Amazon Fine Food Reviews Analysis

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

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

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

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

Number of Attributes/Columns in data: 10

Attribute Information:

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

Objective:

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

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

[Ans] We could use 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 will be set to "positive". Otherwise, it will be set to "negative".

```
%matplotlib inline
In [1]:
        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
```

```
# 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 computi
ng power
# filtered data = pd.read sql query(""" SELECT * FROM Reviews WHERE S
core != 3 LIMIT 500000"", con)
# for tsne assignment you can take 5k data points
filtered data = pd.read sql query(""" SELECT * FROM Reviews WHERE Sco
re != 3 LIMIT 100000""", con)
# Give reviews with Score>3 a positive rating(1), and reviews with a
 score<3 a negative rating(0).</pre>
def partition(x):
    if x < 3:
        return 0
    return 1
#changing reviews with score less than 3 to be positive and vice-vers
actualScore = filtered data['Score']
positiveNegative = actualScore.map(partition)
filtered data['Score'] = positiveNegative
print("Number of data points in our data", filtered data.shape)
filtered data.head(3)
```

Number of data points in our data (100000, 10)

Out[2]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	HelpfulnessDenor
0	1	B001E4KFG0	A3SGXH7AUHU8GW	delmartian	1	
1	2	B00813GRG4	A1D87F6ZCVE5NK	dll pa	0	
2	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1	
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]:

	UserId	ProductId	ProfileName	Time	Score	Text	COUNT(*)
0	#oc- R115TNMSPFT9I7	B005ZBZLT4	Breyton	1331510400	2	Overall its just OK when considering the price	2
1	#oc- R11D9D7SHXIJB9	B005HG9ESG	Louis E. Emory "hoppy"	1342396800	5	My wife has recurring extreme muscle spasms, u	3
2	#oc- R11DNU2NBKQ23Z	B005ZBZLT4	Kim Cieszykowski	1348531200	1	This coffee is horrible and unfortunately not	2
3	#oc- R11O5J5ZVQE25C	B005HG9ESG	Penguin Chick	1346889600	5	This will be the bottle that you grab from the	3
4	#oc- R12KPBODL2B5ZD	B007OSBEV0	Christopher P. Presta	1348617600	1	I didnt like this coffee. Instead of telling y	2

In [5]: | display[display['UserId']=='AZY10LLTJ71NX']

Out[5]:

	UserId	ProductId	ProfileName	Time	Score	Text	COUNT(*)
80638	AZY10LLTJ71NX	B001ATMQK2	undertheshrine "undertheshrine"	1296691200	5	I bought this 6 pack because for the price tha	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("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND UserId="AR5J8UI46CURR"
    ORDER BY ProductID
    """, con)
    display.head()
```

Out[7]:

	ld	ProductId	UserId	ProfileName	HelpfulnessNumerator	HelpfulnessDen
() 78445	B000HDL1RQ	AR5J8UI46CURR	Geetha Krishnan	2	
1	L 138317	B000HDOPYC	AR5J8UI46CURR	Geetha Krishnan	2	
2	2 138277	B000HDOPYM	AR5J8UI46CURR	Geetha Krishnan	2	
3	3 73791	B000HDOPZG	AR5J8UI46CURR	Geetha Krishnan	2	
4	1 155049	B000PAQ75C	AR5J8UI46CURR	Geetha Krishnan	2	
4						>

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.

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

```
display= pd.read_sql_query("""
In [11]:
          SELECT *
          FROM Reviews
          WHERE Score != 3 AND Id=44737 OR Id=64422
          ORDER BY ProductID
          """, con)
          display.head()
Out[11]:
                ld
                      ProductId
                                       UserId ProfileName HelpfulnessNumerator HelpfulnessDen
                                                   J. E.
          0 64422 B000MIDROQ A161DK06JJMCYF
                                                Stephens
                                                                        3
                                                "Jeanne"
          1 44737 B001EQ55RW A2V0I904FH7ABY
                                                                        3
                                                   Ram
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()
          (87773, 10)
Out[13]: 1
               73592
               14181
          Name: Score, dtype: int64
```

[3] Preprocessing

[3.1]. Preprocessing Review Text

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

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

- 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("="*50)

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

My dogs loves this chicken but its a product from China, so we wont be buying it anymore. Its very hard to find any chicken products made in the USA but they are out there, but this one isnt. Its too bad to o because its a good product but I wont take any chances till they kn ow what is going on with the china imports.

The Candy Blocks were a nice visual for the Lego Birthday party but the candy has little taste to it. Very little of the 2 lbs that I bought were eaten and I threw the rest away. I would not buy the candy again.

was way to hot for my blood, took a bite and did a jig lol

My dog LOVES these treats. They tend to have a very strong fish oil s mell. So if you are afraid of the fishy smell, don't get it. But I th ink my dog likes it because of the smell. These treats are really small in size. They are great for training. You can give your dog several of these without worrying about him over eating. Amazon's price was much more reasonable than any other retailer. You can buy a 1 pound bag on Amazon for almost the same price as a 6 ounce bag at other retailers. It's definitely worth it to buy a big bag if your dog eats the malot.

My dogs loves this chicken but its a product from China, so we wont be buying it anymore. Its very hard to find any chicken products made in the USA but they are out there, but this one isnt. Its too bad to o because its a good product but I wont take any chances till they kn ow what is going on with the china imports.

https://stackoverflow.com/questions/16206380/python-beautifulsoup-h ow-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)

My dogs loves this chicken but its a product from China, so we wont be buying it anymore. Its very hard to find any chicken products made in the USA but they are out there, but this one isnt. Its too bad to o because its a good product but I wont take any chances till they kn ow what is going on with the china imports.

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

```
In [18]: sent_1500 = decontracted(sent_1500)
    print(sent_1500)
    print("="*50)
```

was way to hot for my blood, took a bite and did a jig lol

```
In [19]: #remove words with numbers python: https://stackoverflow.com/a/180823
70/4084039
sent_0 = re.sub("\S*\d\S*", "", sent_0).strip()
print(sent_0)
```

My dogs loves this chicken but its a product from China, so we wont be buying it anymore. Its very hard to find any chicken products made in the USA but they are out there, but this one isnt. Its too bad to o because its a good product but I wont take any chances till they kn ow what is going on with the china imports.

```
In [20]: #remove spacial character: https://stackoverflow.com/a/5843547/408403
g
sent_1500 = re.sub('[^A-Za-z0-9]+', ' ', sent_1500)
print(sent_1500)
```

was way to hot for my blood took a bite and did a jig lol

```
In [21]: | # https://gist.github.com/sebleier/554280
      # we are removing the words from the stop words list: 'no', 'nor', 'n
      ot'
      # <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 i
      n the 1st step
      'its', 'itself', 'they', 'them', 'their',\
               'theirs', 'themselves', 'what', 'which', 'who', 'whom',
       'this', 'that', "that'll", 'these', 'those', \
      'then', 'once', 'here', 'there', 'when', 'where', 'why',
       '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', "i
  'mightn', "mightn't", 'mustn',\
               "mustn't", 'needn', "needn't", 'shan', "shan't", 'should
      n', "shouldn't", 'wasn', "wasn't", 'weren', "weren't", \
               'won', "won't", 'wouldn', "wouldn't"])
```

```
In [22]: # 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
        sentance = ' '.join(e.lower() for e in sentance.split() if e.lowe
    r() not in stopwords)
        preprocessed_reviews.append(sentance.strip())
```

100%| 87773/87773 [00:45<00:00, 1942.43it/s]

```
In [23]: preprocessed_reviews[1500]
Out[23]: 'way hot blood took bite jig lol'
```

[3.2] Preprocessing Review Summary

```
In [24]: ## Similartly you can do preprocessing for review summary also.
```

[4] Featurization

Processing Texts from summary

```
In [25]: preprocessed_reviews_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.lowe
r() not in stopwords)
    preprocessed_reviews_summary.append(sentance.strip())

100%[ 87773/87773 [00:28<00:00, 3083.28it/s]</pre>
```

Including Texts from summary in the main text

Adding length of text as a feature

```
In [27]: preprocessed_reviews_fe_final = []
    for i in tqdm(preprocessed_reviews_fe):
        count = 0;
        for j in i.split(" "):
            count += 1;
        i = i + " " + str(count);
        preprocessed_reviews_fe_final.append(i)
100% | 87773/87773 [00:01<00:00, 84100.63it/s]
```

Test - Train Split

```
In [28]: from sklearn.model_selection import train_test_split

X_train, X_test, y_train, y_test = train_test_split(preprocessed_revi
ews_fe_final, final['Score'], test_size=0.33) # this is random splitt
ing
X_train, X_cv, y_train, y_cv = train_test_split(X_train, y_train, test_size=0.33) # this is random splitting
```

[4.1] BAG OF WORDS

```
In [29]: count_vect = CountVectorizer( min_df = 10) #in scikit-learn

X_train_vect = count_vect.fit_transform(X_train)
# X_train_vect = X_train_vect.toarray()
X_cv_vect = count_vect.transform(X_cv)
# X_cv_vect = X_cv_vect.toarray()
X_test_vect = Count_vect.transform(X_test)
# X_test_vect = X_test_vect.toarray()
```

[4.2] Bi-Garams and n-Grams.

[4.3] TF-IDF

```
In [39]: tf_idf = TfidfVectorizer(min_df = 10)
    X_train_vect_tfidf = tf_idf.fit_transform(X_train)
    # X_train_vect_tfidf = X_train_vect_tfidf.toarray()
    X_test_vect_tfidf = tf_idf.transform(X_test)
    # X_test_vect_tfidf = X_test_vect_tfidf.toarray()
    X_cv_vect_tfidf = tf_idf.transform(X_cv)
    # X_cv_vect_tfidf = X_cv_vect_tfidf.toarray()
```

[4.4] Word2Vec

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

[4.4.1.1] Avg W2v

```
In [40]:
         # average Word2Vec
         # compute average word2vec for each review.
         i=0
         list of sent=[]
         X_train_list=[]
         X test list=[]
         X_cv_list=[]
         for sent in X train:
              X train list.append(sent.split())
         for sent in X cv:
              X_cv_list.append(sent.split())
         for sent in X test:
              X test list.append(sent.split())
         w2v model=Word2Vec(X train list,min count=0,size=50, workers=4)
         w2v words = list(w2v model.wv.vocab)
         X train vectors = [];
         for sent in tqdm(X train list):
              sent_vec = np.zeros(50)
              cnt words =0;
              for word in sent:
                  if word in w2v_words:
                      vec = w2v_model.wv[word]
                      sent vec += vec
                      cnt words += 1
              if cnt words != 0:
                  sent vec /= cnt words
              X train vectors.append(sent vec)
         X cv vectors = []
         for sent in tqdm(X_cv_list):
              sent_vec = np.zeros(50)
              cnt words =0;
              for word in sent:
                  if word in w2v words:
                      vec = w2v model.wv[word]
                      sent_vec += vec
                      cnt words += 1
              if cnt_words != 0:
                  sent vec /= cnt words
              X cv vectors.append(sent vec)
         X_test_vectors = []
         for sent in tqdm(X_test_list):
              sent vec = np.zeros(50)
              cnt words =0;
              for word in sent:
                  if word in w2v_words:
                      vec = w2v_model.wv[word]
                      sent_vec += vec
                      cnt words += 1
              if cnt words != 0:
```

```
sent_vec /= cnt_words
X_test_vectors.append(sent_vec)
```

```
100%| 39400/39400 [02:28<00:00, 265.34it/s]
100%| 19407/19407 [01:20<00:00, 240.40it/s]
100%| 28966/28966 [02:01<00:00, 239.24it/s]
```

[4.4.1.2] TFIDF weighted W2v

```
In [55]:
         dictionary = dict(zip(tf idf.get feature names(), list(tf idf.idf )))
         tfidf_feat = tf_idf.get_feature_names()
         tfidf X train vectors = [];
         tfidf X test vectors = [];
         tfidf X cv vectors = [];
         for sent in tqdm(X_train_list):
             sent vec = np.zeros(50)
             weight sum =0;
              for word in sent:
                  if word in (w2v words and tfidf feat):
                      vec = w2v model.wv[word]
                      tf idf count = dictionary[word]*sent.count(word)
                      sent_vec += (vec * tf_idf_count)
                      weight sum += tf idf count
             if weight_sum != 0:
                  sent vec /= weight sum
             tfidf_X_train_vectors.append(sent_vec)
         for sent in tqdm(X test list):
             sent vec = np.zeros(50)
             weight sum =0;
             for word in sent:
                  if word in (w2v_words and tfidf_feat):
                      vec = w2v model.wv[word]
                      tf idf count = dictionary[word]*sent.count(word)
                      sent vec += (vec * tf idf count)
                      weight sum += tf idf count
             if weight sum != 0:
                  sent vec /= weight sum
              tfidf X test vectors.append(sent vec)
         for sent in tqdm(X cv list):
             sent vec = np.zeros(50)
             weight sum =0;
              for word in sent:
                  if word in (w2v words and tfidf feat):
                      vec = w2v model.wv[word]
                      tf idf count = dictionary[word]*sent.count(word)
                      sent vec += (vec * tf idf count)
                      weight_sum += tf_idf_count
             if weight sum != 0:
                  sent_vec /= weight sum
             tfidf X cv vectors.append(sent vec)
```

```
100%| 39400/39400 [03:07<00:00, 210.00it/s]
100%| 28966/28966 [02:17<00:00, 210.75it/s]
100%| 19407/19407 [01:29<00:00, 218.00it/s]
```

[5] Assignment 8: Decision Trees

1. Apply Decision Trees 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 (best `depth` in range [1, 5, 10, 50, 100, 500, 100], and the best `min_samples_split` in range [5, 10, 100, 500])

- Find the best hyper parameter which will give the maximum <u>AUC</u>
 (https://www.appliedaicourse.com/course/applied-ai-course-online/lessons/receiver-operating-characteristic-curve-roc-curve-and-auc-1/) 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. Graphviz

- Visualize your decision tree with Graphviz. It helps you to understand how a decision is being made, given a new vector.
- Since feature names are not obtained from word2vec related models, visualize only BOW & TFIDF decision trees using Graphviz
- Make sure to print the words in each node of the decision tree instead of printing its index.
- Just for visualization purpose, limit max_depth to 2 or 3 and either embed the generated images of graphviz in your notebook, or directly upload them as .png files.

4. Feature importance

Find the top 20 important features from both feature sets Set 1 and Set 2 using `feature_importances_`
method of <u>Decision Tree Classifier (https://scikit-learn.org/stable/modules/generated/sklearn.tree.DecisionTreeClassifier.html)</u> and print their corresponding feature names

5. 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.

6. 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.

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.

 <u>tnr-1/)</u> with predicted and original labels of test data points. Please visualize your confusion matrices using <u>seaborn heatmaps</u>.

(https://seaborn.pydata.org/generated/seaborn.heatmap.html)

(https://seaborn.pydata.org/generated/seaborn.heatmap.html)

(https://seaborn.pydata.org/generated/seaborn.heatmap.html)

(https://seaborn.pydata.org/generated/seaborn.heatmap.html)

7. Conclusion (https://seaborn.pydata.org/generated/seaborn.heatmap.html)

(https://seaborn.pydata.org/generated/seaborn.heatmap.html)

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

(https://seaborn.pydata.org/generated/seaborn.heatmap.html) link

(http://zetcode.com/python/prettytable/)



Note: Data Leakage

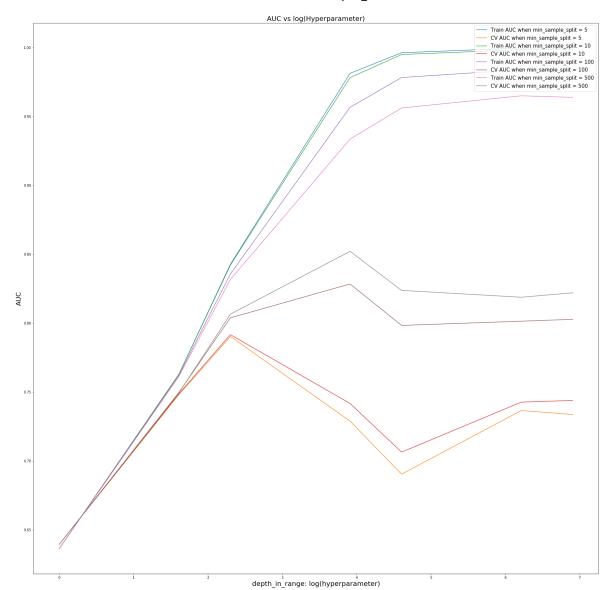
- 1. There will be an issue of data-leakage if you vectorize the entire data and then split it into train/cv/test.
- 2. To avoid the issue of data-leakag, make sure to split your data first and then vectorize it.
- 3. While vectorizing your data, apply the method fit_transform() on you train data, and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

Applying Decision Trees

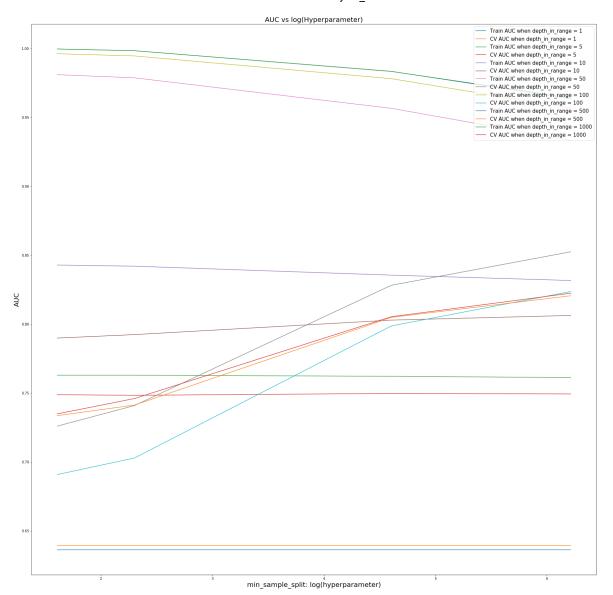
[5.1] Applying Decision Trees on BOW, SET 1

```
from sklearn.tree import DecisionTreeClassifier
In [31]:
         from sklearn.model selection import GridSearchCV
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         parameters = [{'max_depth': [1,5,10,50,100,500,1000]},{'min samples s
         plit':[5,10,100,500]}]
         model = GridSearchCV(DecisionTreeClassifier(),parameters, scoring =
         'roc auc')
         model.fit(X_train_vect, y_train)
         print(model.best estimator )
         print(model.score(X_test_vect, y_test))
         DecisionTreeClassifier(class weight=None, criterion='gini', max depth
         =None,
                     max features=None, max leaf nodes=None,
                     min impurity decrease=0.0, min impurity split=None,
                     min_samples_leaf=1, min_samples_split=500,
                     min weight fraction leaf=0.0, presort=False, random state
         =None,
                     splitter='best')
         0.8222850251641352
```

```
from sklearn.metrics import accuracy score
In [32]:
         from sklearn.metrics import roc auc score
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         ind = []
         train_auc = []
         cv auc = []
         plt.figure(figsize=(30,30))
         for i in (min sample split):
             cv auc plot = []
             train auc plot = []
             for j in (depth in range):
                 dtc = DecisionTreeClassifier(max depth = j,min samples split
         = i)
                 dtc.fit(X train vect, y train)
                 y train pred = dtc.predict proba(X train vect)[:,1]
                 y_cv_pred = dtc.predict_proba(X_cv_vect)[:,1]
                 train auc.append(roc auc score(y train,y_train_pred))
                  train auc plot.append(roc auc score(y train,y train pred))
                  cv_auc.append(roc_auc_score(y_cv, y_cv_pred))
                  cv auc plot.append(roc auc score(y cv, y cv pred))
             plt.plot(np.log(depth in range), train auc plot, label='Train AUC
         when min sample split = ' + str(i))
             plt.plot(np.log(depth in range), cv auc plot, label='CV AUC when
          min sample split = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("depth in range: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```



```
depth in range = [1,5,10,50,100,500,1000]
In [34]:
         min sample split = [5,10,100,500]
         count = 0;
         ind = []
         train auc = []
         cv auc = []
         hyp = []
         plt.figure(figsize=(30,30))
         for i in (depth_in_range):
             cv auc plot = []
             train_auc_plot = []
             for j in (min sample split):
                 dtc = DecisionTreeClassifier(max depth = i,min samples split
         = j
                 dtc.fit(X_train_vect, y_train)
                  ind.append(count)
                 hyp.append([i,j])
                 y_train_pred = dtc.predict_proba(X_train vect)[:,1]
                 v cv pred = dtc.predict proba(X cv vect)[:,1]
                  train auc.append(roc auc score(y train,y train pred))
                  train_auc_plot.append(roc_auc_score(y_train,y_train_pred))
                  cv auc.append(roc auc score(y cv, y cv pred))
                  cv auc plot.append(roc auc score(y cv, y cv pred))
                  count += 1
             plt.plot(np.log(min sample split), train auc plot, label='Train A
         UC when depth in range = ' + str(i))
             plt.plot(np.log(min sample split), cv auc plot, label='CV AUC whe
         n depth in range = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("min_sample_split: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```

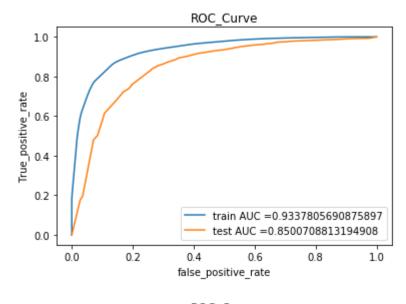


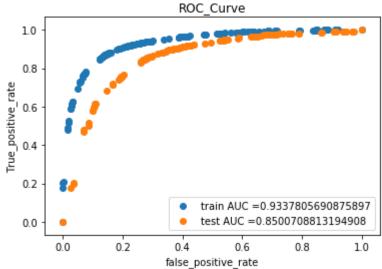
The optimal max_depth is (according to auc curve (max auc)): 50

The optimal min_sample_split is (according to auc curve (max auc)): 500

```
In [70]:
         dtc = DecisionTreeClassifier(max depth = optimal alpha auc[0],min sam
         ples split = optimal alpha auc[1])
         dtc.fit(X train vect, y train)
         pred = dtc.predict(X test vect)
         acc = accuracy score(y test, pred, normalize=True) * float(100)
         print('\setminus n^{***}Test accuracy formax depth = %f and min samples split =
         %f is %f%%' % (optimal_alpha_auc[0],optimal_alpha auc[1],acc))
         train_fpr, train_tpr, thresholds = roc_curve(y_train, dtc.predict_pro
         ba(X train vect)[:,1])
         test_fpr, test_tpr, thresholds = roc_curve(y_test, dtc.predict proba(
         X test vect)[:,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, tes
         t tpr)))
         plt.legend()
         plt.xlabel("false positive rate")
         plt.ylabel("True positive rate")
         plt.title("ROC Curve")
         plt.show()
         plt.scatter(train fpr, train tpr, label="train AUC ="+str(auc(train f
         pr, train tpr)))
         plt.scatter(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("ROC Curve")
         plt.show()
         print("="*100)
```

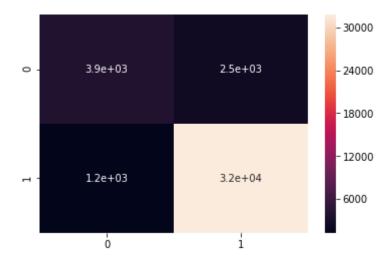
****Test accuracy formax_depth = 50.000000 and min_samples_split = 50 0.000000 is 86.712007%





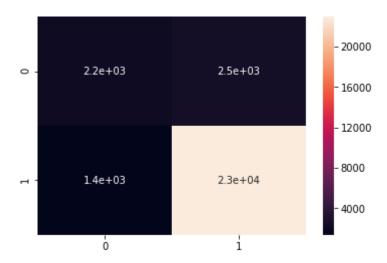
```
In [73]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_train, dtc.predict(X_train_vect))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
```

```
Train confusion matrix [[ 3881 2494] [ 1234 31791]]
```



```
In [71]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_test, dtc.predict(X_test_vect))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
```

```
Train confusion matrix [[ 2197 2474] [ 1375 22920]]
```



[5.1.1] Top 20 important features from SET 1

```
In [37]: # Please write all the code with proper documentation
imp = dtc.feature_importances_

vocab = list(count_vect.get_feature_names())
imp_features = list(imp)
dict_new = {'Words':vocab, 'imp_features':imp_features}
df_new = pd.DataFrame(dict_new)
df_pos_sorted =df_new.sort_values('imp_features', axis=0, ascending=F
alse, inplace=False, kind='quicksort', na_position='last')
print("The top 20 important features are the following:")
print(df_pos_sorted[0:20])
print("\n")
```

```
The top 20 important features are the following:
                     imp features
             Words
4820
                         0.116744
                not
3217
             great
                         0.082265
2148
      disappointed
                         0.042088
746
              best
                         0.038781
589
             awful
                         0.036649
         delicious
                         0.033273
1985
3496
          horrible
                         0.032353
4232
              love
                         0.030641
3156
                         0.029527
              good
7280
          terrible
                         0.021394
5211
           perfect
                         0.019686
                         0.018714
605
                bad
4237
             loves
                         0.017881
2567
         excellent
                         0.016515
4781
                         0.016468
              nice
8054
             worst
                         0.014755
7218
                         0.012683
             tasty
7886
             waste
                         0.012661
2711
                         0.011340
          favorite
```

0.010273

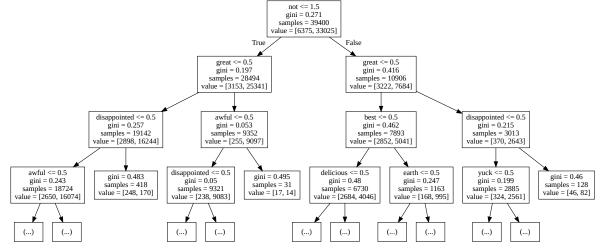
[5.1.2] Graphviz visualization of Decision Tree on BOW, SET 1

money

4616

```
In [38]: import graphviz
    from sklearn import tree
    import os
    dot_data = tree.export_graphviz(dtc, out_file= None, max_depth=3, fea
    ture_names = vocab)
    graph = graphviz.Source(dot_data)
    graph
```

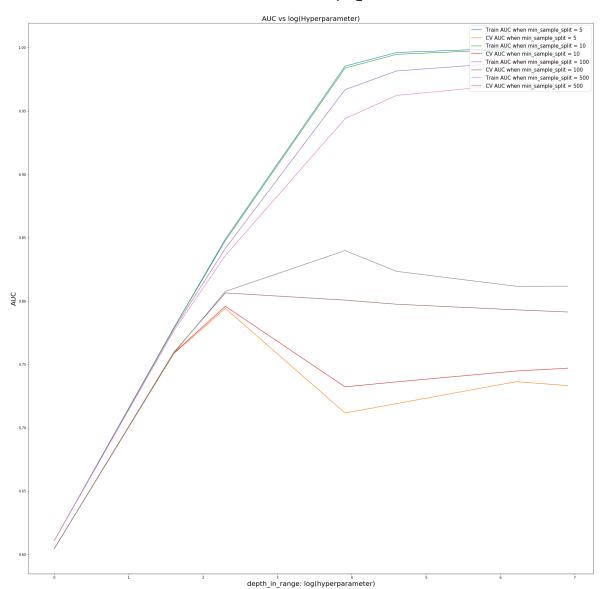
Out[38]:



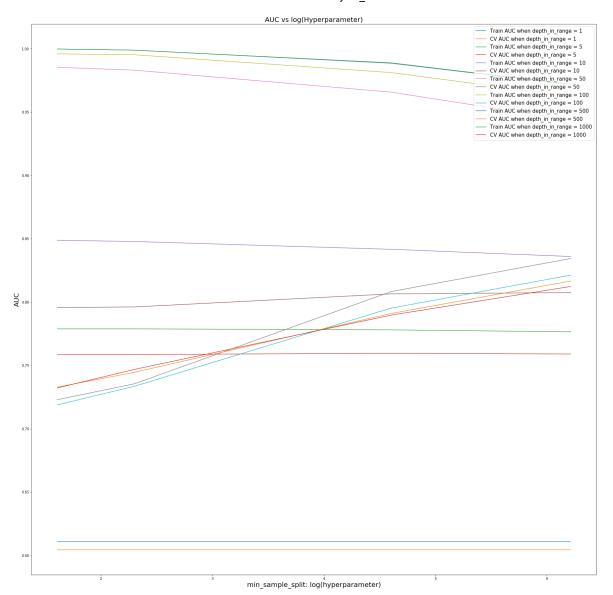
[5.2] Applying Decision Trees on TFIDF, SET 2

```
from sklearn.tree import DecisionTreeClassifier
In [42]:
         from sklearn.model selection import GridSearchCV
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         parameters = [{\text{max depth'}}: [1,5,10,50,100,500,1000]},{\text{min samples s}}
         plit':[5,10,100,500]}]
         model = GridSearchCV(DecisionTreeClassifier(),parameters, scoring =
          'roc auc')
         model.fit(X train vect tfidf, y train)
         print(model.best estimator )
         print(model.score(X_test_vect_tfidf, y_test))
         DecisionTreeClassifier(class weight=None, criterion='gini', max depth
         =None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=500,
                      min weight fraction leaf=0.0, presort=False, random state
         =None,
                      splitter='best')
         0.8121771397203316
```

```
from sklearn.metrics import accuracy score
In [43]:
         from sklearn.metrics import roc auc score
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         ind = []
         train auc = []
         cv auc = []
         plt.figure(figsize=(30,30))
         for i in (min sample split):
             cv auc plot = []
             train auc plot = []
             for j in (depth in range):
                 dtc = DecisionTreeClassifier(max depth = j,min samples split
         = i)
                 dtc.fit(X train vect tfidf, y train)
                 y train pred = dtc.predict proba(X train vect tfidf)[:,1]
                 y_cv_pred = dtc.predict_proba(X_cv_vect_tfidf)[:,1]
                  train auc.append(roc auc score(y train,y train pred))
                  train auc plot.append(roc auc score(y train,y train pred))
                  cv_auc.append(roc_auc_score(y_cv, y_cv_pred))
                  cv auc plot.append(roc auc score(y cv, y cv pred))
             plt.plot(np.log(depth in range), train auc plot, label='Train AUC
         when min sample split = ' + str(i))
             plt.plot(np.log(depth in range), cv auc plot, label='CV AUC when
          min sample split = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("depth in range: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```



```
depth in range = [1,5,10,50,100,500,1000]
In [44]:
         min sample split = [5,10,100,500]
         count = 0;
         ind = []
         train auc = []
         cv auc = []
         hyp = []
         plt.figure(figsize=(30,30))
         for i in (depth_in_range):
             cv auc plot = []
             train_auc_plot = []
             for j in (min sample split):
                 dtc = DecisionTreeClassifier(max_depth = i,min_samples_split
         = j
                 dtc.fit(X train vect tfidf, y train)
                  ind.append(count)
                 hyp.append([i,j])
                 y train pred = dtc.predict proba(X train vect tfidf)[:,1]
                 v cv pred = dtc.predict proba(X cv vect tfidf)[:,1]
                 train auc.append(roc auc score(y train,y train pred))
                  train_auc_plot.append(roc_auc_score(y_train,y_train_pred))
                  cv auc.append(roc auc score(y cv, y cv pred))
                  cv auc plot.append(roc auc score(y cv, y cv pred))
                  count += 1
             plt.plot(np.log(min sample split), train auc plot, label='Train A
         UC when depth in range = ' + str(i))
             plt.plot(np.log(min sample split), cv auc plot, label='CV AUC whe
         n depth in range = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("min_sample_split: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```

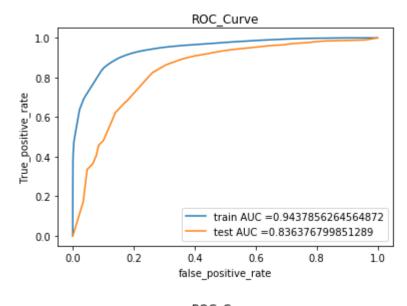


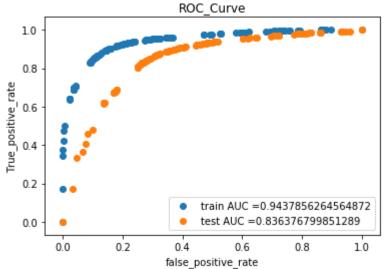
The optimal max_depth is (according to auc curve (max auc)): 50

The optimal min_sample_split is (according to auc curve (max auc)): 500

```
dtc = DecisionTreeClassifier(max depth = optimal alpha auc[0],min sam
ples split = optimal alpha auc[1])
dtc.fit(X train vect tfidf, y train)
pred = dtc.predict(X test vect tfidf)
acc = accuracy score(y test, pred, normalize=True) * float(100)
print('\setminus n^{***}Test accuracy formax depth = %f and min samples split =
%f is %f%%' % (optimal_alpha_auc[0],optimal_alpha auc[1],acc))
train_fpr, train_tpr, thresholds = roc_curve(y_train, dtc.predict_pro
ba(X train vect tfidf)[:,1])
test_fpr, test_tpr, thresholds = roc curve(y test, dtc.predict proba(
X_test_vect_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, tes
t tpr)))
plt.legend()
plt.xlabel("false positive rate")
plt.ylabel("True positive rate")
plt.title("ROC Curve")
plt.show()
plt.scatter(train fpr, train tpr, label="train AUC ="+str(auc(train f
pr, train tpr)))
plt.scatter(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("ROC Curve")
plt.show()
print("="*100)
```

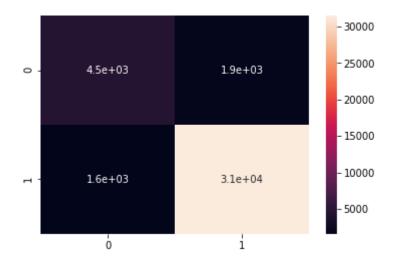
****Test accuracy formax_depth = 50.000000 and min_samples_split = 50 0.000000 is 86.321895%





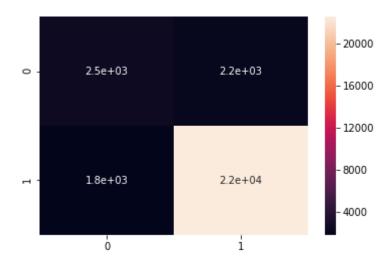
```
In [68]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_train, dtc.predict(X_train_vect_tfidf))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
Train confusion matrix
```

Train confusion matrix [[4508 1867] [1610 31415]]



```
In [69]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_test, dtc.predict(X_test_vect_tfidf))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
```

Train confusion matrix [[2514 2157] [1805 22490]]



[5.2.1] Top 20 important features from SET 2

```
In [47]: # Please write all the code with proper documentation
imp = dtc.feature_importances_

vocab = list(tf_idf.get_feature_names())
imp_features = list(imp)
dict_new = {'Words':vocab, 'imp_features':imp_features}
df_new = pd.DataFrame(dict_new)
df_pos_sorted =df_new.sort_values('imp_features', axis=0, ascending=F
alse, inplace=False, kind='quicksort', na_position='last')
print("The top 20 important features are the following:")
print(df_pos_sorted[0:20])
print("\n")
```

```
The top 20 important features are the following:
                     imp features
             Words
4820
                         0.131908
                not
3217
                         0.080218
             great
2148
      disappointed
                         0.046510
746
              best
                         0.036840
8054
                         0.035300
             worst
4232
                         0.030247
              love
3156
              good
                         0.029220
1985
         delicious
                         0.026794
                         0.021072
605
                bad
3496
          horrible
                         0.019477
589
             awful
                         0.019030
4616
             money
                         0.016564
4781
              nice
                         0.015166
5211
           perfect
                         0.013846
4237
             loves
                         0.013613
2711
          favorite
                         0.013069
2567
         excellent
                         0.012932
7886
             waste
                         0.011778
7218
             tasty
                         0.011646
```

0.011513

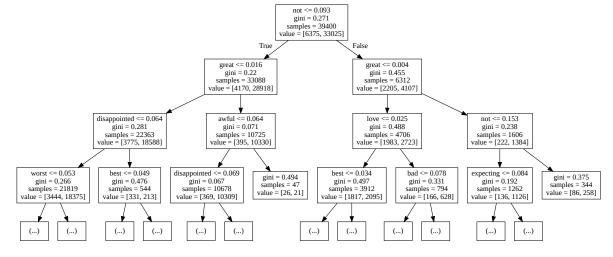
[5.2.2] Graphviz visualization of Decision Tree on TFIDF, SET 2

terrible

7280

```
In [48]: # Please write all the code with proper documentation
    import graphviz
    from sklearn import tree
    import os
    dot_data = tree.export_graphviz(dtc, out_file= None, max_depth=3, fea
    ture_names = vocab)
    graph = graphviz.Source(dot_data)
    graph
```

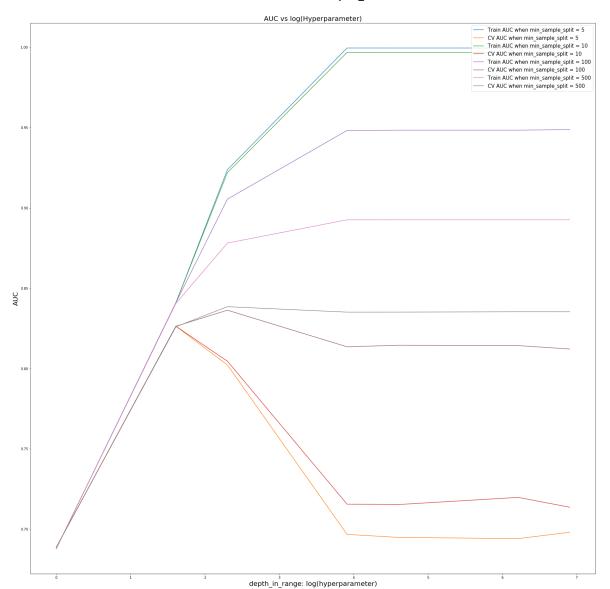
Out[48]:



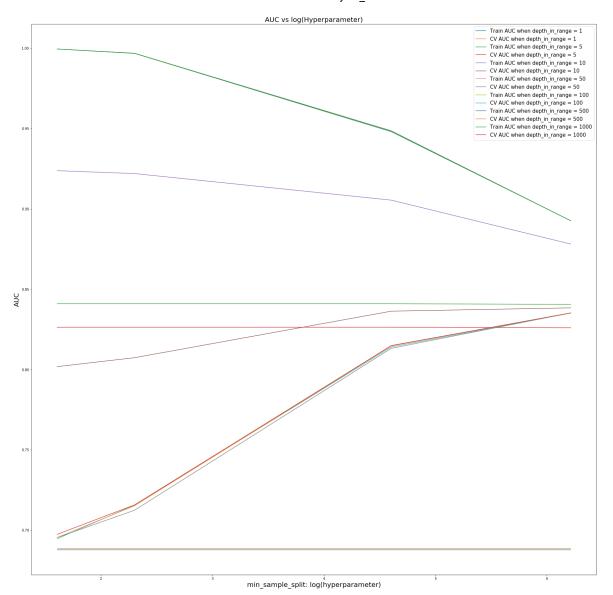
[5.3] Applying Decision Trees on AVG W2V, SET 3

```
In [49]:
         # Please write all the code with proper documentation
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.model selection import GridSearchCV
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         parameters = [{\text{max depth'}}: [1,5,10,50,100,500,1000]}, {\text{min samples s}}
         plit':[5,10,100,500]}]
         model = GridSearchCV(DecisionTreeClassifier(),parameters, scoring =
         'roc auc')
         model.fit(X_train_vectors, y_train)
         print(model.best estimator )
         print(model.score(X test vectors, y test))
         DecisionTreeClassifier(class_weight=None, criterion='gini', max depth
         =None,
                      max features=None, max leaf nodes=None,
                      min impurity decrease=0.0, min impurity split=None,
                      min samples leaf=1, min samples split=500,
                      min weight fraction leaf=0.0, presort=False, random state
         =None,
                      splitter='best')
         0.839527635871944
```

```
from sklearn.metrics import accuracy score
In [50]:
         from sklearn.metrics import roc auc score
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         ind = []
         train_auc = []
         cv_auc = []
         plt.figure(figsize=(30,30))
         for i in (min sample split):
             cv auc plot = []
             train auc plot = []
             for j in (depth in range):
                 dtc = DecisionTreeClassifier(max depth = j,min samples split
         = i)
                 dtc.fit(X train vectors, y train)
                 y train pred = dtc.predict proba(X train vectors)[:,1]
                 y_cv_pred = dtc.predict_proba(X_cv_vectors)[:,1]
                  train auc.append(roc auc score(y train,y train pred))
                  train auc plot.append(roc auc score(y train,y train pred))
                  cv_auc.append(roc_auc_score(y_cv, y_cv_pred))
                 cv auc plot.append(roc auc score(y cv, y cv pred))
             plt.plot(np.log(depth in range), train auc plot, label='Train AUC
         when min sample split = ' + str(i))
             plt.plot(np.log(depth in range), cv auc plot, label='CV AUC when
          min sample split = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("depth in range: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```



```
depth in range = [1,5,10,50,100,500,1000]
In [51]:
         min sample split = [5,10,100,500]
         count = 0;
         ind = []
         train auc = []
         cv auc = []
         hyp = []
         plt.figure(figsize=(30,30))
         for i in (depth_in_range):
             cv auc plot = []
             train_auc_plot = []
             for j in (min sample split):
                 dtc = DecisionTreeClassifier(max_depth = i,min_samples_split
         = j
                 dtc.fit(X_train_vectors, y_train)
                  ind.append(count)
                 hyp.append([i,j])
                 y train pred = dtc.predict proba(X train vectors)[:,1]
                 v cv pred = dtc.predict proba(X cv vectors)[:,1]
                  train auc.append(roc auc score(y train,y train pred))
                  train_auc_plot.append(roc_auc_score(y_train,y_train_pred))
                  cv auc.append(roc auc score(y cv, y cv pred))
                  cv auc plot.append(roc auc score(y cv, y cv pred))
                  count += 1
             plt.plot(np.log(min sample split), train auc plot, label='Train A
         UC when depth in range = ' + str(i))
             plt.plot(np.log(min sample split), cv auc plot, label='CV AUC whe
         n depth in range = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("min_sample_split: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```



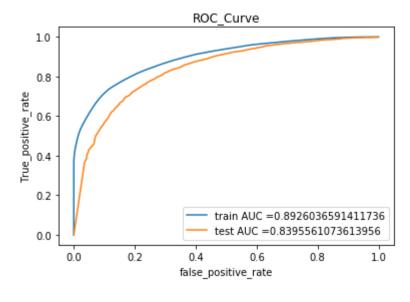
```
In [52]: optimal_alpha_auc = hyp[cv_auc.index(max(cv_auc))]
    print('\nThe optimal max_depth is (according to auc curve (max auc)):
    ' , optimal_alpha_auc[0])
    print('\nThe optimal min_sample_split is (according to auc curve (max auc)): ' , optimal_alpha_auc[1])
```

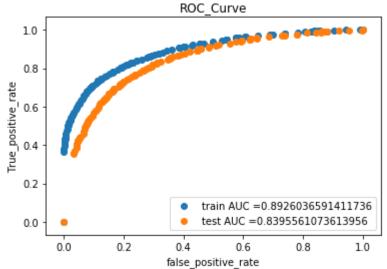
The optimal max_depth is (according to auc curve (max auc)): 10

The optimal min_sample_split is (according to auc curve (max auc)): 500

```
dtc = DecisionTreeClassifier(max depth = optimal alpha auc[0], min sam
ples split = optimal alpha auc[1])
dtc.fit(X train vectors, y train)
pred = dtc.predict(X test vectors)
acc = accuracy score(y test, pred, normalize=True) * float(100)
print('\setminus n^{***}Test accuracy formax depth = %f and min samples split =
%f is %f%%' % (optimal_alpha_auc[0],optimal_alpha auc[1],acc))
train_fpr, train_tpr, thresholds = roc_curve(y_train, dtc.predict_pro
ba(X train vectors)[:,1])
test_fpr, test_tpr, thresholds = roc_curve(y_test, dtc.predict proba(
X test vectors)[:,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, tes
t tpr)))
plt.legend()
plt.xlabel("false positive rate")
plt.ylabel("True positive rate")
plt.title("ROC Curve")
plt.show()
plt.scatter(train fpr, train tpr, label="train AUC ="+str(auc(train f
pr, train tpr)))
plt.scatter(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("ROC Curve")
plt.show()
print("="*100)
```

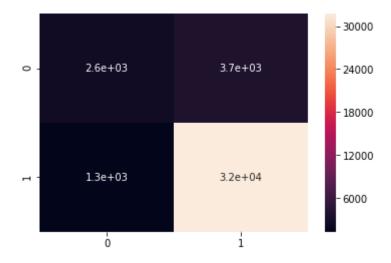
****Test accuracy formax_depth = 50.000000 and min_samples_split = 50 0.000000 is 85.938687%





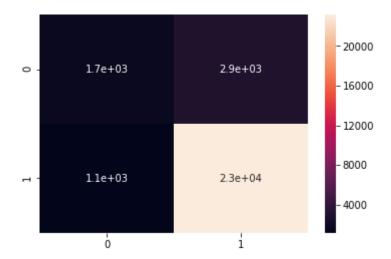
```
In [65]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_train, dtc.predict(X_train_vectors))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
Train confusion matrix
```

Train confusion matrix [[2650 3725] [1328 31697]]



```
In [66]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_test, dtc.predict(X_test_vectors))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
```

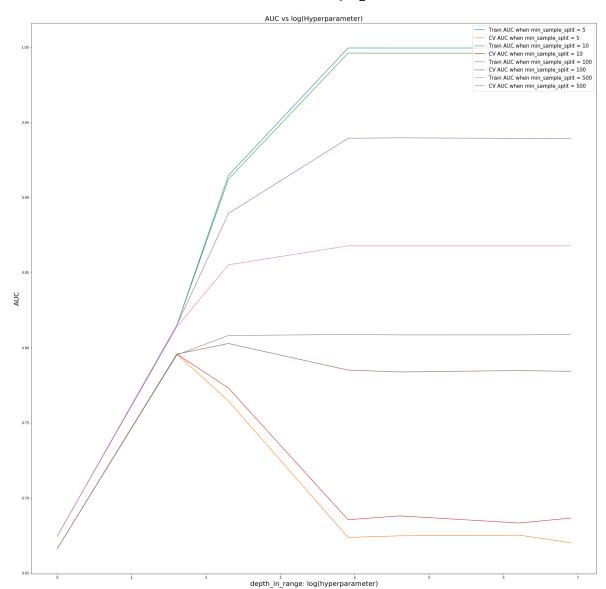
Train confusion matrix [[1745 2926] [1147 23148]]



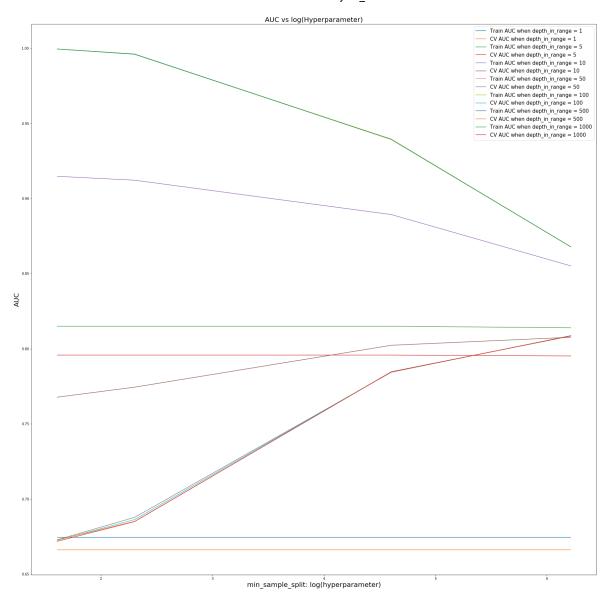
[5.4] Applying Decision Trees on TFIDF W2V, SET 4

```
# Please write all the code with proper documentation
In [56]:
         from sklearn.tree import DecisionTreeClassifier
         from sklearn.model selection import GridSearchCV
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         parameters = [{'max depth': [1,5,10,50,100,500,1000]},{'min samples s
         plit':[5,10,100,500]}]
         model = GridSearchCV(DecisionTreeClassifier(),parameters, scoring =
         model.fit(tfidf_X_train_vectors, y_train)
         print(model.best estimator )
         print(model.score(tfidf_X_test_vectors, y_test))
         DecisionTreeClassifier(class weight=None, criterion='gini', max depth
         =None,
                     max_features=None, max_leaf_nodes=None,
                     min impurity decrease=0.0, min impurity split=None,
                     min_samples_leaf=1, min_samples_split=500,
                     min weight fraction leaf=0.0, presort=False, random state
         =None,
                     splitter='best')
         0.8096382468594454
```

```
from sklearn.metrics import accuracy score
In [57]:
         from sklearn.metrics import roc auc score
         depth in range = [1,5,10,50,100,500,1000]
         min sample split = [5,10,100,500]
         ind = []
         train_auc = []
         cv_auc = []
         plt.figure(figsize=(30,30))
         for i in (min sample split):
             cv auc plot = []
             train auc plot = []
             for j in (depth in range):
                 dtc = DecisionTreeClassifier(max depth = j,min samples split
         = i)
                 dtc.fit(tfidf X train vectors, y train)
                 y train pred = dtc.predict proba(tfidf X train vectors)[:,1]
                 y_cv_pred = dtc.predict_proba(tfidf_X_cv_vectors)[:,1]
                  train auc.append(roc auc score(y train,y train pred))
                  train auc plot.append(roc auc score(y train,y train pred))
                  cv_auc.append(roc_auc_score(y_cv, y_cv_pred))
                 cv auc plot.append(roc auc score(y cv, y cv pred))
             plt.plot(np.log(depth in range), train auc plot, label='Train AUC
         when min sample split = ' + str(i))
             plt.plot(np.log(depth in range), cv auc plot, label='CV AUC when
          min sample split = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("depth in range: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```



```
depth in range = [1,5,10,50,100,500,1000]
In [58]:
         min sample split = [5,10,100,500]
         count = 0;
         ind = []
         train auc = []
         cv auc = []
         hyp = []
         plt.figure(figsize=(30,30))
         for i in (depth_in_range):
             cv auc plot = []
             train_auc_plot = []
             for j in (min sample split):
                 dtc = DecisionTreeClassifier(max_depth = i,min_samples_split
         = j
                 dtc.fit(tfidf X train vectors, y_train)
                  ind.append(count)
                 hyp.append([i,j])
                 y train pred = dtc.predict_proba(tfidf_X_train_vectors)[:,1]
                 v cv pred = dtc.predict proba(tfidf X cv vectors)[:,1]
                 train auc.append(roc auc score(y train,y train pred))
                  train_auc_plot.append(roc_auc_score(y_train,y_train_pred))
                  cv auc.append(roc auc score(y cv, y cv pred))
                  cv auc plot.append(roc auc score(y cv, y cv pred))
                  count += 1
             plt.plot(np.log(min sample split), train auc plot, label='Train A
         UC when depth in range = ' + str(i))
             plt.plot(np.log(min sample split), cv auc plot, label='CV AUC whe
         n depth in range = ' + str(i))
         plt.legend(loc=1, prop={'size': 15})
         plt.xlabel("min_sample_split: log(hyperparameter)", fontsize = 20)
         plt.ylabel("AUC", fontsize = 20)
         plt.title("AUC vs log(Hyperparameter)",fontsize = 20)
         plt.show()
```



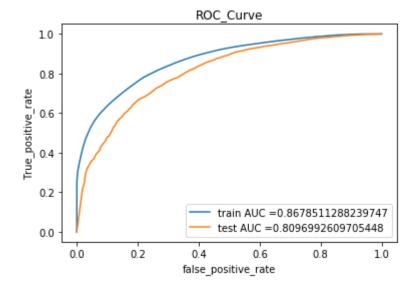
```
In [59]: optimal_alpha_auc = hyp[cv_auc.index(max(cv_auc))]
    print('\nThe optimal max_depth is (according to auc curve (max auc)):
    ' , optimal_alpha_auc[0])
    print('\nThe optimal min_sample_split is (according to auc curve (max auc)): ' , optimal_alpha_auc[1])
```

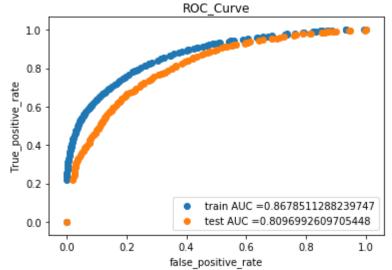
The optimal max_depth is (according to auc curve (max auc)): 50

The optimal min_sample_split is (according to auc curve (max auc)): 500

```
In [60]:
         dtc = DecisionTreeClassifier(max depth = optimal alpha auc[0], min sam
         ples split = optimal alpha auc[1])
         dtc.fit(tfidf X train vectors, y train)
         pred = dtc.predict(tfidf X test vectors)
         acc = accuracy score(y test, pred, normalize=True) * float(100)
         print('\setminus n^{***}Test accuracy formax depth = %f and min samples split =
         %f is %f%%' % (optimal_alpha_auc[0],optimal_alpha auc[1],acc))
         train_fpr, train_tpr, thresholds = roc_curve(y_train, dtc.predict_pro
         ba(tfidf X train vectors)[:,1])
         test fpr, test tpr, thresholds = roc curve(y test, dtc.predict proba(
         tfidf_X_test_vectors)[:,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, tes
         t tpr)))
         plt.legend()
         plt.xlabel("false positive rate")
         plt.ylabel("True positive rate")
         plt.title("ROC Curve")
         plt.show()
         plt.scatter(train fpr, train tpr, label="train AUC ="+str(auc(train f
         pr, train tpr)))
         plt.scatter(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("ROC Curve")
         plt.show()
         print("="*100)
```

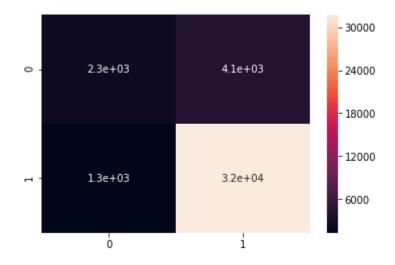
****Test accuracy formax_depth = 50.000000 and min_samples_split = 50 0.000000 is 85.044535%





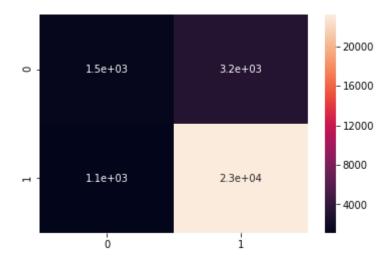
```
In [61]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_train, dtc.predict(tfidf_X_train_vectors))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
```

Train confusion matrix [[2309 4066] [1290 31735]]



```
In [62]: from sklearn.metrics import confusion_matrix
    print("Train confusion matrix")
    cm_tr = confusion_matrix(y_test, dtc.predict(tfidf_X_test_vectors))
    sns.heatmap(cm_tr, annot=True)
    print(cm_tr)
```

Train confusion matrix [[1465 3206] [1126 23169]]



[6] Conclusions

```
In [74]: from prettytable import PrettyTable
x = PrettyTable()

x.field_names = ["Vectorizer", "max_depth", "min_samples_split", "AU
C"]
x.add_row(["BOW", 50, 500, 0.85])
x.add_row(["TFIDF", 50, 500, 0.84])
x.add_row(["W2V", 50, 500, 0.84])
x.add_row(["TFIDFW2V", 50, 500, 0.80])
print(x)
```

Vectorizer	 max_depth 	min_samples_split	++ AUC ++
BOW TFIDF W2V TFIDFW2V	50	500	0.85
	50	500	0.84
	50	500	0.84
	50	500	0.8

```
In [ ]:
```