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

Objective:

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

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

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

[1]. Reading Data

[1.1] Loading the data

The dataset is available in two forms

- 1. .csv file
- 2. SQLite Database

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

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

```
In [256]: %matplotlib inline
import warnings
warnings.filterwarnings("ignore")

import sqlite3
import pandas as pd
import numpy as np
import nltk
import string
```

```
import matplotlib.pyplot as plt
import seaborn as sns
from sklearn.feature extraction.text import TfidfTransformer
from sklearn.feature extraction.text import TfidfVectorizer
from sklearn.feature extraction.text import CountVectorizer
from sklearn.metrics import confusion matrix
from sklearn import metrics
from sklearn.metrics import roc curve, auc
from nltk.stem.porter import PorterStemmer
import re
# Tutorial about Python regular expressions: https://pymotw.com/2/re/
import string
from nltk.corpus import stopwords
from nltk.stem import PorterStemmer
from nltk.stem.wordnet import WordNetLemmatizer
from gensim.models import Word2Vec
from gensim.models import KeyedVectors
import pickle
from tadm import tadm
import os
from sklearn.cross validation import train test split
from sklearn.neighbors import KNeighborsClassifier
from sklearn.metrics import accuracy score
from sklearn.cross validation import cross val score
from collections import Counter
from sklearn.metrics import accuracy score
from sklearn import cross validation
con = sqlite3.connect('/Users/puravshah/Downloads/amazon-fine-food-revi
```

```
In [257]: # using SQLite Table to read data.
con = sqlite3.connect('/Users/puravshah/Downloads/amazon-fine-food-revi
ews/database.sqlite')

# filtering only positive and negative reviews i.e.
# not taking into consideration those reviews with Score=3
```

```
# SELECT * FROM Reviews WHERE Score != 3 LIMIT 500000, will give top 50
0000 data points
# you can change the number to any other number based on your computing
# filtered data = pd.read sql query(""" SELECT * FROM Reviews WHERE Sco
re != 3 LIMIT 500000""", con)
# for tsne assignment you can take 5k data points
filtered data = pd.read sql query(""" SELECT * FROM Reviews WHERE Score
!= 3 LIMIT 100000""", con)
# Give reviews with Score>3 a positive rating(1), and reviews with a sc
ore<3 a negative rating(0).
def partition(x):
    if x < 3:
        return 0
    return 1
#changing reviews with score less than 3 to be positive and vice-versa
actualScore = filtered data['Score']
positiveNegative = actualScore.map(partition)
filtered data['Score'] = positiveNegative
print("Number of data points in our data", filtered data.shape)
filtered data.head(3)
```

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

Out[257]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfulnes
0	1	B001E4KFG0	A3SGXH7AUHU8GW	delmartian	1	1

	ld	ProductId	Userl	d ProfileName	Helpfuln	essNumerator	Help	fulnes
1	2	B00813GRG4	A1D87F6ZCVE5NK	dll pa	0		0	
2	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1		1	
4 ■					1			•
SE FR GR HA	LEC OM OUF		d_sql_query(""" roductId, Profile	eName, Time,	Score,	Text, COUNT(*)	
		(display.sha	ape)					
(8	066	88, 7)						
		Userl	d Productid Pr	ofileName	Time	Score	Text	COU

In [258]:

In [259]:

Out[259]:

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

In [260]: display[display['UserId']=='AZY10LLTJ71NX']

Out[260]:

UserId ProductId ProfileName Time Score Text
--

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

In [261]: display['COUNT(*)'].sum()

Out[261]: 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 [262]: display= pd.read_sql_query("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND UserId="AR5J8UI46CURR"
    ORDER BY ProductID
    """, con)
    display.head()
```

Out[262]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfuln
--	----	-----------	--------	-------------	----------------------	----------

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

As 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

```
In [266]: display= pd.read_sql_query("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND Id=44737 OR Id=64422
    ORDER BY ProductID
    """, con)
    display.head()
```

Out[266]:

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

In [267]: final=final[final.HelpfulnessNumerator<=final.HelpfulnessDenominator]</pre>

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

(87773, 10)

Out[268]: 1 73592
0 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 [269]: # 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 too bec ause its a good product but I wont take any chances till they know 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 w ere 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 sme ll. So if you are afraid of the fishy smell, don't get it. But I think 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 the se 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 them a lot.

```
In [270]: # remove urls from text python: https://stackoverflow.com/a/40823105/40 84039 sent_0 = re.sub(r"http\S+", "", sent_0) \\ sent_1000 = re.sub(r"http\S+", "", sent_1000)
```

```
sent_150 = re.sub(r"http\S+", "", sent_1500)
sent_4900 = re.sub(r"http\S+", "", sent_4900)
print(sent_0)
```

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 too bec ause its a good product but I wont take any chances till they know what is going on with the china imports.

```
In [271]: # https://stackoverflow.com/questions/16206380/python-beautifulsoup-how
          -to-remove-all-tags-from-an-element
          from bs4 import BeautifulSoup
          soup = BeautifulSoup(sent 0, 'lxml')
          text = soup.get text()
          print(text)
          print("="*50)
          soup = BeautifulSoup(sent 1000, 'lxml')
          text = soup.get text()
          print(text)
          print("="*50)
          soup = BeautifulSoup(sent 1500, 'lxml')
          text = soup.get text()
          print(text)
          print("="*50)
          soup = BeautifulSoup(sent 4900, 'lxml')
          text = soup.get text()
          print(text)
```

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 too bec ause its a good product but I wont take any chances till they know what is going on with the china imports.

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My dog LOVES these treats. They tend to have a very strong fish oil sme ll. So if you are afraid of the fishy smell, don't get it. But I think 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 the se 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 them a lot.

```
In [272]: # https://stackoverflow.com/a/47091490/4084039
import re

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

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

```
In [273]: 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

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 too bec ause its a good product but I wont take any chances till they know what is going on with the china imports.

```
In [275]: #remove spacial character: https://stackoverflow.com/a/5843547/4084039
    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 [276]: # https://gist.github.com/sebleier/554280
          # we are removing the words from the stop words list: 'no', 'nor', 'no
          # <br /><br /> ==> after the above steps, we are getting "br br"
          # we are including them into stop words list
          # instead of <br /> if we have <br/> these tags would have revmoved in
           the 1st step
          stopwords= set(['br', 'the', 'i', 'me', 'my', 'myself', 'we', 'our', 'o
          urs', 'ourselves', 'you', "you're", "you've",\
                      "you'll", "you'd", 'your', 'yours', 'yourself', 'yourselve
          s', 'he', 'him', 'his', 'himself', \
                      'she', "she's", 'her', 'hers', 'herself', 'it', "it's", 'it
          s', 'itself', 'they', 'them', 'their',\
                      'theirs', 'themselves', 'what', 'which', 'who', 'whom', 'th
          is', 'that', "that'll", 'these', 'those', \
                      'am', 'is', 'are', 'was', 'were', 'be', 'been', 'being', 'h
          ave', 'has', 'had', 'having', 'do', 'does', \
                      'did', 'doing', 'a', 'an', 'the', 'and', 'but', 'if', 'or',
           'because', 'as', 'until', 'while', 'of', \
```

```
'at', 'by', 'for', 'with', 'about', 'against', 'between',
          'into', 'through', 'during', 'before', 'after',\
                      'above', 'below', 'to', 'from', 'up', 'down', 'in', 'out',
          'on', 'off', 'over', 'under', 'again', 'further',\
                      'then', 'once', 'here', 'there', 'when', 'where', 'why', 'h
          ow', 'all', 'any', 'both', 'each', 'few', 'more',\
                      'most', 'other', 'some', 'such', 'only', 'own', 'same', 's
          o', 'than', 'too', 'very', \
                      's', 't', 'can', 'will', 'just', 'don', "don't", 'should',
          "should've", 'now', 'd', 'll', 'm', 'o', 're', \
                      've', 'y', 'ain', 'aren', "aren't", 'couldn', "couldn't",
          'didn', "didn't", 'doesn', "doesn't", 'hadn',\
                      "hadn't", 'hasn', "hasn't", 'haven', "haven't", 'isn', "is
          n't", 'ma', 'mightn', "mightn't", 'mustn',\
                      "mustn't", 'needn', "needn't", 'shan', "shan't", 'shouldn',
           "shouldn't", 'wasn', "wasn't", 'weren', "weren't", \
                      'won', "won't", 'wouldn', "wouldn't"])
In [277]: # 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.lower
          () not in stopwords)
              preprocessed reviews.append(sentance.strip())
                | 87773/87773 [00:25<00:00, 3422.82it/s]
In [278]: preprocessed reviews[1500]
Out[278]: 'way hot blood took bite jig lol'
```

[3.2] Preprocessing Review Summary

In [0]: ## Similartly you can do preprocessing for review summary also.

[4] Featurization

[4.1] BAG OF WORDS

```
In [311]: X 1, X test, y 1, y test = cross validation.train test split(preprocess
          ed reviews, final['Score'], test size=0.3, random state=0)
          X tr, X cv, y tr, y cv = cross validation.train test split(X 1, y 1, te
          st size=0.3)
          count vect = CountVectorizer() #in scikit-learn
          count vect.fit(X tr)
          count vect.fit(X cv)
          count vect.fit(X test)
          print("some feature names ", count vect.get feature names()[:10])
          print('='*50)
          final counts = count vect.transform(X tr)
          X cv bow=count vect.transform(X cv)
          X test bow=count vect.transform(X test)
          print("the type of count vectorizer ", type(final counts))
          print("the shape of out text BOW vectorizer ",final counts.get shape())
          print("the number of unique words ", final counts.get shape()[1])
          some feature names ['aa', 'aaa', 'aaaa', 'aaaand', 'aaah', 'aaahs', 'a
          afco', 'aahhhs', 'aahing', 'aamazon']
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
          the shape of out text BOW vectorizer (43008, 30512)
          the number of unique words 30512
```

[4.2] Bi-Grams and n-Grams.

```
In [280]: #bi-gram, tri-gram and n-gram
          #removing stop words like "not" should be avoided before building n-gra
          # count vect = CountVectorizer(ngram range=(1,2))
          # please do read the CountVectorizer documentation http://scikit-learn.
          org/stable/modules/generated/sklearn.feature extraction.text.CountVecto
          rizer.html
          # you can choose these numebrs min df=10, max features=5000, of your ch
          oice
          count vect = CountVectorizer(ngram range=(1,2), min df=10, max features)
          =5000)
          final bigram counts = count vect.fit transform(X tr)
          X cv ngram = count vect.fit_transform(X_cv)
          X test ngram=count vect.fit transform(X test)
          print("the type of count vectorizer ", type(final bigram counts))
          print("the shape of out text BOW vectorizer ",final bigram counts.get s
          hape())
          print("the number of unique words including both unigrams and bigrams "
           , final bigram counts.get shape()[1])
```

the type of count vectorizer <class 'scipy.sparse.csr.csr_matrix'> the shape of out text BOW vectorizer (43008, 5000) the number of unique words including both uniquems and bigrams 5000

[4.3] TF-IDF

```
In [281]: tf_idf_vect = TfidfVectorizer(ngram_range=(1,2), min_df=10)
    tf_idf_vect.fit(X_tr)
    tf_idf_vect.fit(X_cv)
    tf_idf_vect.fit(X_test)
    print("some sample features(unique words in the corpus)",tf_idf_vect.ge
    t_feature_names()[0:10])
    print('='*50)
```

```
final tf idf = tf idf vect.transform(X tr)
          X cv tfidf=tf idf vect.transform(X cv)
          X test tfidf=tf idf vect.transform(X test)
          print("the type of count vectorizer ", type(final tf idf))
          print("the shape of out text TFIDF vectorizer ",final tf idf.get shape
          ())
          print("the number of unique words including both unigrams and bigrams "
          , final tf idf.get shape()[1])
          some sample features(unique words in the corpus) ['ability', 'able', 'a
          ble buy', 'able drink', 'able eat', 'able enjoy', 'able find', 'able ge
          t', 'able give', 'able make']
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
          the shape of out text TFIDF vectorizer (43008, 15763)
          the number of unique words including both unigrams and bigrams 15763
          [4.4] Word2Vec
In [282]: # Train your own Word2Vec model using your own text corpus
          i=0
          list of sentance=[]
          for sentance in preprocessed reviews:
              list of sentance.append(sentance.split())
In [283]: # 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 val
          # To use this code-snippet, download "GoogleNews-vectors-negative300.bi
```

from https://drive.google.com/file/d/0B7XkCwpI5KDYNlNUTTlSS21pQmM/edi

```
# it's 1.9GB in size.
# http://kavita-ganesan.com/gensim-word2vec-tutorial-starter-code/#.W17
SRFA77PY
# 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,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=True)
        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 trai
n w2v = True, to train your own w2v ")
[('fantastic', 0.8536819815635681), ('awesome', 0.8350322842597961),
('excellent', 0.8242692351341248), ('good', 0.8191254734992981), ('terr
ific', 0.7985398173332214), ('wonderful', 0.7982454299926758), ('perfec
t', 0.770243763923645), ('amazing', 0.7415543794631958), ('fabulous',
0.7378154993057251), ('nice', 0.7201124429702759)]
[('greatest', 0.8203911781311035), ('tastiest', 0.7233695983886719),
('best', 0.7149415016174316), ('nastiest', 0.6551864147186279), ('ive',
0.6354540586471558), ('closest', 0.6261584758758545), ('surpass', 0.602
5659441947937), ('weakest', 0.6011222004890442), ('horrible', 0.5994516
611099243), ('awful', 0.5992503762245178)]
```

```
In [284]: 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 17386
    sample words ['dogs', 'loves', 'chicken', 'product', 'china', 'wont',
    'buying', 'anymore', 'hard', 'find', 'products', 'made', 'usa', 'one',
    'isnt', 'bad', 'good', 'take', 'chances', 'till', 'know', 'going', 'imp
    orts', 'love', 'saw', 'pet', 'store', 'tag', 'attached', 'regarding',
    'satisfied', 'safe', 'infestation', 'literally', 'everywhere', 'flyin
    g', 'around', 'kitchen', 'bought', 'hoping', 'least', 'get', 'rid', 'we
    eks', 'fly', 'stuck', 'squishing', 'buggers', 'success', 'rate']
```

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

[4.4.1.1] Avg W2v

```
In [285]: # average Word2Vec
          # compute average word2vec for each review.
          sent vectors = []; # the avg-w2v for each sentence/review is stored in
           this list
          for sent in tqdm(list of sentance): # for each review/sentence
              sent vec = np.zeros(50) # as word vectors are of zero length 50, yo
          u 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/re
          view
              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.append(sent vec)
```

[4.4.1.2] TFIDF weighted W2v

```
In [286]: \# S = ["abc def pqr", "def def def abc", "pqr pqr def"]
          model = TfidfVectorizer()
          tf idf matrix = model.fit transform(preprocessed reviews)
          # we are converting a dictionary with word as a key, and the idf as a v
          alue
          dictionary = dict(zip(model.get feature names(), list(model.idf )))
In [287]: # 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 ce
          ll\ val = tfidf
          tfidf sent vectors = []; # the tfidf-w2v for each sentence/review is st
          ored in this list
          row=0;
          for sent in tqdm(list of sentance): # 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/r
          eview
              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.append(sent_vec)
    row += 1

100%| 87773/87773 [24:15<00:00, 56.02it/s]</pre>
```

[5] Assignment 4: Apply Naive Bayes

1. Apply Multinomial NaiveBayes 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)

2. The hyper paramter tuning(find best Alpha)

- Find the best hyper parameter which will give the maximum AUC value
- Consider a wide range of alpha values for hyperparameter tuning, start as low as 0.00001
- 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

 Find the top 10 features of positive class and top 10 features of negative class for both feature sets Set 1 and Set 2 using values of `feature_log_prob_` parameter of <u>MultinomialNB</u> and print their corresponding feature names

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. Here on X-axis you will have alpha values, since they have a wide range, just to represent those alpha values on the graph, apply log function on those alpha values.

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 confusion matrix with predicted and original labels of test data points. Please visualize your confusion matrices using seaborn heatmaps.



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.

Applying Multinomial Naive Bayes

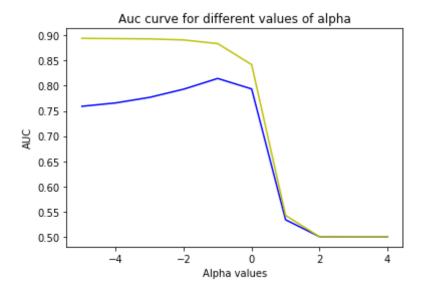
[5.1] Applying Naive Bayes on BOW, SET 1

```
In [323]: #Some parts of code taken from the reference ipynb provided
          from sklearn.naive bayes import MultinomialNB
          from sklearn.metrics import roc auc score
          auc=[]
          auc2=[1
          #defining alpha values
          print(np.log10(alpha))
          #for different alpha values finding the auc values to be plotted
          for i in alpha:
             model=MultinomialNB(alpha=i)
             model.fit(final counts,y tr)
             pred=model.predict(X cv bow)
             probab=model.predict proba(X cv bow)
             fpr,tpr,thresholds=metrics.roc curve(y cv,pred)
             auc.append(roc auc score(y cv,pred))
             acc=accuracy score(y cv,pred,normalize=True)*float(100)
             pred2=model.predict(final counts)
             probab2=model.predict proba(final counts)
             fpr train,tpr train,thresholds=roc curve(y tr,pred2)
             auc2.append(roc auc score(y tr,pred2))
              print('Accuracy score at alpha value %d is %d'%(i,acc))
          plt.plot(np.log10(alpha),auc,'b')
          plt.plot(np.log10(alpha),auc2,'y')
          plt.title('Auc curve for different values of alpha')
          plt.xlabel('Alpha values')
          plt.vlabel('AUC')
          #it can be seen from the plot that the auc values for the train and cv
          dataset overlap and there is no clear distinction between the two
          [-5, -4, -3, -2, -1, 0, 1, 2, 3, 4,]
```

Accuracy score at alpha value 0 is 88

Accuracy score at alpha value 0 is 88
Accuracy score at alpha value 0 is 89
Accuracy score at alpha value 0 is 89
Accuracy score at alpha value 0 is 90
Accuracy score at alpha value 1 is 90
Accuracy score at alpha value 10 is 84
Accuracy score at alpha value 1000 is 83
Accuracy score at alpha value 1000 is 83
Accuracy score at alpha value 10000 is 83

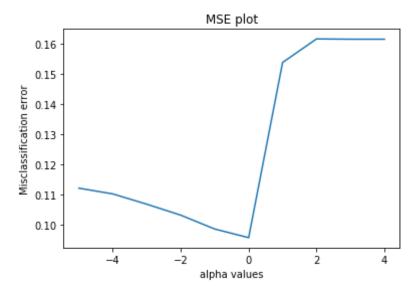
Out[323]: Text(0,0.5,'AUC')



```
In [324]: #This part of the code is to caluclate the optimal alpha by calculating
    the mis-classification error
    cv_scores=[]
    alpha=[0.00001,0.0001,0.001,0.1,1,10,100,1000,10000]
    for i in alpha:
        model=MultinomialNB(alpha=i)
        scores=cross_val_score(model,final_counts,y_tr,cv=10,scoring='accuracy')
        cv_scores.append(scores.mean())
    MSE=[1-x for x in cv_scores]
```

```
plt.plot(np.log10(alpha),MSE)
plt.title('MSE plot')
plt.xlabel('alpha values')
plt.ylabel('Misclassification error')
optimal_alpha=alpha[MSE.index(min(MSE))]
print('The optimal alpha vale for low error is=%d'%optimal_alpha)
```

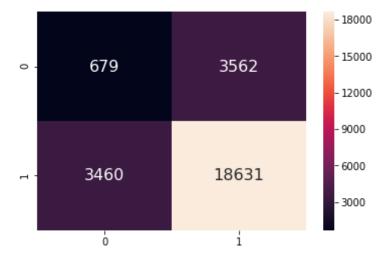
The optimal alpha vale for low error is=1

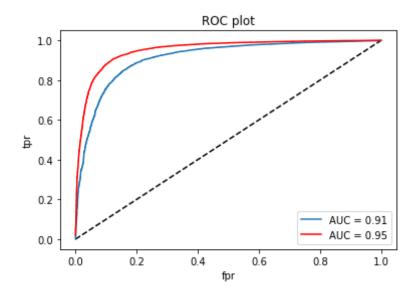


```
In [325]: #Using the optimal value of alpha that was calculated we run our model
   with that alpha value and check the accuracy
   model=MultinomialNB(alpha=optimal_alpha)
   model.fit(final_counts,y_tr)
   predict=model.predict(X_test_bow)
   probab=model.predict_proba(X_test_bow)
   probab=probab[:,1]
   predict2=model.predict(final_counts)
   probab2=model.predict_proba(final_counts)
   probab2=probab2[:,1]
   acc=accuracy_score(y_test,predict,normalize=True)*float(100)
   print('The accuracy of the model with optimal value of alpha is=%d'%opt imal_alpha)
```

The accuracy of the model with optimal value of alpha is=1

```
In [326]: from sklearn.metrics import confusion matrix as cm
          from sklearn.metrics import roc auc score
          #plotting the confusion matrix and the ROC curve
          df cm = pd.DataFrame(confusion matrix(Y test, predict), range(2), range(
          2))
              #heatman for visualization of matrix
          sns.heatmap(df cm, annot=True, annot kws={"size": 16}, fmt='g')
          plt.show()
          fpr,tpr,thresholds=roc curve(y test,probab)
          auc=roc auc score(y test,probab)
          fpr train,tpr train,thresholds=roc curve(y tr,probab2)
          auc2=roc auc score(y tr,probab2)
          plt.plot([0,1],[0,1],'k--')
          plt.plot(fpr,tpr,label = 'AUC = %0.2f' %auc)
          plt.plot(fpr train,tpr train,'r',label = 'AUC = %0.2f' %auc2)
          plt.title('ROC plot')
          plt.xlabel('fpr')
          plt.ylabel('tpr')
          plt.legend(loc='lower right')
          plt.show()
```





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

```
In [327]: #calculating the feature log probabilities
          feature log prob=model.feature log prob
          #print(feature_log_prob.shape)
          #concerting it into a data frame
          df=pd.DataFrame(feature log prob)
          #print(df.head())
          #Getting the feature names from bow vectorizer
          bow=count vect.get feature names()
          #Making the feature names as the index(matching the probabilities to th
          e respective values)
          df=pd.DataFrame(feature log prob,columns=bow)
          #print(df.head())
          df=df.T
          #print(df.head())
          #printing the top 10 positive features
          print("Top 10 Positive Features:-\n",df[1].sort values(ascending = Fals
          e)[0:10])
```

```
Top 10 Positive Features:-
         -3.720529
 not
        -4.530450
like
        -4.668716
good
        -4.755367
great
        -4.895308
one
        -4.959547
taste
        -5.006954
coffee
        -5.063586
flavor
        -5.072091
love
would
        -5.076085
Name: 1, dtype: float64
```

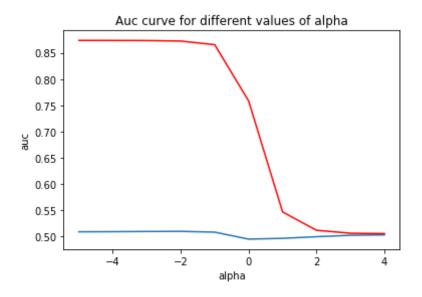
[5.1.2] Top 10 important features of negative class from SET 1

```
In [328]: # printing the top 10 negative features
         print("Top 10 negative Features:-\n",df[0].sort values(ascending = Fals
         e)[0:10])
         Top 10 negative Features:-
          not
                    -3.343000
         like
               -4.469759
         would
                 -4.732575
                 -4.759215
         taste
         product -4.781486
                  -4.960399
         one
         coffee -5.164503
                 -5.214759
         aood
         flavor
                   -5.232849
                   -5.259440
         no
         Name: 0, dtype: float64
```

[5.2] Applying Naive Bayes on TFIDF, SET 2

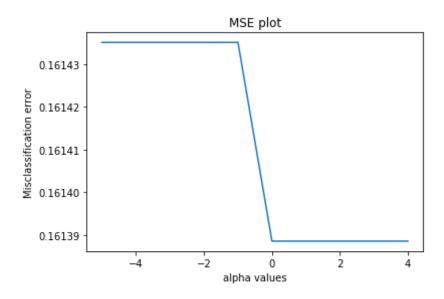
In [341]: #The below code is exactly same as the previous one with the only difference being that it is applied to TF-IDF vectorizer

```
auc=[]
          auc2=[]
          for i in alpha:
             model=MultinomialNB(alpha=i)
             model.fit(final tf idf,y tr)
              pred1=model.predict(X_cv_tfidf)
              probabl=model.predict proba(X cv tfidf)
              probab1=probab1[:,1]
             fpr,tpr,thresholds=metrics.roc curve(y cv,probab1)
             auc.append(roc auc score(y cv,probab1))
              acc=accuracy score(y cv,pred1,normalize=True)*float(100)
              pred4=model.predict(final tf idf)
              probab4=model.predict proba(final tf idf)
              probab4=probab4[:,1]
             fpr train,tpr train,thresholds=metrics.roc curve(y tr,probab4)
              auc2.append(roc auc score(y tr,probab4))
              print('Accuracy score at alpha value %d is %d'%(i,acc))
          plt.plot(np.log10(alpha),auc)
          plt.plot(np.log10(alpha),auc2,'r')
          plt.title('Auc curve for different values of alpha')
          plt.xlabel('alpha')
          plt.vlabel('auc')
          Accuracy score at alpha value 0 is 83
          Accuracy score at alpha value 1 is 83
         Accuracy score at alpha value 10 is 83
          Accuracy score at alpha value 100 is 83
          Accuracy score at alpha value 1000 is 83
         Accuracy score at alpha value 10000 is 83
Out[341]: Text(0,0.5,'auc')
```



```
In [342]: # Please write all the code with proper documentation
    cv_scores=[]
    alpha=[0.00001,0.0001,0.001,0.1,1,10,100,1000,10000]
    for i in alpha:
        model=MultinomialNB(alpha=i)
        scores=cross_val_score(model,final_tf_idf,y_tr,cv=10,scoring='accuracy')
        cv_scores.append(scores.mean())
    MSE=[1-x for x in cv_scores]
    plt.plot(np.log10(alpha),MSE)
    plt.title('MSE plot')
    plt.xlabel('alpha values')
    plt.ylabel('Misclassification error')
    optimal_alpha=alpha[MSE.index(min(MSE))]
    print('The optimal alpha vale for low error is=%d'%optimal_alpha)
```

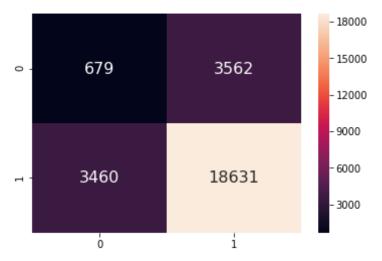
The optimal alpha vale for low error is=1

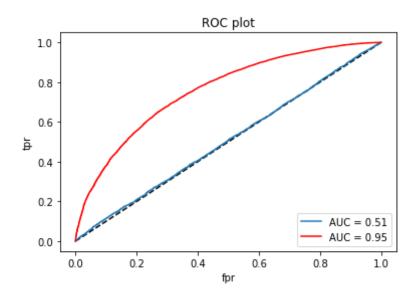


```
In [343]: model=MultinomialNB(alpha=optimal_alpha)
    model.fit(final_tf_idf,y_tr)
    predict1=model.predict(X_test_tfidf)
    probab1=model.predict_proba(X_test_tfidf)
    probab1=probab1[:,1]
    predict3=model.predict(final_tf_idf)
    probab3=model.predict_proba(final_tf_idf)
    probab3=probab3[:,1]
    acc=accuracy_score(y_test,predict1,normalize=True)*float(100)
    print('The accuracy of the model with optimal value of alpha is=%d'%opt
    imal_alpha)
```

The accuracy of the model with optimal value of alpha is=1

```
fpr_train,tpr_train,thresholds=roc_curve(y_tr,probab3)
auc3=roc_auc_score(y_tr,probab2)
plt.plot([0,1],[0,1],'k--')
plt.plot(fpr,tpr,label = 'AUC = %0.2f' %auc1)
plt.plot(fpr_train,tpr_train,'r',label = 'AUC = %0.2f' %auc3)
plt.title('ROC plot')
plt.xlabel('fpr')
plt.ylabel('tpr')
plt.legend(loc='lower right')
plt.show()
```





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

```
In [338]: feature log prob tfidf=model.feature log prob
          #print(feature log prob.shape)
          df=pd.DataFrame(feature log prob tfidf)
          #print(df.shape)
          tfidf_features=tf_idf_vect.get_feature_names()
          dfl=pd.DataFrame(feature log prob tfidf,columns=tfidf features)
          #print(df1.shape)
          #print(df.head())
          df1=df1.T
          #print(df1.head())
          print("Top 10 Positive Features:-\n",df1[1].sort values(ascending = Fal
          se)[0:10])
          Top 10 Positive Features:-
           not
                     -4.999648
          like
                    -5.551137
                    -5.583325
          good
                    -5.597252
          great
```

```
coffee -5.624187
taste -5.753221
product -5.753313
tea -5.771656
one -5.809017
love -5.812956
Name: 1, dtype: float64
```

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

```
In [339]: # Please write all the code with proper documentation
         print("Top 10 negative Features:-\n",df1[0].sort values(ascending = Fal
         se)[0:10])
         Top 10 negative Features:-
                   -5.016674
          not
         like -5.555650
         great -5.564681
                 -5.579038
         good
                 -5.596584
         coffee
         product -5.727963
                 -5.751141
         taste
                -5.760627
         tea
                 -5.788780
         love
              -5.820209
         one
         Name: 0, dtype: float64
```

[6] Conclusions

```
In [340]: # Please compare all your models using Prettytable library
    from prettytable import PrettyTable
    x=PrettyTable()
    x.field_names=['Vectorizer','Model','Hyperparameter','AUC']
    x.add_row(["Bag Of Words","Multinomial Naive bayes",1,0.91])
    x.add_row(["TF-IDF","Multinomial Naive Bayes",0.01,0.48])
    print(x)
```

Vectorizer	Model	Hyperparameter	AUC
•	Multinomial Naive bayes Multinomial Naive Bayes	1 1 0.01	0.91 0.95