.9GB in size.
# http://kavita-ganesan.com/gensim-word2vec-tutorial-starter-code/#.W17SRFAzZPY
# you can comment this whole cell
# or change these varible according to your need
is_your_ram_gt_16g=False
want to use google w2v = False
want_to_train_w2v = True
if want to train w2v:
     # min_count = 5 considers only words that occured atleast 5 times
     w2v model=Word2Vec(list of sentance train,min count=5,size=50, workers=4)
     print(w2v model.wv.most similar('great'))
     print('='*50)
    print(w2v model.wv.most similar('worst'))
elif want_to_use_google_w2v and is_your_ram_gt_16g:
     if os.path.isfile('GoogleNews-vectors-negative300.bin'):
         w2v model=KeyedVectors.load word2vec format('GoogleNews-vectors-negative300.bin', binary=Tr
ue)
         print(w2v model.wv.most similar('great'))
         print(w2v_model.wv.most_similar('worst'))
     else:
         print("you don't have gogole's word2vec file, keep want to train w2v = True, to train your
own w2v ")
4
[('awesome', 0.8440885543823242), ('terrific', 0.825779378414154), ('wonderful',
0.8151693344116211), ('fantastic', 0.8132066130638123), ('excellent', 0.8085969686508179),
('good', 0.8001235127449036), ('perfect', 0.7367382645606995), ('amazing', 0.7261995077133179), ('
fabulous', 0.7111331224441528), ('incredible', 0.6996779441833496)]
[('nastiest', 0.7835044264793396), ('tastiest', 0.7758582830429077), ('best', 0.7460561990737915),
('greatest', 0.7407208681106567), ('saltiest', 0.6663205027580261), ('superior',
0.6384615302085876), ('disgusting', 0.6267989277839661), ('healthiest', 0.5846405029296875),
('grossest', 0.5758904218673706), ('smoothest', 0.5751399397850037)]
In [36]:
w2v words = list(w2v model.wv.vocab)
print("number of words that occured minimum 5 times ",len(w2v words))
print("sample words ", w2v words[0:50])
number of words that occured minimum 5 times 16192
sample words ['pure', 'toxic', 'garbage', 'eat', 'poison', 'want', 'metals', 'go', 'ahead', 'buy', 'junk', 'feed', 'look', 'high', 'ratings', 'apparently', 'people', 'love', 'eating', 'not', 'harmful', 'would', 'sodium', 'mercury', 'phosphate', 'enjoy', 'got', 'oil', 'free', 'ordered', 's tuff', 'started', 'okay', 'based', 'smell', 'think', 'turned', 'rancid', 'time', 'let', 'tell', 'c ooking', 'coconut', 'no', 'picnic', 'edit', 'year', 'later', 'may', 'lowered']
```

[4.4.1] Converting text into vectors using Avg W2V, TFIDF-W2V

[4.4.1.1] Avg W2v

```
In [37]:
```

```
# average Word2Vec
# compute average word2vec for each review.
```

```
sent vectors train = []; # the avg-w2v for each sentence/review is stored in this list
for sent in tqdm(list of sentance train): # for each review/sentence
   sent vec = np.zeros(50) # as word vectors are of zero length 50, you might need to change this
to 300 if you use google's w2v
   cnt words =0; # num of words with a valid vector in the sentence/review
    for word in sent: # for each word in a review/sentence
        if word in w2v words:
            vec = w2v model.wv[word]
            sent vec += vec
           cnt_words += 1
    if cnt words != 0:
       sent vec /= cnt words
    sent vectors train.append(sent_vec)
print(len(sent vectors train))
print(len(sent vectors train[0]))
100%|
        | 70000/70000 [03:16<00:00, 356.77it/s]
70000
```

In [38]:

50

```
# average Word2Vec
# compute average word2vec for each review.
sent vectors test = []; # the avg-w2v for each sentence/review is stored in this list
for sent in tqdm(list of sentance test): # for each review/sentence
   sent vec = np.zeros(50) # as word vectors are of zero length 50, you might need to change this
to 300 if you use google's w2v
   cnt words =0; # num of words with a valid vector in the sentence/review
    for word in sent: # for each word in a review/sentence
       if word in w2v words:
           vec = w2v_model.wv[word]
           sent vec += vec
           cnt_words += 1
    if cnt words != 0:
       sent vec /= cnt words
    sent_vectors_test.append(sent_vec)
print(len(sent_vectors_test))
print(len(sent_vectors_test[0]))
100%| 30000/30000 [01:19<00:00, 377.26it/s]
```

30000 50

```
In [39]:
```

```
x_train_avgw2v=sent_vectors_train
x_test_avgw2v=sent_vectors_test
```

In [40]:

```
from sklearn import preprocessing
x_train_avgw2v = preprocessing.normalize(x_train_avgw2v)
x_test_avgw2v = preprocessing.normalize(x_test_avgw2v)
```

[4.4.1.2] TFIDF weighted W2v

```
In [41]:
```

```
# S = ["abc def pqr", "def def def abc", "pqr pqr def"]
model = TfidfVectorizer(min_df=10, max_features=500)
tf_idf_matrix = model.fit_transform(x_train)
# we are converting a dictionary with word as a key, and the idf as a value
dictionary = dict(zip(model.get_feature_names(), list(model.idf_)))
```

```
In [42]:
```

```
# TF-IDF weighted Word2Vec
tfidf feat = model.get feature names() # tfidf words/col-names
# final tf idf is the sparse matrix with row= sentence, col=word and cell val = tfidf
tfidf sent vectors train = []; # the tfidf-w2v for each sentence/review is stored in this list
for sent in tqdm(list_of_sentance_train): # for each review/sentence
   sent vec = np.zeros(50) # as word vectors are of zero length
    weight sum =0; # num of words with a valid vector in the sentence/review
    for word in sent: # for each word in a review/sentence
       if word in w2v words and word in tfidf feat:
           vec = w2v model.wv[word]
             tf_idf = tf_idf_matrix[row, tfidf_feat.index(word)]
            # to reduce the computation we are
            # dictionary[word] = idf value of word in whole courpus
            # sent.count(word) = tf valeus of word in this review
           tf idf = dictionary[word] * (sent.count(word) /len(sent))
           sent vec += (vec * tf idf)
           weight sum += tf idf
    if weight sum != 0:
       sent vec /= weight sum
    tfidf sent vectors train.append(sent vec)
    row += 1
100%| 70000/70000 [03:25<00:00, 341.30it/s]
```

In [43]:

```
tfidf sent vectors test = []; # the tfidf-w2v for each sentence/review is stored in this list
for sent in tqdm(list of sentance test): # for each review/sentence
   sent vec = np.zeros(50) # as word vectors are of zero length
    weight sum =0; # num of words with a valid vector in the sentence/review
    for word in sent: # for each word in a review/sentence
       if word in w2v words and word in tfidf feat:
           vec = w2v model.wv[word]
             tf idf = tf idf matrix[row, tfidf feat.index(word)]
            # to reduce the computation we are
            # dictionary[word] = idf value of word in whole courpus
            # sent.count(word) = tf valeus of word in this review
           tf idf = dictionary[word] * (sent.count(word) /len(sent))
           sent_vec += (vec * tf idf)
           weight sum += tf idf
    if weight sum != 0:
       sent_vec /= weight_sum
    tfidf sent vectors test.append(sent vec)
    row += 1
100%| 30000/30000 [01:29<00:00, 334.85it/s]
```

In [44]:

```
x_train_tfidfw2v = tfidf_sent_vectors_train
x_test_tfidfw2v = tfidf_sent_vectors_test
```

In [45]:

```
from sklearn import preprocessing
x_train_tfidfw2v = preprocessing.normalize(x_train_tfidfw2v)
x_test_tfidfw2v = preprocessing.normalize(x_test_tfidfw2v)
```

[5] Assignment 9: Random Forests

1. Apply Random Forests & GBDT on these feature sets

- SET 1: Review text, preprocessed one converted into vectors using (BOW)
- SET 2:Review text, preprocessed one converted into vectors using (TFIDF)
- SET 3:Review text, preprocessed one converted into vectors using (AVG W2v)
- SET 4:Review text, preprocessed one converted into vectors using (TFIDF W2v)

2. The hyper paramter tuning (Consider two hyperparameters: n_estimators & max_depth)

- Find the best hyper parameter which will give the maximum AUC value
- Find the best hyper paramter using k-fold cross validation or simple cross validation data
- Use gridsearch cv or randomsearch cv or you can also write your own for loops to do this task of hyperparameter tuning

3. Feature importance

• Get top 20 important features and represent them in a word cloud. Do this for BOW & TFIDF.

4. Feature engineering

- To increase the performance of your model, you can also experiment with with feature engineering like:
 - Taking length of reviews as another feature.
 - Considering some features from review summary as well.

5. Representation of results

• You need to plot the performance of model both on train data and cross validation data for each hyper parameter, like shown in the figure

with X-axis as **n_estimators**, Y-axis as **max_depth**, and Z-axis as **AUC Score**, we have given the notebook which explains how to plot this 3d plot, you can find it in the same drive 3d_scatter_plot.ipynb



• You need to plot the performance of model both on train data and cross validation data for each hyper parameter, like shown in the figure

<u>seaborn heat maps</u> with rows as n_estimators, columns as max_depth, and values inside the cell representing AUC Score

- You choose either of the plotting techniques out of 3d plot or heat map
- Once after you found the best hyper parameter, you need to train your model with it, and find the AUC on test data and plot the ROC curve on both train and test.
- Along with plotting ROC curve, you need to print the <u>confusion matrix</u> with predicted and original labels of test data points. Please visualize your confusion matrices using <u>seaborn heatmaps</u>.

6. Conclusion

• You need to summarize the results at the end of the notebook, summarize it in the table format. To print out a table please refer to this prettytable library link

Note: Data Leakage

- 1. There will be an issue of data-leakage if you vectorize the entire data and then split it into train/cv/test.
- 2. To avoid the issue of data-leakag, make sure to split your data first and then vectorize it.
- 3. While vectorizing your data, apply the method fit_transform() on you train data, and apply the method transform() on cv/test data.
- 4. For more details please go through this link.

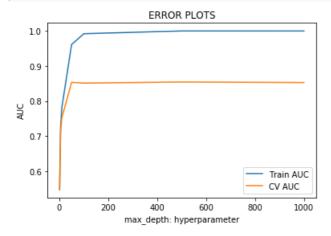
[5.1] Applying RF

[5.1.1] Applying Random Forests on BOW, SET 1

In [46]:

```
## find hyperparameter using roc_auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc_auc_score
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]
```

```
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for d in max depth values:
    clf = RandomForestClassifier(max depth=d)
    clf.fit(x train bow, y train)
    y train pred = clf.predict proba(x train bow)[:,1]
    y test pred = clf.predict proba(x test bow)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv scores.append(roc auc score(y test, y test pred))
#plot cross-validated score, training score vs alpha
plt.plot(max_depth_values, training_scores, label='Train AUC')
plt.plot(max_depth_values, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("max depth: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [45]:

```
## find hyperparameter using roc_auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc auc score
base learners = [40,60,80,100,120]
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for n in base learners:
   clf = RandomForestClassifier(n estimators=n)
   clf.fit(x_train_bow, y_train)
   y train pred = clf.predict proba(x train bow)[:,1]
   y_test_pred = clf.predict_proba(x_test_bow)[:,1]
    training scores.append(roc auc score(y train,y train pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base learners, cv scores, label='CV AUC')
plt.legend()
plt.xlabel("n estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```

```
0.98 - 0.96 - 0.94 - 0.92 - Train AUC CV AUC - CV AUC n_estimator: hyperparameter
```

In [47]:

```
### Using GridsearchCV
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import GridSearchCV
base learners = [40,60,80,100,120]
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]
param grid = {'n estimators': base learners , "max depth":max depth values}
RFC = RandomForestClassifier(max_features='sqrt')
model = GridSearchCV(RFC, param_grid, scoring = 'roc_auc', cv=3 , n_jobs = -1,pre_dispatch=2)
model.fit(x train bow, y train)
print("Model with best parameters :\n", model.best_estimator_)
print("Accuracy of the model : ", model.score(x test bow, y test))
Model with best parameters :
 RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
            max depth=1000, max features='sqrt', max leaf nodes=None,
            min_impurity_decrease=0.0, min_impurity_split=None,
            min samples leaf=1, min samples split=2,
            min weight fraction leaf=0.0, n estimators=120, n jobs=None,
            oob score=False, random state=None, verbose=0,
            warm start=False)
Accuracy of the model : 0.9285171888288075
```

In [46]:

```
#### 3D Scatter Plot
import plotly.offline as offline
import plotly.graph_objs as go
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
y1 = [1, 5, 10, 50, 100]
z1 = training_scores

x2 = [40,60,80,100,120]
y2 = [1, 5, 10, 50, 100]
z2 = cv_scores
```

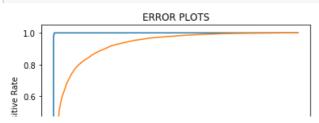
In [47]:

Observation: Using GridsearchCV max_depth is 1000 and n_esimators is 120.

ROC Curve using false positive rate versus true positive rate

```
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
from sklearn.ensemble import RandomForestClassifier
clf = RandomForestClassifier(max_depth=1000,n_estimators=120)
clf.fit(x_train_bow, y_train)
y pred = clf.predict(x test bow)
# roc_auc_score(y_true, y_score) the 2nd parameter should be probability estimates of the positive
class
# not the predicted outputs
train fpr, train tpr, thresholds = roc curve(y train, clf.predict proba(x train bow)[:,1])
test_fpr, test_tpr, thresholds = roc_curve(y_test, clf.predict_proba(x_test_bow)[:,1])
plt.plot(train fpr, train tpr, label="train AUC ="+str(auc(train fpr, train tpr)))
plt.plot(test fpr, test tpr, label="test AUC ="+str(auc(test fpr, test tpr)))
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
print("="*100)
from sklearn.metrics import confusion_matrix
print("Train confusion matrix")
print(confusion matrix(y train, clf.predict(x train bow)))
print("Test confusion matrix")
print(confusion matrix(y test, clf.predict(x test bow)))
```



```
0.2 - train AUC = 0.9999683095615042 test AUC = 0.9283141154625595 0.0 0.0 0.2 0.4 0.6 0.8 1.0 False Positive Rate
```

```
Train confusion matrix
[[10786 26]
[ 0 59188]]
Test confusion matrix
[[ 918 3917]
[ 46 25119]]
```

)

In [49]:

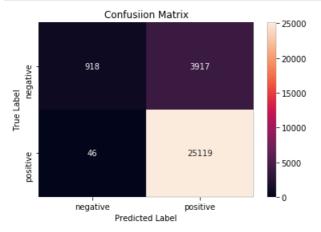
```
# Confusion Matrix
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test, y_pred)
cm
```

Out[49]:

```
array([[ 918, 3917],
        [ 46, 25119]])
```

In [50]:

```
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class_label = ["negative", "positive"]
df_cm = pd.DataFrame(cm, index = class_label, columns = class_label)
sns.heatmap(df_cm, annot = True, fmt = "d")
plt.title("Confusion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
```



[5.1.2] Wordcloud of top 20 important features from SET 1

In [51]:

```
# Calculate feature importances from decision trees
importances = clf.feature_importances_

# Sort feature importances in descending order
indices = np.argsort(importances)[::-1][:20]

# Rearrange feature names so they match the sorted feature importances
names = count vect get feature names()
```

```
type (names)
Out[51]:
list
In [53]:
# Python program to generate WordCloud
###https://www.geeksforgeeks.org/generating-word-cloud-python/
# importing all necessery modules
from wordcloud import WordCloud, STOPWORDS
import matplotlib.pyplot as plt
import pandas as pd
comment_words = ' '
stopwords = set(STOPWORDS)
# iterate through the csv file
for val in names:
# typecaste each val to string
    val = str(val)
# split the value
    tokens = val.split()
# Converts each token into lowercase
    for i in range(len(tokens)):
       tokens[i] = tokens[i].lower()
    for words in tokens:
        comment_words = comment_words + words + ' '
wordcloud = WordCloud (width = 800, height = 800, background color ='white', stopwords = stopwords,
min font size = 10).generate(comment words)
# plot the WordCloud image
plt.figure(figsize = (8, 8), facecolor = None)
plt.imshow(wordcloud)
plt.axis("off")
plt.tight layout(pad = 0)
plt.show()
```

mames - comme_vecc.dec_teachte_mames()

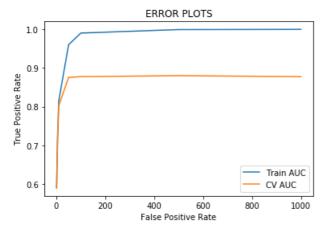




[5.1.3] Applying Random Forests on TFIDF, SET 2

In [48]:

```
## find hyperparameter using roc_auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc auc score
max depth values = [1, 5, 10, 50, 100, 500, 1000]
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for d in max depth values:
   clf = RandomForestClassifier(max depth=d)
   clf.fit(x_train_tfidf, y_train)
   y train pred = clf.predict proba(x train tfidf)[:,1]
   y_test_pred = clf.predict_proba(x_test_tfidf)[:,1]
    training scores.append(roc auc score(y train,y train pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(max depth values, training scores, label='Train AUC')
plt.plot(max depth values, cv scores, label='CV AUC')
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
```



In [55]:

```
## find hyperparameter using roc_auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc_auc_score
base_learners = [40,60,80,100,120]

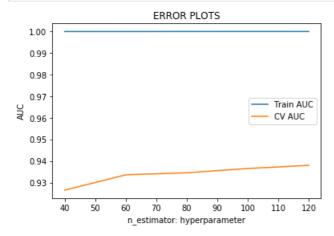
#empty lists that stores cv scores and training_scores
cv_scores = []
training_scores = []

#perform k fold cross validation
for n in base_learners:
    clf = RandomForestClassifier(n_estimators=n)
    clf.fit(x_train_tfidf, y_train)
    y_train_pred = clf.predict_proba(x_train_tfidf)[:,1]
```

```
y_test_pred = clf.predict_proba(x_test_tfidf)[:,1]

training_scores.append(roc_auc_score(y_train,y_train_pred))
cv_scores.append(roc_auc_score(y_test, y_test_pred))

#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base_learners, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("n_estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [51]:

```
## Using GridsearchCV
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import GridSearchCV
base learners = [40,60,80,100,120]
max depth values = [1, 5, 10, 50, 100, 500, 1000]
param_grid = {'n_estimators': base_learners , "max_depth":max_depth_values}
RFC = RandomForestClassifier(max features='sqrt')
model = GridSearchCV(RFC, param_grid, scoring = 'accuracy', cv=3 , n_jobs = -1,pre_dispatch=2)
model.fit(x train tfidf, y train)
print("Model with best parameters :\n", model.best estimator )
print("Accuracy of the model : ",model.score(x test tfidf, y test))
Model with best parameters :
 RandomForestClassifier(bootstrap=True, class_weight=None, criterion='gini',
            max_depth=500, max_features='sqrt', max_leaf_nodes=None,
            min impurity decrease=0.0, min impurity split=None,
            min_samples_leaf=1, min_samples_split=2,
            min_weight_fraction_leaf=0.0, n_estimators=40, n_jobs=None,
            oob score=False, random state=None, verbose=0,
            warm start=False)
Accuracy of the model : 0.890666666666667
```

In [56]:

```
#### 3D Scatter Plot
import plotly.offline as offline
import plotly.graph_objs as go
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
y1 = [1, 5, 10, 50, 100]
z1 = training_scores

x2 = [40,60,80,100,120]
y2 = [1, 5, 10, 50, 100]
z2 = cv_scores
```

In [57]:

Observation: Using GridsearchCV max_depth is 500 and n_esimators is 40.

```
In [ ]:
```

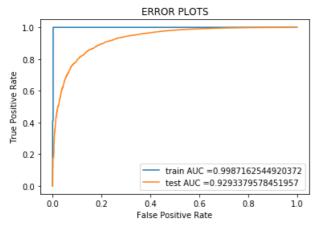
```
### ROC Curve using false positive rate versus true positive rate
```

In [52]:

```
# https://scikit-
learn.org/stable/modules/generated/sklearn.metrics.roc curve.html#sklearn.metrics.roc curve
from sklearn.metrics import roc curve, auc
clf = RandomForestClassifier(max depth=500, n estimators=40)
clf.fit(x train tfidf, y train)
y pred = clf.predict(x test tfidf)
# roc auc score(y true, y score) the 2nd parameter should be probability estimates of the positive
class
# not the predicted outputs
train fpr, train tpr, thresholds = roc curve(y train, clf.predict proba(x train tfidf)[:,1])
test_fpr, test_tpr, thresholds = roc_curve(y_test, clf.predict_proba(x_test_tfidf)[:,1])
plt.plot(train fpr, train tpr, label="train AUC ="+str(auc(train fpr, train tpr)))
plt.plot(test fpr, test tpr, label="test AUC ="+str(auc(test fpr, test tpr)))
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
```

```
plt.show()
print("="*100)

from sklearn.metrics import confusion_matrix
print("Train confusion matrix")
print(confusion_matrix(y_train, clf.predict(x_train_tfidf)))
print("Test confusion matrix")
print(confusion_matrix(y_test, clf.predict(x_test_tfidf)))
```



```
Train confusion matrix
[[10962 55]
[ 0 58983]]
Test confusion matrix
[[ 1515 3137]
[ 131 25217]]
```

In [53]:

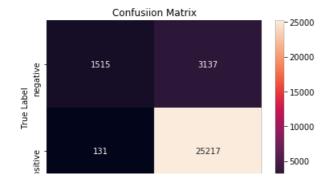
```
# Confusion Matrix
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test, y_pred)
cm
```

Out[53]:

```
array([[ 1515, 3137], [ 131, 25217]])
```

In [54]:

```
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class_label = ["negative", "positive"]
df_cm = pd.DataFrame(cm, index = class_label, columns = class_label)
sns.heatmap(df_cm, annot = True, fmt = "d")
plt.title("Confusiion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
```



[5.1.4] Wordcloud of top 20 important features from SET 2

In [70]:

```
# Calculate feature importances from decision trees
importances = clf.feature_importances_

# Sort feature importances in descending order
indices = np.argsort(importances)[::-1][:20]

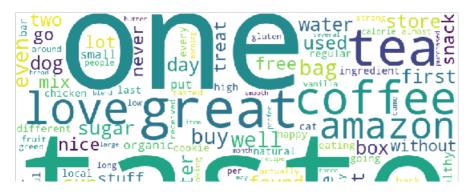
# Rearrange feature names so they match the sorted feature importances
names = tf_idf_vect.get_feature_names()
type(names)
```

Out[70]:

list

In [71]:

```
####https://www.geeksforgeeks.org/generating-word-cloud-python/
# importing all necessery modules
from wordcloud import WordCloud, STOPWORDS
import matplotlib.pyplot as plt
import pandas as pd
comment words = ' '
stopwords = set(STOPWORDS)
# iterate through the csv file
for val in names:
# typecaste each val to string
    val = str(val)
# split the value
   tokens = val.split()
# Converts each token into lowercase
   for i in range(len(tokens)):
       tokens[i] = tokens[i].lower()
    for words in tokens:
       comment words = comment words + words + ' '
wordcloud = WordCloud (width = 800, height = 800, background color ='white', stopwords = stopwords,
min font size = 10).generate(comment words)
# plot the WordCloud image
plt.figure(figsize = (8, 8), facecolor = None)
plt.imshow(wordcloud)
plt.axis("off")
plt.tight_layout(pad = 0)
plt.show()
```

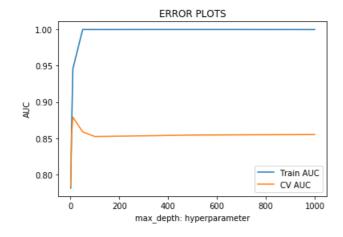




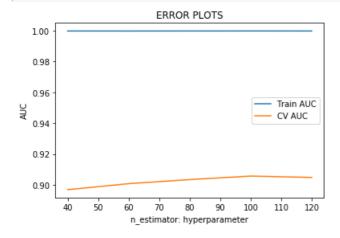
[5.1.5] Applying Random Forests on AVG W2V, SET 3

```
In [58]:
```

```
## find hyperparameter using roc_auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc auc score
max depth values = [1, 5, 10, 50, 100, 500, 1000]
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for d in max depth values:
   clf = RandomForestClassifier(max depth=d)
   clf.fit(x train_avgw2v, y_train)
   y train pred = clf.predict proba(x train avgw2v)[:,1]
    y_test_pred = clf.predict_proba(x_test_avgw2v)[:,1]
    training scores.append(roc auc score(y train,y train pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(max depth values, training scores, label='Train AUC')
plt.plot(max_depth_values, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
```



```
## find hyperparameter using roc auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc auc score
base learners = [40,60,80,100,120]
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for n in base learners:
    clf = RandomForestClassifier(n_estimators=n)
    clf.fit(x_train_avgw2v, y_train)
    y train pred = clf.predict proba(x train avgw2v)[:,1]
    y test_pred = clf.predict_proba(x_test_avgw2v)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base_learners, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("n_estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [59]:

```
## Using GridsearchCV
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import GridSearchCV
base learners = [40,60,80,100,120]
max depth values = [1, 5, 10, 50, 100, 500, 1000]
param grid = {'n estimators': base learners , "max depth":max depth values}
RFC = RandomForestClassifier(max features='sqrt')
model = GridSearchCV(RFC, param grid,cv=3 ,pre dispatch=2,scoring='roc auc')
model.fit(x_train_avgw2v, y_train)
print("Model with best parameters :\n", model.best estimator )
print("Accuracy of the model : ", model.score(x_test_avgw2v, y_test))
Model with best parameters :
 RandomForestClassifier(bootstrap=True, class weight=None, criterion='gini',
            max depth=100, max features='sqrt', max leaf nodes=None,
            min_impurity_decrease=0.0, min_impurity_split=None,
            min samples leaf=1, min samples split=2,
            min weight fraction leaf=0.0, n estimators=120, n jobs=None,
            oob score=False, random state=None, verbose=0,
            warm start=False)
Accuracy of the model: 0.8997360015989295
```

In [60]:

```
#### 3D Scatter Plot
import plotly.offline as offline
import plotly.graph_objs as go
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
y1 = [1, 5, 10, 50, 100]
z1 = training_scores

x2 = [40,60,80,100,120]
y2 = [1, 5, 10, 50, 100]
z2 = cv_scores
```

In [61]:

Observation: Using GridsearchCV max_depth is 100 and n_esimators is 120.

In [60]:

```
# https://scikit-
learn.org/stable/modules/generated/sklearn.metrics.roc_curve.html#sklearn.metrics.roc_curve
from sklearn.metrics import roc_curve, auc
clf = RandomForestClassifier(max_depth=100,n_estimators=120)
clf.fit(x_train_avgw2v, y_train)
y_pred = clf.predict(x_test_avgw2v)
# roc_auc_score(y_true, y_score) the 2nd parameter should be probability estimates of the positive
class
```

```
| # not the predicted outputs
train fpr, train tpr, thresholds = roc curve(y train, clf.predict proba(x train avgw2v)[:,1])
test fpr, test tpr, thresholds = roc curve(y test, clf.predict proba(x test avgw2v)[:,1])
plt.plot(train fpr, train tpr, label="train AUC ="+str(auc(train fpr, train tpr)))
plt.plot(test_fpr, test_tpr, label="test AUC ="+str(auc(test_fpr, test_tpr)))
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
print("="*100)
from sklearn.metrics import confusion matrix
print("Train confusion matrix")
print(confusion matrix(y train, clf.predict(x train avgw2v)))
print("Test confusion matrix")
print(confusion_matrix(y_test, clf.predict(x_test_avgw2v)))
```

```
Train confusion matrix
[[10993 24]
[ 0 58983]]
Test confusion matrix
[[ 1599 3053]
[ 458 24890]]
```

In [61]:

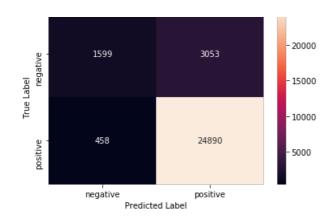
```
# Confusion Matrix
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test, y_pred)
cm
```

Out[61]:

```
array([[ 1599, 3053], [ 458, 24890]])
```

In [62]:

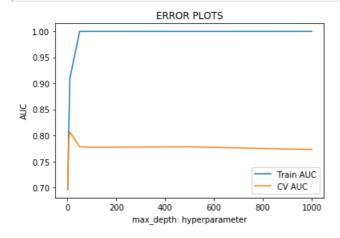
```
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class_label = ["negative", "positive"]
df_cm = pd.DataFrame(cm, index = class_label, columns = class_label)
sns.heatmap(df_cm, annot = True, fmt = "d")
plt.title("Confusion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
```



[5.1.6] Applying Random Forests on TFIDF W2V, SET 4

```
In [62]:
```

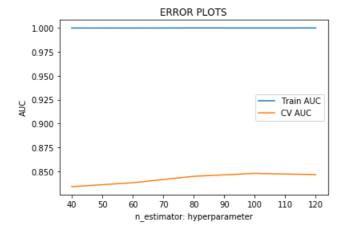
```
## find hyperparameter using roc auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc_auc_score
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]
#empty lists that stores cv scores and training scores
cv scores = []
training_scores = []
#perform k fold cross validation
for d in max depth values:
   clf = RandomForestClassifier (max depth=d)
   clf.fit(x train tfidfw2v, y train)
   y_train_pred = clf.predict_proba(x_train_tfidfw2v)[:,1]
   y_test_pred = clf.predict_proba(x_test_tfidfw2v)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv scores.append(roc auc score(y test, y test pred))
#plot cross-validated score, training score vs alpha
plt.plot(max_depth_values, training_scores, label='Train AUC')
plt.plot(max_depth_values, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("max depth: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [63]:

```
## find hyperparameter using roc_auc score and plot AUC
from sklearn.ensemble import RandomForestClassifier
from sklearn.metrics import roc_auc_score
base_learners = [40,60,80,100,120]
```

```
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for n in base learners:
    clf = RandomForestClassifier(n estimators=n)
   clf.fit(x_train_tfidfw2v, y_train)
   y_train_pred = clf.predict_proba(x_train_tfidfw2v)[:,1]
   y test pred = clf.predict proba(x test tfidfw2v)[:,1]
    training scores.append(roc auc score(y train, y train pred))
    cv scores.append(roc auc score(y test, y test pred))
#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base learners, cv scores, label='CV AUC')
plt.legend()
plt.xlabel("n estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [67]:

```
## Using GridsearchCV
from sklearn.ensemble import RandomForestClassifier
from sklearn.model_selection import GridSearchCV

base_learners = [40,60,80,100,120]
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]

param_grid = {'n_estimators': base_learners , "max_depth":max_depth_values}
RFC = RandomForestClassifier (max_features='sqrt')
model = GridSearchCV (RFC, param_grid,cv=3 ,pre_dispatch=2,scoring='roc_auc')
model.fit(x_train_tfidfw2v, y_train)
```

Out[67]:

In [64]:

```
import plotly.offline as offline
import plotly.graph_objs as go
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
y1 = [1, 5, 10, 50, 100]
z1 = training_scores

x2 = [40,60,80,100,120]
y2 = [1, 5, 10, 50, 100]
z2 = cv_scores
```

In [65]:

Observation: By observing curve getting high AUC at 100 max depth and 120 n estimator.

```
In [52]:
```

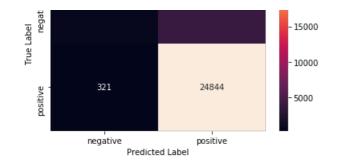
```
# https://scikit-
learn.org/stable/modules/generated/sklearn.metrics.roc_curve.html#sklearn.metrics.roc_curve
from sklearn.metrics import roc_curve, auc
clf = RandomForestClassifier(max_depth=100,n_estimators=120)
clf.fit(x_train_tfidfw2v, y_train)
y_pred = clf.predict(x_test_tfidfw2v)
# roc_auc_score(y_true, y_score) the 2nd parameter should be probability estimates of the positive
class
# not the predicted outputs

train_fpr, train_tpr, thresholds = roc_curve(y_train, clf.predict_proba(x_train_tfidfw2v)[:,1])
test_fpr, test_tpr, thresholds = roc_curve(v_test_clf.predict_proba(x_test_tfidfw2v)[:,1])
```

```
plt.plot(train_fpr, train_tpr, label="train AUC ="+str(auc(train_fpr, train_tpr)))
plt.plot(test_fpr, test_tpr, label="test AUC ="+str(auc(test_fpr, test_tpr)))
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
print("="*100)
from sklearn.metrics import confusion matrix
print("Train confusion matrix")
print(confusion_matrix(y_train, clf.predict(x_train_tfidfw2v)))
print("Test confusion matrix")
print(confusion_matrix(y_test, clf.predict(x_test_tfidfw2v)))
                    ERROR PLOTS
  1.0
  0.8
True Positive Rate
  0.6
  0.4
  0.2
                       train AUC = 0.9998867160582465
                       test AUC =0.8454159322001162
  0.0
                            0.6
                                   0.8
                                          1.0
      0.0
             0.2
                    0.4
                   False Positive Rate
______
Train confusion matrix
[[10782 30]
 [ 0 59188]]
Test confusion matrix
[[ 780 4055]
[ 321 24844]]
In [53]:
# Confusion Matrix
from sklearn.metrics import confusion_matrix
cm = confusion matrix(y test, y pred)
Out[53]:
array([[ 780, 4055],
       [ 321, 24844]])
In [54]:
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class_label = ["negative", "positive"]
df cm = pd.DataFrame(cm, index = class label, columns = class label)
sns.heatmap(df_cm, annot = True, fmt = "d")
plt.title("Confusiion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
```

Confusiion Matrix

4055

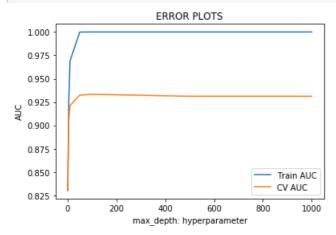


[5.2] Applying GBDT using XGBOOST

[5.2.1] Applying XGBOOST on BOW, SET 1

```
In [80]:
```

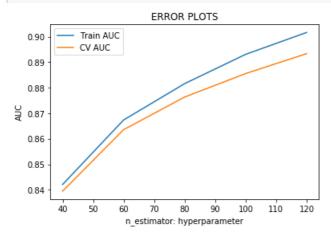
```
## find hyperparameter using roc_auc score and plot AUC
from sklearn.metrics import roc_auc_score
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]
#empty lists that stores cv scores and training_scores
cv scores = []
training_scores = []
#perform k fold cross validation
for d in max depth values:
    clf = XGBClassifier(max depth=d)
    clf.fit(x train bow, y train)
    y train pred = clf.predict proba(x train bow)[:,1]
    y_test_pred = clf.predict_proba(x_test_bow)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(max_depth_values, training_scores, label='Train AUC')
plt.plot(max_depth_values, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("max_depth: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [81]:

```
## find hyperparameter using roc_auc score and plot AUC
from xgboost import XGBClassifier
from sklearn.metrics import roc_auc_score
base_learners = [40,60,80,100,120]
#empty lists that stores cv scores and training_scores
```

```
cv scores = []
training_scores = []
#perform k fold cross validation
for n in base learners:
    clf = XGBClassifier(n estimators=n)
    clf.fit(x train bow, y train)
    y_train_pred = clf.predict_proba(x_train_bow)[:,1]
   y test pred = clf.predict proba(x test bow)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv scores.append(roc auc score(y test, y test pred))
#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base learners, cv scores, label='CV AUC')
plt.legend()
plt.xlabel("n estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [68]:

```
## Using GridsearchCV
from xgboost import XGBClassifier
from sklearn.model_selection import GridSearchCV

base_learners = [40,60,80,100,120]
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]

param_grid = {'n_estimators': base_learners , "max_depth":max_depth_values}
XGB = XGBClassifier(max_features='sqrt')
model = GridSearchCV(XGB, param_grid,cv=3 ,pre_dispatch=2,scoring='roc_auc')
model.fit(x_train_bow, y_train)
```

Out[68]:

In [82]:

```
#### 3D Scatter Plot
import plotly.offline as offline
```

```
import plotly.graph_objs as go
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
y1 = [1, 5, 10, 50, 100]
z1 = training_scores

x2 = [40,60,80,100,120]
y2 = [1, 5, 10, 50, 100]
z2 = cv_scores
```

In [83]:

Observation: By GridsearchCV getting high AUC at 3 max_depth and 100 n_estimator.

In [46]:

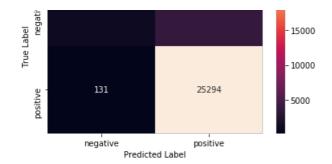
```
# https://scikit-
learn.org/stable/modules/generated/sklearn.metrics.roc_curve.html#sklearn.metrics.roc_curve
from sklearn.metrics import roc_curve, auc
from xgboost import XGBClassifier
clf = XGBClassifier(max_depth=3,n_estimators=100)
clf.fit(x_train_bow, y_train)
y_pred = clf.predict(x_test_bow)
# roc_auc_score(y_true, y_score) the 2nd parameter should be probability estimates of the positive
class
# not the predicted outputs

train_fpr, train_tpr, thresholds = roc_curve(y_train, clf.predict_proba(x_train_bow)[:,1])
test_fpr_test_tpr_thresholds = roc_curve(y_train, clf.predict_proba(x_test_bow)[:,1])
```

```
plt.plot(train_fpr, train_tpr, label="train AUC ="+str(auc(train_fpr, train_tpr)))
plt.plot(test_fpr, test_tpr, label="test AUC ="+str(auc(test fpr, test tpr)))
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
print("="*100)
from sklearn.metrics import confusion matrix
print("Train confusion matrix")
print(confusion matrix(y_train, clf.predict(x_train_bow)))
print("Test confusion matrix")
print(confusion_matrix(y_test, clf.predict(x_test_bow)))
                     ERROR PLOTS
  1.0
  0.8
True Positive Rate
  0.6
  0.4
  0.2
                        train AUC = 0.8957094517467553
                         test AUC = 0.8763261451499375
  0.0
              0.2
                             0.6
                                     0.8
                    False Positive Rate
Train confusion matrix
[[ 2558 8364]
[ 230 58848]]
Test confusion matrix
[[ 990 3585]
 [ 131 25294]]
4
In [47]:
# Confusion Matrix
from sklearn.metrics import confusion matrix
cm = confusion matrix(y test, y pred)
Out[47]:
In [48]:
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class label = ["negative", "positive"]
df_cm = pd.DataFrame(cm, index = class_label, columns = class_label)
sns.heatmap(df cm, annot = True, fmt = "d")
plt.title("Confusiion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
               Confusiion Matrix
```

20000

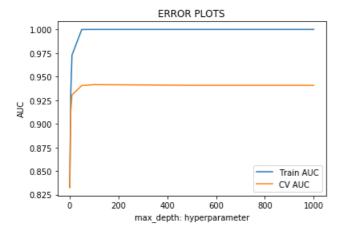
cest_tpr, test_tpr, threshords - roc_curve(y_test, trr.predict_proba(x_test_bow)[.,i])



[5.2.2] Applying XGBOOST on TFIDF, SET 2

In [84]:

```
## find hyperparameter using roc_auc score and plot AUC
from xgboost import XGBClassifier
from sklearn.metrics import roc_auc_score
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for d in max depth values:
   clf = XGBClassifier(max_depth=d)
   clf.fit(x train tfidf, y train)
    y_train_pred = clf.predict_proba(x_train_tfidf)[:,1]
   y_test_pred = clf.predict_proba(x_test_tfidf)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(max depth values, training scores, label='Train AUC')
plt.plot(max_depth_values, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("max_depth: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```

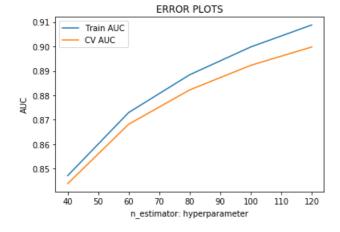


In [85]:

```
## find hyperparameter using roc_auc score and plot AUC
from xgboost import XGBClassifier
from sklearn.metrics import roc_auc_score
base_learners = [40,60,80,100,120]

#empty lists that stores cv scores and training_scores
cv_scores = []
training_scores = []
```

```
#perform k fold cross validation
for n in base_learners:
   clf = XGBClassifier(n_estimators=n)
   clf.fit(x train tfidf, y train)
    y_train_pred = clf.predict_proba(x_train_tfidf)[:,1]
   y test pred = clf.predict proba(x test tfidf)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base learners, cv scores, label='CV AUC')
plt.legend()
plt.xlabel("n estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [49]:

```
## Using GridsearchCV
 from xgboost import XGBClassifier
 from sklearn.model selection import GridSearchCV
base learners = [40,60,80,100,120]
max depth values = [1, 5, 10, 50, 100, 500, 1000]
 param_grid = {'n_estimators': base_learners , "max_depth":max_depth_values}
 XGB = XGBClassifier(max features='sqrt')
model = GridSearchCV(XGB, param_grid,cv=3 ,pre_dispatch=2,scoring='roc_auc')
model.fit(x_train_tfidf, y_train)
Out[49]:
GridSearchCV(cv=3, error score='raise-deprecating',
                    \verb|estimator=XGBClassifier(base\_score=0.5, booster='gbtree', colsample\_bylevel=1, booster='gbtree', colsample\_bylevel=1, booster='gbtree', booster='gbtree'
                    colsample bytree=1, gamma=0, learning rate=0.1, max delta step=0,
                    max depth=3, max features='sqrt', min child weight=1, missing=None,
                    n estimators=100, n jobs=1, nthread=None,
                    objective='binary:logistic', random state=0, reg alpha=0,
                    reg_lambda=1, scale_pos_weight=1, seed=None, silent=True,
                    subsample=1),
                    fit params=None, iid='warn', n jobs=None,
                    param_grid={'n_estimators': [40, 60, 80, 100, 120], 'max_depth': [1, 5, 10, 50, 100, 500, 1
000]},
                    pre_dispatch=2, refit=True, return_train_score='warn',
                    scoring='roc auc', verbose=0)
```

Observation: By Observing Curve, getting high AUC at 3 max_depth and 100 n_estimator.

```
In [86]:
```

```
#### 3D Scatter Plot
import plotly.offline as offline
import plotly.graph_objs as go
```

```
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
y1 = [1, 5, 10, 50, 100]
z1 = training_scores

x2 = [40,60,80,100,120]
y2 = [1, 5, 10, 50, 100]
z2 = cv_scores
```

In [87]:

In [56]:

```
# https://scikit-
learn.org/stable/modules/generated/sklearn.metrics.roc_curve.html#sklearn.metrics.roc_curve
from sklearn.metrics import roc_curve, auc
clf = XGBClassifier(max_depth=3,n_estimators=100)
clf.fit(x_train_tfidf, y_train)
y_pred = clf.predict(x_test_tfidf)
# roc_auc_score(y_true, y_score) the 2nd parameter should be probability estimates of the positive
class
# not the predicted outputs

train_fpr, train_tpr, thresholds = roc_curve(y_train, clf.predict_proba(x_train_tfidf)[:,1])
test_fpr, test_tpr, thresholds = roc_curve(y_test, clf.predict_proba(x_test_tfidf)[:,1])
plt.plot(train_fpr, train_tpr, label="train AUC ="+str(auc(train_fpr, train_tpr)))
plt.plot(test_fpr, test_tpr, label="test AUC ="+str(auc(test_fpr, test_tpr)))
```

```
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
print("="*100)
from sklearn.metrics import confusion matrix
print("Train confusion matrix")
print(confusion_matrix(y_train, clf.predict(x_train_tfidf)))
print("Test confusion matrix")
print(confusion_matrix(y_test, clf.predict(x_test_tfidf)))
                    ERROR PLOTS
  1.0
  0.8
True Positive Rate
  0.6
  0.4
  0.2
                       train AUC =0.9025589908019453
                     test AUC =0.8843992499099999
  0.0
      0.0
             0.2
                            0.6
                                   0.8
                   False Positive Rate
______
Train confusion matrix
[[ 2940 7982]
[ 232 58846]]
Test confusion matrix
[[ 1145 3430]
   129 25296]]
In [57]:
# Confusion Matrix
from sklearn.metrics import confusion matrix
cm = confusion matrix(y test, y pred)
cm
Out[57]:
array([[ 1145, 3430], [ 129, 25296]])
In [58]:
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class label = ["negative", "positive"]
df_cm = pd.DataFrame(cm, index = class_label, columns = class_label)
sns.heatmap(df_cm, annot = True, fmt = "d")
plt.title("Confusiion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
              Confusiion Matrix
                                         25000
                                         - 20000
           1145
                           3430
```

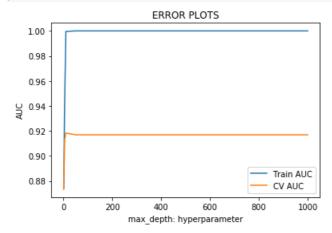
15000



[5.2.3] Applying XGBOOST on AVG W2V, SET 3

In [88]:

```
## find hyperparameter using roc auc score and plot AUC
from xgboost import XGBClassifier
from sklearn.metrics import roc_auc_score
max depth values = [1, 5, 10, 50, 100, 500, 1000]
#empty lists that stores cv scores and training scores
cv scores = []
training scores = []
#perform k fold cross validation
for d in max depth values:
   clf = XGBClassifier(max depth=d)
   clf.fit(x_train_avgw2v, y_train)
   y_train_pred = clf.predict_proba(x_train_avgw2v)[:,1]
   y test pred = clf.predict proba(x test avgw2v)[:,1]
    training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv scores.append(roc auc score(y test, y test pred))
#plot cross-validated score, training score vs alpha
plt.plot(max_depth_values, training_scores, label='Train AUC')
plt.plot(max depth values, cv scores, label='CV AUC')
plt.legend()
plt.xlabel("max_depth: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [89]:

```
### find hyperparameter using roc_auc score and plot AUC
from xgboost import XGBClassifier
from sklearn.metrics import roc_auc_score
base_learners = [40,60,80,100,120]

#empty lists that stores cv scores and training_scores
cv_scores = []
training_scores = []

#perform k fold cross validation
for n in base_learners:
    clf = XGBClassifier(n estimators=n)
```

```
clf.fit(x_train_avgw2v, y_train)
   y_train_pred = clf.predict_proba(x_train_avgw2v)[:,1]
   y_test_pred = clf.predict_proba(x_test_avgw2v)[:,1]

   training_scores.append(roc_auc_score(y_train,y_train_pred))
   cv_scores.append(roc_auc_score(y_test, y_test_pred))

#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base_learners, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("n_estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR_PLOTS")
plt.show()
```

ERROR PLOTS Train AUC 0.915 CV AUC 0.910 0.905 ₽ 0.900 0.895 0.890 0.885 0.880 40 50 70 80 90 100 110 120 n_estimator: hyperparameter

In [69]:

```
## Using GridsearchCV
from xgboost import XGBClassifier
from sklearn.model_selection import GridSearchCV

base_learners = [40,60,80,100,120]
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]

param_grid = {'n_estimators': base_learners , "max_depth":max_depth_values}
RFC = XGBClassifier(max_features='sqrt')
model = GridSearchCV(XGB, param_grid,cv=3 ,pre_dispatch=2,scoring='roc_auc')
model.fit(x_train_avgw2v, y_train)
```

Out[69]:

In [90]:

```
#### 3D Scatter Plot
import plotly.offline as offline
import plotly.graph_objs as go
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
v1 = [1. 5. 10. 50. 100]
```

```
z1 = training_scores

x2 = [40,60,80,100,120]

y2 = [1, 5, 10, 50, 100]

z2 = cv_scores
```

In [91]:

Observation: By observing curve getting high AUC at 3 max_depth and 100 n_estimator.

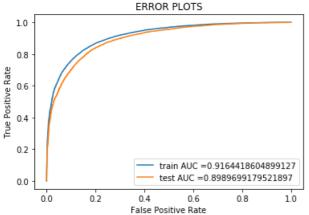
In [53]:

```
# https://scikit-
learn.org/stable/modules/generated/sklearn.metrics.roc curve.html#sklearn.metrics.roc curve
from sklearn.metrics import roc_curve, auc
clf = XGBClassifier(max depth=3,n estimators=100)
clf.fit(x_train_avgw2v, y_train)
y_pred = clf.predict(x_test_avgw2v)
# roc auc score(y true, y score) the 2nd parameter should be probability estimates of the positive
class
# not the predicted outputs
train_fpr, train_tpr, thresholds = roc_curve(y_train, clf.predict_proba(x_train_avgw2v)[:,1])
test fpr, test tpr, thresholds = roc curve(y test, clf.predict proba(x test avgw2v)[:,1])
plt.plot(train_fpr, train_tpr, label="train AUC ="+str(auc(train_fpr, train_tpr)))
plt.plot(test fpr, test tpr, label="test AUC ="+str(auc(test fpr, test tpr)))
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
```

```
plt.title("ERROR PLOTS")
plt.show()

print("="*100)

from sklearn.metrics import confusion_matrix
print("Train confusion matrix")
print(confusion_matrix(y_train, clf.predict(x_train_avgw2v)))
print("Test confusion matrix")
print(confusion_matrix(y_test, clf.predict(x_test_avgw2v)))
```



```
Train confusion matrix
[[ 4798 6124]
  [ 1331 57747]]
Test confusion matrix
[[ 1844 2731]
  [ 663 24762]]
```

In [54]:

```
# Confusion Matrix
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test, y_pred)
cm
```

Out[54]:

```
array([[ 1844, 2731], [ 663, 24762]])
```

In [55]:

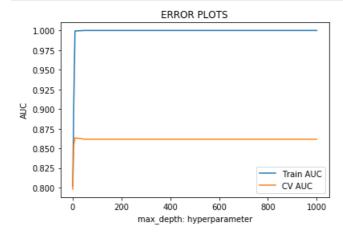
```
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class_label = ["negative", "positive"]
df_cm = pd.DataFrame(cm, index = class_label, columns = class_label)
sns.heatmap(df_cm, annot = True, fmt = "d")
plt.title("Confusiion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
```



[5.2.4] Applying XGBOOST on TFIDF W2V, SET 4

```
In [92]:
```

```
## find hyperparameter using roc_auc score and plot AUC
from xgboost import XGBClassifier
from sklearn.metrics import roc auc score
max depth values = [1, 5, 10, 50, 100, 500, 1000]
#empty lists that stores cv scores and training scores
cv scores = []
training_scores = []
#perform k fold cross validation
for d in max depth values:
    clf = XGBClassifier(max depth=d)
    clf.fit(x_train_tfidfw2v, y_train)
    y train pred = clf.predict proba(x train tfidfw2v)[:,1]
    y test_pred = clf.predict_proba(x_test_tfidfw2v)[:,1]
    training scores.append(roc auc score(y train,y train pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))
#plot cross-validated score, training score vs alpha
plt.plot(max depth values, training scores, label='Train AUC')
plt.plot(max_depth_values, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("max depth: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In [93]:

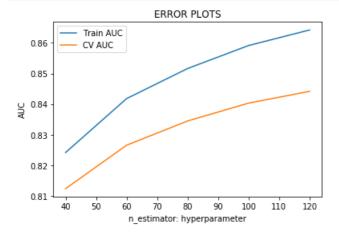
```
## find hyperparameter using roc_auc score and plot AUC
from xgboost import XGBClassifier
from sklearn.metrics import roc_auc_score
base_learners = [40,60,80,100,120]

#empty lists that stores cv scores and training_scores
cv_scores = []
training_scores = []

#perform k fold cross validation
for n in base_learners:
    clf = XGBClassifier(n_estimators=n)
    clf.fit(x_train_tfidfw2v, y_train)
    y_train_pred = clf.predict_proba(x_train_tfidfw2v)[:,1]
    y_test_pred = clf.predict_proba(x_test_tfidfw2v)[:,1]
```

```
training_scores.append(roc_auc_score(y_train,y_train_pred))
    cv_scores.append(roc_auc_score(y_test, y_test_pred))

#plot cross-validated score, training score vs alpha
plt.plot(base_learners, training_scores, label='Train AUC')
plt.plot(base_learners, cv_scores, label='CV AUC')
plt.legend()
plt.xlabel("n_estimator: hyperparameter")
plt.ylabel("AUC")
plt.title("ERROR PLOTS")
plt.show()
```



In []:

```
## Using GridsearchCV
from xgboost import XGBClassifier
from sklearn.model_selection import GridSearchCV

base_learners = [40,60,80,100,120]
max_depth_values = [1, 5, 10, 50, 100, 500, 1000]

param_grid = {'n_estimators': base_learners , "max_depth":max_depth_values}
RFC = XGBClassifier(max_features='sqrt')
model = GridSearchCV(XGB, param_grid,cv=3 ,pre_dispatch=2,scoring='roc_auc')
model.fit(x_train_tfidfw2v, y_train)
Out[]:
```


In [94]:

```
#### 3D Scatter Plot
import plotly.offline as offline
import plotly.graph_objs as go
offline.init_notebook_mode()
import numpy as np

x1 = [40,60,80,100,120]
y1 = [1, 5, 10, 50, 100]
z1 = training_scores

x2 = [40,60,80,100,120]
```

```
y2 = [1, 5, 10, 50, 100]
z2 = cv_scores
```

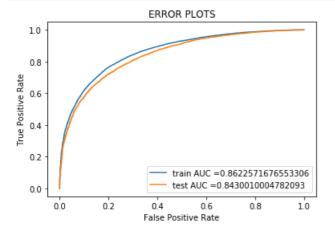
In [95]:

Observation: By observing curve getting high AUC at 3 max_depth and 100 n_estimator.

In [50]:

```
# https://scikit-
learn.org/stable/modules/generated/sklearn.metrics.roc curve.html#sklearn.metrics.roc curve
from sklearn.metrics import roc curve, auc
clf = XGBClassifier(max_depth=3,n_estimators=100)
clf.fit(x_train_tfidfw2v, y_train)
y pred = clf.predict(x test tfidfw2v)
\# roc_auc_score(y_true, y_score) the 2nd parameter should be probability estimates of the positive
class
# not the predicted outputs
train fpr, train tpr, thresholds = roc curve(y train, clf.predict proba(x train tfidfw2v)[:,1])
test_fpr, test_tpr, thresholds = roc_curve(y_test, clf.predict_proba(x_test_tfidfw2v)[:,1])
plt.plot(train fpr, train tpr, label="train AUC ="+str(auc(train fpr, train tpr)))
plt.plot(test_fpr, test_tpr, label="test AUC ="+str(auc(test_fpr, test tpr)))
plt.legend()
plt.xlabel("False Positive Rate")
plt.ylabel("True Positive Rate")
plt.title("ERROR PLOTS")
plt.show()
print("="*100)
```

```
from sklearn.metrics import confusion_matrix
print("Train confusion matrix")
print(confusion_matrix(y_train, clf.predict(x_train_tfidfw2v)))
print("Test confusion matrix")
print(confusion_matrix(y_test, clf.predict(x_test_tfidfw2v)))
```



```
Train confusion matrix
[[ 2579 8343]
[ 918 58160]]
Test confusion matrix
[[ 987 3588]
[ 429 24996]]
```

In [51]:

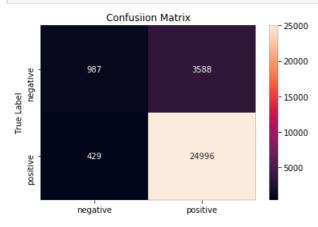
```
# Confusion Matrix
from sklearn.metrics import confusion_matrix
cm = confusion_matrix(y_test, y_pred)
cm
```

Out[51]:

```
array([[ 987, 3588], [ 429, 24996]])
```

In [52]:

```
# plot confusion matrix to describe the performance of classifier.
import seaborn as sns
class_label = ["negative", "positive"]
df_cm = pd.DataFrame(cm, index = class_label, columns = class_label)
sns.heatmap(df_cm, annot = True, fmt = "d")
plt.title("Confusiion Matrix")
plt.xlabel("Predicted Label")
plt.ylabel("True Label")
plt.show()
```



[6] Conclusions

In []:

```
# Please compare all your models using Prettytable library
```

In [55]

```
from prettytable import PrettyTable

x = PrettyTable()

x.field_names = ["Vectorizer","MOdel","max_depth","n_estimator","AUC"]

x.add_row(["BOW","RF","1000","120","92.83%"])

x.add_row(["TFIDF","RF","500","40","92.93%"])

x.add_row(["AvgW2V","RF","100","120","89.89%"])

x.add_row(["TFIDF W2V","RF","100","120","84.54%"])

x.add_row(["BOW","XGB","3","100","87.63%"])

x.add_row(["TFIDF","XGB","3","100","88.43%"])

x.add_row(["AvgW2V","XGB","3","100","89.89%"])

x.add_row(["TFIDF W2V","XGB","3","100","84.30%"])

print(x)
```

+	Vectorizer	+ - 	MOdel	+- +-	max_depth	+- +-	n_estimator	+-	AUC
	BOW	 	RF	т- 	1000	- 	120		92.83%
	TFIDF		RF		500		40		92.93%
	AvgW2V		RF		100		120		89.89%
	TFIDF W2V		RF		100		120		84.54%
	BOW		XGB		3		100		87.63%
	TFIDF		XGB		3		100		88.43%
	AvgW2V		XGB		3		100		89.89%
	TFIDF W2V		XGB		3		100		84.30%
+		+-		+-		+-		+-	+

1) RandomForest with TFIDF getting highewr accuracy. 2) GridsearcvCV takes a long time to train. 3) In XGBoost classifier geting same hyperparameters for all 4 featurization techniques. 4) In both RandomForest and XGBoost classifier, with AVGW2v and TFIDFW2V getting same accuracy. 5) In RandomForest Classifier with TFIDFW2V, Hyperparameters are not getting with GridseachCV.