Amazon Fine Food Reviews Analysis

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

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

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

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

Number of Attributes/Columns in data: 10

Attribute Information:

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

Objective:

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

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

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

[1]. Reading Data

[1.1] Loading the data

The dataset is available in two forms

- 1. .csv file
- 2. SQLite Database

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

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

In [1]:

```
%matplotlib inline
import warnings
warnings.filterwarnings("ignore")

import sqlite3
import pandas as pd
import numpy as np
import nltk
import string
import matplotlib.pyplot as plt
```

```
import seaborn as sns
from sklearn.feature extraction.text import TfidfTransformer
from sklearn.feature_extraction.text import TfidfVectorizer
from sklearn.feature extraction.text import CountVectorizer
from sklearn.metrics import confusion matrix
from sklearn import metrics
from sklearn.metrics import roc curve, auc
from nltk.stem.porter import PorterStemmer
import re
# Tutorial about Python regular expressions: https://pymotw.com/2/re/
import string
from nltk.corpus import stopwords
from nltk.stem import PorterStemmer
from nltk.stem.wordnet import WordNetLemmatizer
from gensim.models import Word2Vec
from gensim.models import KeyedVectors
import pickle
from tqdm import tqdm
import os
C:\Users\lenovo\Anaconda3\lib\site-packages\gensim\utils.py:1197: UserWarning: detected Windows; a
liasing chunkize to chunkize_serial
 warnings.warn("detected Windows; aliasing chunkize to chunkize_serial")
In [2]:
# using SQLite Table to read data.
con = sqlite3.connect('database.sqlite')
# filtering only positive and negative reviews i.e.
# not taking into consideration those reviews with Score=3
# SELECT * FROM Reviews WHERE Score != 3 LIMIT 500000, will give top 500000 data points
```

```
# you can change the number to any other number based on your computing power
# filtered data = pd.read sql query(""" SELECT * FROM Reviews WHERE Score != 3 LIMIT 500000""", co
n)
# for tsne assignment you can take 5k data points
filtered_data = pd.read_sql_query(""" SELECT * FROM Reviews WHERE Score != 3""", con)
# Give reviews with Score>3 a positive rating, and reviews with a score<3 a negative rating.
def partition(x):
    if x < 3:
       return 'negative'
    return 'positive'
#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)
print("Data Points in Each class :")
print(filtered data['Score'].value counts())
filtered data.head(3)
```

Number of data points in our data (525814, 10)
Data Points in Each class:
positive 443777
negative 82037
Name: Score, dtype: int64

Out[2]:

Id ProductId UserId ProfileName HelpfulnessNumerator HelpfulnessDenominator Score Time Summary

	ld	ProductId	UserId	ProfileName	HelpfulnessNumerator	HelpfulnessDenominator	Score	Time	Summary
1	2	B00813GRG4	A1D87F6ZCVE5NK	dll pa	0	0	negative	1346976000	Not as Advertisec
2	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1	1	positive	1219017600	"Delight' says it al
4									Þ

[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 [3]:

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

Out[3]:

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

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 Productld belonged to the same product just having different flavour or quantity. Hence in order to reduce redundancy it was decided to eliminate the rows having same parameters.

The method used for the same was that we first sort the data according to ProductId and then just keep the first similar product review and delelte the others. for eg. in the above just the review for ProductId=B000HDL1RQ remains. This method ensures that there is only one representative for each product and deduplication without sorting would lead to possibility of different representatives still existing for the same product.

```
In [4]:
```

```
#Sorting the data taking productid as the parameter
sorted data=filtered data.sort values('ProductId', axis=0, ascending=True, inplace=False, kind='qui
cksort', na_position='last')
sorted_data.shape
Out[4]:
(525814, 10)
In [5]:
#Deleting the dublicates reviews which is created when user writed a review for the product, it au
tomatically generates for the same product of different color etc
final = sorted data.drop duplicates(subset={"UserId","ProfileName","Time","Text"}, keep='first', in
place=False)
final.shape
Out[5]:
(364173, 10)
In [6]:
#Checking to see how much % of data still remains
(final['Id'].size*1.0)/(filtered data['Id'].size*1.0)*100
Out[6]:
69.25890143662969
In [7]:
display= pd.read_sql_query("""
SELECT *
FROM Reviews
WHERE Score != 3 AND Id=44737 OR Id=64422
ORDER BY ProductID
""", con)
display.head()
Out[7]:
```

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	HelpfulnessDenominator	Score	Time	Summary
C	64422	B000MIDROQ	A161DK06JJMCYF	J. E. Stephens "Jeanne"	3	1	5	1224892800	Bought This for My Son at College
1	44737	B001EQ55RW	A2V0I904FH7ABY	Ram	3	2	4	1212883200	Pure cocoa taste with crunchy almonds inside
4									Þ

In [8]:

#Dropping the data which has HelpfulnessNumerator<HelpfulnessDenominator which is impossible final=final[final.HelpfulnessNumerator<=final.HelpfulnessDenominator]

#Before starting the next phase of preprocessing lets see the number of entries left
print(final shape)

```
brinc (rinar. snabe)
#How many positive and negative reviews are present in our dataset?
final['Score'].value_counts()
(364171, 10)
Out[8]:
positive
          307061
negative
           57110
Name: Score, dtype: int64
In [9]:
#Checking to see how much % of data still remains
print("Percentage of data still remains", (final['Id'].size*1.0)/(filtered data['Id'].size*1.0)*100)
print("Final Data", final.shape)
Percentage of data still remains 69.25852107399194
Final Data (364171, 10)
```

[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 [5]:

```
stop = set(stopwords.words('english')) #set of stopwords
sno = nltk.stem.SnowballStemmer('english') #initialising the snowball stemmer

def cleanhtml(sentence): #function to clean the word of any html-tags
    cleanr = re.compile('<.*?>')
    cleantext = re.sub(cleanr, ' ', sentence)
    return cleantext

def cleanpunc(sentence): #function to clean the word of any punctuation or special characters
    cleaned = re.sub(r'[?!!\\'"|#]',r'',sentence)
    cleaned = re.sub(r'[.|,|)|(|\\|/]',r' ',cleaned)
    return cleaned
```

In [11]:

```
#Code for implementing step-by-step the checks mentioned in the pre-processing phase
# this code takes a while to run as it needs to run on 500k sentences.
from tqdm import tqdm
i=0
strl=' '
final_string=[]
all_positive_words=[] # store words from +ve reviews here
all_negative_words=[] # store words from -ve reviews here.
s=''
for sent in tqdm(final['Text'].values):
```

```
#print(sent);
    sent=cleanhtml(sent) # remove HTMl tags
    for w in sent.split():
        for cleaned words in cleanpunc(w).split():
            if((cleaned words.isalpha()) & (len(cleaned words)>2)):
                 if(cleaned_words.lower() not in stop):
                     s=(sno.stem(cleaned_words.lower())).encode('utf8')
                     filtered sentence.append(s)
                     if (final['Score'].values)[i] == 'positive':
                         all positive words.append(s) \#list of all words used to describe positive r
eviews
                     if(final['Score'].values)[i] == 'negative':
                         all negative words.append(s) #list of all words used to describe negative r
eviews reviews
                 else:
                     continue
            else:
                continue
    #print(filtered_sentence)
    str1 = b" ".join(filtered_sentence) #final string of cleaned words
    #print("*
    final_string.append(str1)
    i+=1
4
                                                                              364171/364171
100%|
[09:06<00:00, 665.88it/s]
In [12]:
final['CleanedText']=final_string #adding a column of CleanedText which displays the data after pr
e-processing of the review
final['CleanedText']=final['CleanedText'].str.decode("utf-8")
final.head(3)
Out[12]:
          ld
               ProductId
                                UserId ProfileName HelpfulnessNumerator HelpfulnessDenominator
                                                                                                  Time
                                                                                                        S
                                                                                       Score
                                            shari
138706 150524 0006641040
                          ACITT7DI6IDDI
                                                                0
                                                                                    0 positive
                                                                                              939340800
                                         zychinski
                                                                                                       edi
138688 150506 0006641040 A2IW4PEEKO2R0U
                                            Tracy
                                                                                    1 positive 1194739200
                                          sally sue
138689 150507 0006641040 A1S4A3IQ2MU7V4
                                                                 1
                                                                                    1 positive 1191456000
                                         "sally sue"
In [13]:
final data=final.sort values('Time', axis=0, ascending=True, inplace=False, kind='quicksort', na po
sition='last')
In [3]:
final = final data.head(100000)
X train data = final[:60000]
X test data = final[60000:100000]
y_train = X_train_data['Score']
y test = X test data['Score']
```

filtered sentence=[]

```
print("Data")
print(X_train_data.shape)
print(X test data.shape)
print("Label")
print(y_train.shape)
print(y_test.shape)
Data
(60000, 11)
(40000, 11)
(60000,)
(40000,)
In [4]:
X test data['Score'].value counts()
Out[4]:
positive 34582
negative
           5418
Name: Score, dtype: int64
```

[3.2] Preprocessing Review Summary

```
In [16]:
```

```
## Similartly you can do preprocessing for review summary also.
```

[4] Featurization

[4.1] BAG OF WORDS

```
In [11]:

#BoW on Text
print("**Bow Vectorizer**")
print("="*50)
count_vect = CountVectorizer()
X_train_BoW = count_vect.fit_transform(X_train_data['CleanedText'])
X_test_BoW = count_vect.transform(X_test_data['CleanedText'])
print(X_train_BoW.shape)
print(X_test_BoW.shape)
**Bow Vectorizer**
```

(60000, 29132) (40000, 29132)

[4.2] Bi-Grams and n-Grams.

```
In [ ]:
```

[4.3] TF-IDF

```
In [33]:
```

```
#TFIDF on Text
print("**TFIDF Vectorizer**")
```

[4.4] Word2Vec

```
In [6]:
```

In [7]:

In [8]:

is', 'is', 'a', 'classic', 'book', 'i', 'am', 'willing', 'to', 'bet', 'my', 'son', 'will', 'still', 'be', 'able', 'to', 'recite', 'from', 'memory', 'when', 'he', 'is', 'in', 'college']

```
w2v model=gensim.models.Word2Vec(list of sent train,min count=5,size=50, workers=6)
In [10]:
w2v words = list(w2v model.wv.vocab)
print(len(w2v_words))
14907
In [11]:
w2v model.wv.most similar('good')
Out[11]:
[('great', 0.8218839168548584),
 ('fantastic', 0.7583197951316833),
 ('decent', 0.7440043687820435),
 ('fine', 0.7032303810119629),
 ('amazing', 0.6977575421333313),
 ('wonderful', 0.6968648433685303),
 ('yummy', 0.6901988983154297),
 ('bad', 0.6877514123916626),
 ('terrific', 0.6834275722503662),
 ('tasty', 0.6785352230072021)]
In [12]:
w2v model.wv.most similar('tasty')
Out[12]:
[('satisfying', 0.8392266631126404),
 ('filling', 0.8139099478721619),
 ('yummy', 0.806728720664978),
 ('delicious', 0.7997151017189026),
 ('tastey', 0.7370927333831787),
 ('flavorful', 0.7332741022109985),
 ('light', 0.7140116691589355),
 ('versatile', 0.7100734710693359),
 ('hearty', 0.6937506794929504),
 ('sweet', 0.6828324794769287)]
In [13]:
w2v model.wv.most similar('horrible')
Out[13]:
[('terrible', 0.9139878749847412),
 ('awful', 0.8891698122024536),
 ('disgusting', 0.8430461883544922),
 ('funny', 0.8422303199768066),
 ('weird', 0.8200281858444214),
 ('gross', 0.7890602350234985),
 ('strange', 0.7720152139663696),
 ('nasty', 0.7229914665222168),
 ('bad', 0.7111765146255493),
 ('ok', 0.7051739692687988)]
```

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

[4.4.1.1] Avg W2v

In [9]:

```
III [14]:
#TRATN
# average Word2Vec
# compute average word2vec for each review.
sent vectors train = [];
for sent in tqdm(list of sent train):
   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
    sent_vectors_train.append(sent_vec)
print(len(sent vectors train))
print(len(sent_vectors_train[0]))
100%|
                                                                                  | 60000/60000 [03:
25<00:00, 291.40it/s]
60000
50
In [15]:
# average Word2Vec
# compute average word2vec for each review.
sent vectors test = [];
for sent in tqdm(list_of_sent_test):
   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
    sent vectors test.append(sent vec)
print(len(sent vectors test))
print(len(sent_vectors_test[0]))
100%|
                                                                                  | 40000/40000 [02:
28<00:00, 268.90it/s]
40000
50
[4.4.1.2] TFIDF weighted W2v
In [16]:
tfidf vect = TfidfVectorizer(min df = 50)
train tfidf w2v = tfidf vect.fit transform(X train data["CleanedText"])
test tfidf w2v = tfidf vect.transform(X test data["CleanedText"])
dictionary = dict(zip(tfidf_vect.get_feature_names(), list(tfidf_vect.idf_)))
print(train tfidf w2v.shape)
print(test tfidf w2v.shape)
(60000, 2951)
(40000, 2951)
In [17]:
# TF-IDF weighted Word2Vec
tfidf_feat = tfidf_vect.get_feature_names() # tfidf words/col-names
```

```
# final tf idf is the sparse matrix with row= sentence, col=word and cell val = tfidf
tfidf sent vectors train = []; # the tfidf-w2v for each sentence/review is stored in this list
row=0;
for sent in tqdm(list_of sent train): # for each review/sentence
   sent vec = np.zeros(50) # as word vectors are of zero length
    weight sum =0; # num of words with a valid vector in the sentence/review
   for word in sent: # for each word in a review/sentence
       if word in w2v words and word in tfidf feat:
           vec = w2v_model.wv[word]
             tf idf = tf idf matrix[row, tfidf feat.index(word)]
            # to reduce the computation we are
            # dictionary[word] = idf value of word in whole courpus
            # sent.count(word) = tf valeus of word in this review
            tf idf = dictionary[word] * (sent.count(word) /len(sent))
            sent vec += (vec * tf idf)
           weight sum += tf idf
    if weight sum != 0:
       sent vec /= weight sum
    tfidf_sent_vectors_train.append(sent_vec)
    row += 1
100%|
                                                                          | 60000/60000 [07:
15<00:00, 137.74it/s]
```

In [18]:

```
# TF-IDF weighted Word2Vec
tfidf_feat = tfidf_vect.get_feature_names() # tfidf words/col-names
# final tf idf is the sparse matrix with row= sentence, col=word and cell val = tfidf
tfidf_sent_vectors_test = []; # the tfidf-w2v for each sentence/review is stored in this list
for sent in tqdm(list of sent test): # for each review/sentence
   sent_vec = np.zeros(50) # as word vectors are of zero length
    weight sum =0; # num of words with a valid vector in the sentence/review
    for word in sent: # for each word in a review/sentence
        if word in w2v words and word in tfidf_feat:
            vec = w2v model.wv[word]
             tf idf = tf idf matrix[row, tfidf feat.index(word)]
            # to reduce the computation we are
            # dictionary[word] = idf value of word in whole courpus
            # sent.count(word) = tf valeus of word in this review
            tf idf = dictionary[word] * (sent.count (word) /len(sent))
            sent vec += (vec * tf idf)
            weight sum += tf idf
    if weight sum != 0:
       sent vec /= weight sum
    tfidf sent_vectors_test.append(sent_vec)
100%|
                                                                          | 40000/40000 [05:
13<00:00, 127.56it/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 AUC value
 - Find the best hyper paramter using k-fold cross validation or simple cross validation data
 - Use gridsearch cv or randomsearch cv or you can also write your own for loops to do this task of hyperparameter tuning

3. 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</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.
- Along with plotting ROC curve, you need to print the <u>confusion matrix</u> with predicted and original labels of test data points. Please visualize your confusion matrices using <u>seaborn heatmaps</u>.

7. 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.

In [21]:

```
from sklearn.tree import DecisionTreeClassifier
from sklearn.metrics import roc_auc_score
import scikitplot as skplt
from cycler import cycler
from sklearn.model_selection import GridSearchCV
from sklearn import svm
from sklearn.metrics import
accuracy_score,precision_score,recall_score,confusion_matrix,classification_report,fl_score
```

Applying Decision Trees

[5.1] Applying Decision Trees on BOW, SET 1

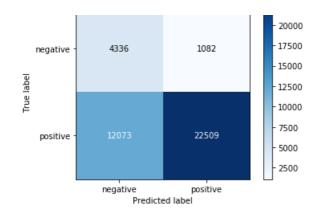
```
In [58]:
```

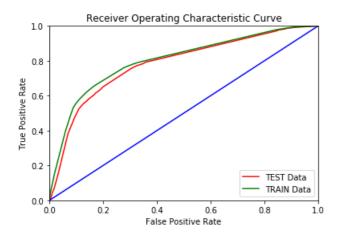
```
X_train = X_train_BOW
X_test = X_test_BOW
```

```
max depths = [3,4,5,6,7,8,9,10,11]
min_split = [5, 10, 100, 500]
param grid = {'max depth': max depths,'min samples split':min split}
model = GridSearchCV(DecisionTreeClassifier(), param grid, scoring = 'roc auc', cv=3 , n jobs = -1)
model.fit(X train, y train)
print("Model with best parameters :\n", model.best estimator )
print("Accuracy of the model : ", model.score(X train, y train))
Model with best parameters :
 DecisionTreeClassifier(class weight=None, criterion='gini', max depth=11,
            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')
Accuracy of the model : 0.7688922695958512
In [60]:
optimal depth = 11
optimal split = 500
In [61]:
lr = DecisionTreeClassifier(max depth=optimal depth, min samples split =optimal split,
class weight='balanced')
lr.fit(X train BOW, y train)
pred = lr.predict(X test BOW)
print("***Test Data Report***")
print("Best max_depth = ",optimal_depth)
print("Best min samples split = ",optimal split)
```

```
fpr, tpr, threshold = metrics.roc curve(y test, lr.predict proba(X test)[:,1],pos label="positive")
auc = metrics.auc(fpr, tpr)
print("AUC = ",auc*100)
skplt.metrics.plot confusion matrix(y test, pred)
plt.show()
fpr, tpr, threshold = metrics.roc curve(y test, lr.predict proba(X test)[:,1],pos label="positive")
fpr2, tpr2, threshold2 = metrics.roc curve(y train, lr.predict proba(X train)
[:,1],pos label="positive")
roc auc = metrics.auc(fpr, tpr)
roc_auc2 = metrics.auc(fpr2, tpr2)
# method I: plt
import matplotlib.pyplot as plt
f, ax = plt.subplots()
plt.title('Receiver Operating Characteristic Curve')
cy = cycler('color', ['red', 'green', 'blue'])
ax.set prop cycle(cy)
ax.plot(fpr, tpr, label = 'AUC = %0.2f' % roc auc)
ax.plot(fpr2, tpr2, label = 'AUC = %0.2f' % roc auc2)
plt.legend(['TEST Data', 'TRAIN Data'], loc = 'lower right')
ax.plot([0, 1], [0, 1])
plt.xlim([0, 1])
plt.ylim([0, 1])
plt.ylabel('True Positive Rate')
plt.xlabel('False Positive Rate')
plt.show()
```

```
***Test Data Report***
Best max_depth = 11
Best min_samples_split = 500
AUC = 78.17512034620545
```





[5.1.1] Top 20 important features from SET 1

In [62]:

```
feature_name = count_vect.get_feature_names()
w = lr.feature_importances_
weight=w.reshape(-1)
sorted_feature = np.argsort(weight)
top_20_positive_feature=sorted_feature[:-20:-1]
```

In [63]:

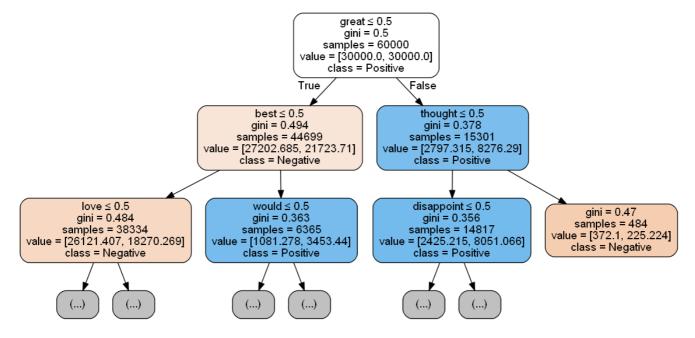
```
print("Top 20 features :")
print("-----")
for i in top_20_positive_feature:
    print("%s\t-->\t%f"%(feature_name[i], weight[i]))
```

```
Top 20 features :
great --> 0.186790
best --> 0.113268
disappoint --> 0.090686
love --> 0.089791
delici --> 0.071927
excel --> 0.041670
perfect --> 0.035028
favorit --> 0.034536
good --> 0.028687
bad --> 0.024647
thought --> 0.024129
nice --> 0.018794
wonder --> 0.015806
howev --> 0.015724
tast --> 0.014607
would --> 0.014598
product --> 0.012589
hope --> 0.012294
terribl --> 0.011665
```

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

In [64]:

Out[64]:



[5.2] Applying Decision Trees on TFIDF, SET 2

```
In [66]:
```

```
X_train = X_train_tfidf
X_test = X_test_tfidf
```

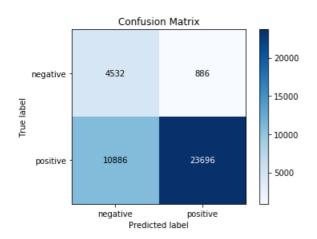
```
In [67]:
max depths = [3,4,5,6,7,8,9,10,11]
min_split = [5, 10, 100, 500]
param grid = {'max depth': max depths,'min samples split':min split}
model = GridSearchCV(DecisionTreeClassifier(), param grid, scoring = 'roc auc', cv=3, n jobs = -1)
model.fit(X_train, y_train)
print("Model with best parameters : \n", model.best estimator)
print("Accuracy of the model : ", model.score(X train, y train))
Model with best parameters :
 DecisionTreeClassifier(class_weight=None, criterion='gini', max_depth=11,
            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')
Accuracy of the model: 0.7561631280345095
```

In [68]: optimal_depth = 11 optimal_split = 500

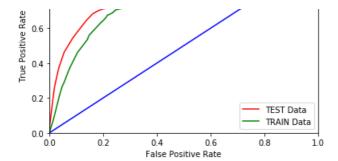
In [69]:

```
lr = DecisionTreeClassifier(max depth=optimal depth, min samples split =optimal split,
class weight='balanced')
lr.fit(X_test,y_test)
pred = lr.predict(X_test)
print("***Test Data Report***")
print("Best max depth = ",optimal depth)
print("Best min samples split = ",optimal split)
fpr, tpr, threshold = metrics.roc_curve(y_test, lr.predict_proba(X_test)[:,1],pos_label="positive")
auc = metrics.auc(fpr, tpr)
print("AUC = ",auc*100)
skplt.metrics.plot_confusion_matrix(y_test, pred)
plt.show()
fpr, tpr, threshold = metrics.roc_curve(y_test, lr.predict_proba(X_test)[:,1],pos_label="positive")
fpr2, tpr2, threshold2 = metrics.roc curve(y train, lr.predict proba(X train)
[:,1],pos_label="positive")
roc auc = metrics.auc(fpr, tpr)
roc_auc2 = metrics.auc(fpr2, tpr2)
# method I: plt
import matplotlib.pyplot as plt
f, ax = plt.subplots()
plt.title('Receiver Operating Characteristic Curve')
cy = cycler('color', ['red', 'green', 'blue'])
ax.set_prop_cycle(cy)
ax.plot(fpr, tpr, label = 'AUC = %0.2f' % roc_auc)
ax.plot(fpr2, tpr2, label = 'AUC = %0.2f' % roc auc2)
plt.legend(['TEST Data', 'TRAIN Data'],loc = 'lower right')
ax.plot([0, 1], [0, 1])
plt.xlim([0, 1])
plt.ylim([0, 1])
plt.ylabel('True Positive Rate')
plt.xlabel('False Positive Rate')
plt.show()
```

Test Data Report
Best max_depth = 11
Best min_samples_split = 500
AUC = 80.45766361745706







[5.2.1] Top 20 important features from SET 2

```
In [70]:
```

```
feature_name = tf_idf_vect.get_feature_names()
w = lr.feature_importances_
weight=w.reshape(-1)
sorted_feature = np.argsort(weight)
top_20_positive_feature=sorted_feature[:-20:-1]
```

In [71]:

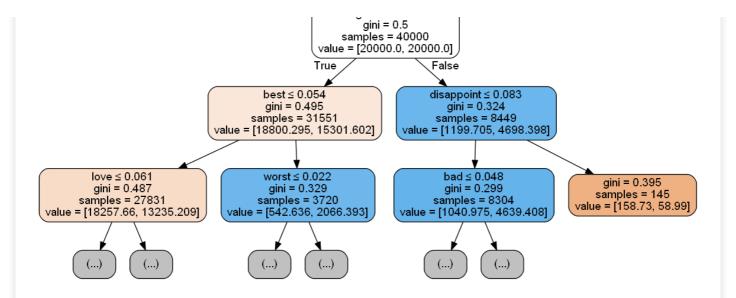
```
print("Top 20 features :")
print("----")
for i in top_20_positive_feature:
    print("%s\t-->\t%f"%(feature_name[i], weight[i]))
```

```
Top 20 features:
great --> 0.201859
love --> 0.109540
best --> 0.105428
disappoint --> 0.090735
delici --> 0.075654
favorit --> 0.045947
good --> 0.039681
excel --> 0.033474
perfect --> 0.032291
easi --> 0.021314
thought --> 0.021107
bad --> 0.020806
amaz --> 0.019545
stale --> 0.018004
worst --> 0.009417
away --> 0.009355
wast --> 0.008710
tast --> 0.008478
would --> 0.007649
```

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

In [72]:

Out[72]:



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

```
In [88]:
```

```
X_train = sent_vectors_train
X_test = sent_vectors_test
```

In [89]:

```
max_depths = [3,4,5,6,7,8,9,10,11]
min_split = [5, 10, 100, 500]
param_grid = {'max_depth': max_depths, 'min_samples_split':min_split}

model = GridSearchCV(DecisionTreeClassifier(), param_grid, scoring = 'roc_auc', cv=3, n_jobs = -1)
model.fit(X_train, y_train)
print("Model with best parameters :\n", model.best_estimator_)
print("Accuracy of the model : ", model.score(X_train, y_train))
```

Accuracy of the model : 0.8346600084783115

In [90]:

```
optimal_depth = 9
optimal_split = 500
```

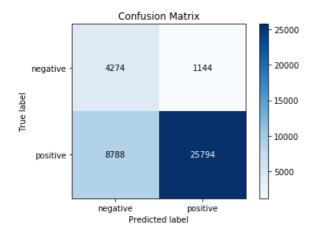
In [91]:

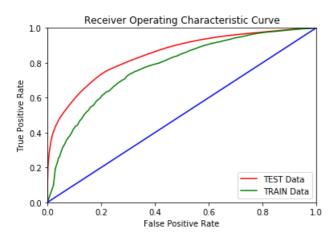
```
lr = DecisionTreeClassifier(max_depth=optimal_depth, min_samples_split =optimal_split,
    class_weight='balanced')
lr.fit(X_test,y_test)
pred = lr.predict(X_test)

print("***Test Data Report***")
print("Best max_depth = ",optimal_depth)
print("Best min_samples_split = ",optimal_split)
fpr, tpr, threshold = metrics.roc_curve(y_test, lr.predict_proba(X_test)[:,1],pos_label="positive")
auc = metrics.auc(fpr, tpr)
print("AUC = ",auc*100)
skplt.metrics.plot_confusion_matrix(y_test, pred)
plt.show()
```

```
fpr, tpr, threshold = metrics.roc_curve(y_test, lr.predict_proba(X_test)[:,1],pos_label="positive")
fpr2, tpr2, threshold2 = metrics.roc curve(y train, lr.predict proba(X train)
[:,1],pos label="positive")
roc auc = metrics.auc(fpr, tpr)
roc_auc2 = metrics.auc(fpr2, tpr2)
# method I: plt
import matplotlib.pyplot as plt
f, ax = plt.subplots()
plt.title('Receiver Operating Characteristic Curve')
cy = cycler('color', ['red', 'green', 'blue'])
ax.set_prop_cycle(cy)
ax.plot(fpr, tpr, label = 'AUC = %0.2f' % roc auc)
ax.plot(fpr2, tpr2, label = 'AUC = %0.2f' % roc auc2)
plt.legend(['TEST Data', 'TRAIN Data'],loc = 'lower right')
ax.plot([0, 1], [0, 1])
plt.xlim([0, 1])
plt.ylim([0, 1])
plt.ylabel('True Positive Rate')
plt.xlabel('False Positive Rate')
plt.show()
```

```
***Test Data Report***
Best max_depth = 9
Best min_samples_split = 500
AUC = 84.5876684749206
```





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

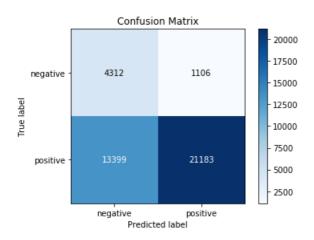
In [34]:

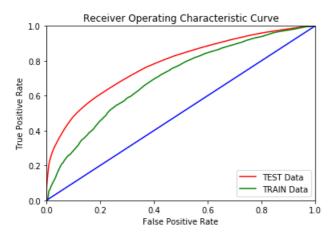
```
X_train = tfidf_sent_vectors_train
X_test = tfidf_sent_vectors_test
```

```
In [26]:
max depths = [3,4,5,6,7,8,9,10,11]
min split = [5, 10, 100, 500]
param_grid = {'max_depth': max_depths,'min_samples_split':min_split}
model = GridSearchCV(DecisionTreeClassifier(), param grid, scoring = 'roc auc', cv=3 , n jobs = -1)
model.fit(X_train, y_train)
 print("Model with best parameters :\n", model.best_estimator_)
print("Accuracy of the model : ", model.score(X_train, y_train))
Model with best parameters :
  {\tt DecisionTreeClassifier(class\_weight=None, criterion='gini', max\_depth=8, criterion='gini', max_depth=8, criterion='gini
                            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')
Accuracy of the model : 0.7623964663907726
In [27]:
optimal depth = 8
optimal split = 500
In [35]:
lr = DecisionTreeClassifier(max depth=optimal depth, min samples split =optimal split,
 class weight='balanced')
lr.fit(X_test,y_test)
pred = lr.predict(X test)
print("***Test Data Report***")
print("Best max_depth = ",optimal_depth)
 print("Best min samples split = ",optimal split)
 fpr, tpr, threshold = metrics.roc curve(y test, lr.predict proba(X test)[:,1],pos label="positive")
auc = metrics.auc(fpr, tpr)
print("AUC = ", auc*100)
skplt.metrics.plot_confusion_matrix(y_test, pred)
plt.show()
 fpr, tpr, threshold = metrics.roc_curve(y_test, lr.predict_proba(X_test)[:,1],pos_label="positive")
 fpr2, tpr2, threshold2 = metrics.roc_curve(y_train, lr.predict_proba(X_train)
 [:,1],pos label="positive")
 roc auc = metrics.auc(fpr, tpr)
```

```
roc auc2 = metrics.auc(fpr2, tpr2)
# method I: plt
import matplotlib.pyplot as plt
f, ax = plt.subplots()
plt.title('Receiver Operating Characteristic Curve')
cy = cycler('color', ['red', 'green', 'blue'])
ax.set prop cycle(cy)
ax.plot(fpr, tpr, label = 'AUC = %0.2f' % roc_auc)
ax.plot(fpr2, tpr2, label = 'AUC = %0.2f' % roc auc2)
plt.legend(['TEST Data', 'TRAIN Data'], loc = 'lower right')
ax.plot([0, 1], [0, 1])
plt.xlim([0, 1])
plt.ylim([0, 1])
plt.ylabel('True Positive Rate')
plt.xlabel('False Positive Rate')
plt.show()
***Test Data Report***
Best max_depth = 8
Best min_samples_split =
```

AUC = 77.87856192734452





[6] Conclusions

In [36]:

```
#importing library
from prettytable import PrettyTable
x = PrettyTable()

#adding Field names
x.field_names = ["SL No.","Vectorizer","Best (max_depth)","Best (min_samples_split)","AUC"]

# adding row to table
x.title = 'Decision Trees'
x.add_row(["1","BoW",11,500,78.1751])
x.add_row(["2","TFIDF",11,500,80.4577])
x.add_row(["3","Avg-W2vec",9,500,84.5877])
x.add_row(["4","TFIDF-W2vec",8,500,77.8786])

#printing the table
print(x)
```

SL No.	Vectorizer	Best (max_depth)	Best (min_samples_split)	AUC
1	BOW TFIDF Avg-W2vec TFIDF-W2vec	11 11 11 9 8	500 500 500 500	78.1751 80.4577 84.5877 77.8786

Conclusion

- AVG W2VECgives the best AUC Score amoung the models on test data i.e 84.5877
- TFIDF-W2vec gives the the least AUC Score among the models.
- Decision trees models have lower accuracy rates as compared to other models like logistic regression, naive bayes