```
In [1]: # Exercise: t-SNE visualization of Amazon reviews with polarity based c
        olor-coding
        %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
```

```
# using the sqllite table to read data
con = sqlite3.connect(r'G:\machine_learning\Real world problem Predict
rating given product reviews on Amazon\amazon\database.sqlite')

# filtering only positive and negative i.e not taking those review with
score = 3

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

C:\Users\hemant\Anaconda\lib\site-packages\gensim\utils.py:1209: UserWa
rning: detected Windows; aliasing chunkize to chunkize_serial
    warnings.warn("detected Windows; aliasing chunkize to chunkize_seria
l")
```

In [2]: filtered_data.head(5)

Out[2]:

| | ld | ProductId | Userld | ProfileName | HelpfulnessNumerator | Helpfulnes |
|---|----|------------|----------------|-------------|----------------------|------------|
| C | 1 | B001E4KFG0 | A3SGXH7AUHU8GW | delmartian | 1 | 1 |
| 1 | 2 | B00813GRG4 | A1D87F6ZCVE5NK | dli pa | 0 | 0 |

| | ld | ProductId | Userld | ProfileName | HelpfulnessNumerator | Helpfulnes |
|---|----|------------|----------------|--|----------------------|------------|
| 2 | 3 | B000LQOCH0 | ABXLMWJIXXAIN | Natalia Corres "Natalia Corres" | 1 | 1 |
| 3 | 4 | B000UA0QIQ | A395BORC6FGVXV | Karl | 3 | 3 |
| 4 | 5 | B006K2ZZ7K | A1UQRSCLF8GW1T | Michael D. Bigham "M. Wassir" | 0 | 0 |

```
In [3]: # give reviews with score>3 a postive rating and reviews with score <3
    negative rating

def partition(x):
    if x > 3:
        return 'positive'
    else:
        return 'negative'
    positiveNegative = filtered_data['Score']
```

actual = positiveNegative.map(partition)
filtered_data['Score'] = actual

In [4]: filtered_data.head(5)

Out[4]:

| | ld | ProductId | Userld | ProfileName | HelpfulnessNumerator | Helpfulnes |
|---|----|------------|----------------|--|----------------------|------------|
| 0 | 1 | B001E4KFG0 | A3SGXH7AUHU8GW | delmartian | 1 | 1 |
| 1 | 2 | B00813GRG4 | A1D87F6ZCVE5NK | dll pa | 0 | 0 |
| 2 | 3 | B000LQOCH0 | ABXLMWJIXXAIN | Natalia Corres "Natalia Corres" | 1 | 1 |
| 3 | 4 | B000UA0QIQ | A395BORC6FGVXV | Karl | 3 | 3 |

| | ld | ProductId | Userld | ProfileName | HelpfulnessNumerator | Helpfulnes |
|---|----|------------|----------------|-------------------------------------|----------------------|------------|
| 4 | 5 | B006K2ZZ7K | A1UQRSCLF8GW1T | Michael D. Bigham "M. Wassir" | 0 | 0 |

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 [5]: display = pd.read_sql_query("""select * from Reviews where Score != 3
    AND Userid = "AR5J8UI46CURR" order by ProductID """,con)
```

In [6]: display

Out[6]:

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

| | ld | ProductId | Userld | ProfileName | HelpfulnessNumerator | Helpfuln |
|---|--------|------------|---------------|--------------------|----------------------|----------|
| 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 can be seen above the same user has multiple reviews of the with the same values for HelpfulnessNumerator, HelpfulnessDenominator, Score, Time, Summary and Text and on doing analysis it was found that

ProductId=B000HDOPZG was Loacker Quadratini Vanilla Wafer Cookies, 8.82-Ounce Packages (Pack of 8)

ProductId=B000HDL1RQ was Loacker Quadratini Lemon Wafer Cookies, 8.82-Ounce Packages (Pack of 8) and so on

It was inferred after analysis that reviews with same parameters other than ProductId belonged to the same product just having different flavour or quantity. Hence in order to reduce redundancy it was decided to eliminate the rows having same parameters.

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

```
In [7]: # sorting data according to product id in ascending order
sorted_data = filtered_data.sort_values('ProductId',axis = 0,ascending
=True)
```

In [8]: sorted_data.head(5)

Out[8]:

| | ld | ProductId | Userld | ProfileName | HelpfulnessNumerator | Нє |
|--------|--------|------------|----------------|--------------------|----------------------|----|
| 138706 | 150524 | 0006641040 | ACITT7DI6IDDL | shari zychinski | 0 | 0 |
| 138688 | 150506 | 0006641040 | A2IW4PEEKO2R0U | Tracy | 1 | 1 |

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

```
In [9]: # deduplication of entries
  final = sorted_data.drop_duplicates(subset = {"UserId","ProfileName","T
    ime","Text"},keep = 'first',inplace = False)
```

```
In [12]: #Checking to see how much % of data still remains
  (final['Id'].size*1.0)/(filtered_data['Id'].size*1.0)*100
```

Out[12]: 69.25890143662969

Observation:- It was also seen that in two rows given below the value of HelpfulnessNumerator is greater than HelpfulnessDenominator which is not practically possible hence these two rows too are removed from calcualtions

```
In [13]: display= pd.read_sql_query("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND Id=44737 OR Id=64422
    ORDER BY ProductID
    """, con)
    display.head()
```

Out[13]:

| | ld | ProductId | Userld | ProfileName | HelpfulnessNumerator | Helpfuln |
|---|-------|------------|----------------|-------------------------------|----------------------|----------|
| 0 | 64422 | B000MIDROQ | A161DK06JJMCYF | J. E. Stephens "Jeanne" | 3 | 1 |

| | ld | ProductId | UserId | ProfileName | HelpfulnessNumerator | Helpfuln |
|---|-------|------------|----------------|-------------|----------------------|----------|
| 1 | 44737 | B001EQ55RW | A2V0I904FH7ABY | Ram | 3 | 2 |

In [14]: final=final[final.HelpfulnessNumerator<=final.HelpfulnessDenominator]</pre>

In [15]: #Before starting the next phase of preprocessing lets see the number of
 entries left
 print(final.shape)

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

(364171, 10)

Out[15]: positive 307061 negative 57110

Name: Score, dtype: int64

Text Preprocessing: Stemming, stop-word removal and Lemmatization. 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:-

Begin by removing the html tags Remove any punctuations or limited set of special characters like, or or # etc. Check if the word is made up of english letters and is not alpha-numeric 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) Convert the word to lowercase Remove Stopwords Finally Snowball Stemming the

word (it was obsereved to be better than Porter Stemming) After which we collect the words used to describe positive and negative reviews

```
In [16]: # find sentences containing HTML tags
import re
i=0;
for sent in final['Text'].values:
    if (len(re.findall('<.*?>', sent))):
        print(i)
        print(sent)
        break;
    i += 1;
```

I set aside at least an hour each day to read to my son (3 y/o). At this point, I consider myself a connoisseur of children's books and this is one of the best. Santa Clause put this under the tree. Since then, we e've read it perpetually and he loves it.

| '>

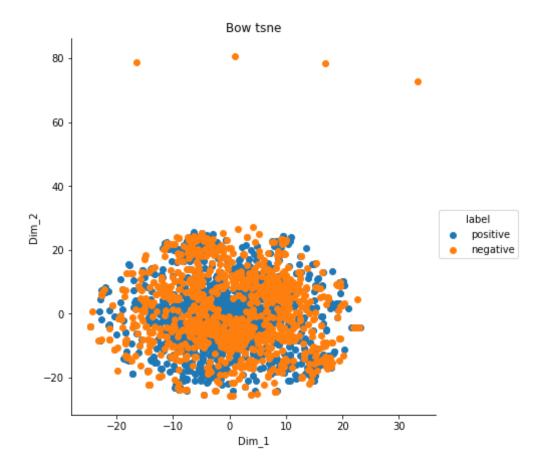
| '><

```
print('***********************************
         print(sno.stem('tasty'))
         {'they', 'can', 'while', 've', "that'll", "couldn't", 'needn', "she's",
         'a', 'him', "you'll", 'down', 'further', "you'd", "shan't", 'ourselve
         s', 'doing', 'if', 'before', 'then', 'ain', 'mightn', 'i', 'does', "do
         n't", 'yours', 'between', 'couldn', 'most', 'from', 'but', 'didn', 'her
         self', 'hers', 'more', "didn't", 'into', 'we', "shouldn't", 'weren', "h
         adn't", 'having', 'being', "haven't", 'both', 'there', 'been', 'wasn',
         'below', "should've", 'myself', 'and', 'those', 'have', 'under', 'o',
         'had', 'only', 'than', 's', 'haven', 'will', 'itself', 'our', 'their',
         'once', "isn't", 't', 'other', 'them', 'were', 'it', 'aren', 'has', 'n
         o', 'yourselves', 'as', 'how', 'very', 'because', 'all', 'not', 'does
         n', 'ours', 'y', 'with', 'such', "mustn't", "weren't", 'above', 'when',
         'shouldn', 'shan', "you're", 'be', 'after', 'about', 'off', "needn't",
         'up', 'me', 'mustn', "mightn't", 'of', 'that', 'now', 'll', 'her', 's
         o', 'd', "doesn't", 'was', 'too', 'did', 'isn', 'until', 'is', 'she',
         'wouldn', 'same', 'your', 'here', 'am', 'in', 'yourself', "wasn't", 'ju
         st', 'again', 'you', 'at', 'on', "wouldn't", 'm', 're', 'own', 'these',
         'himself', 'any', "won't", 'or', 'against', 'through', "you've", 'an',
         'why', 'my', 'his', 'do', "hasn't", 'won', 'should', 'hadn', 'ma', 'the
         irs', 'which', 'are', 'few', 'don', 'each', 'out', "it's", 'over', 'it
         s', 'themselves', 'by', 'the', 'this', 'whom', 'what', "aren't", 'has
         n', 'some', 'for', 'nor', 'who', 'where', 'during', 'he', 'to'}
         ***********
         tasti
In [18]: #Code for implementing step-by-step the checks mentioned in the pre-pro
         cessing phase
         # this code takes a while to run as it needs to run on 500k sentences.
         if not os.path.isfile('final.sqlite'):
             final string=[]
             all positive words=[] # store words from +ve reviews here
             all negative words=[] # store words from -ve reviews here.
             for i, sent in enumerate(tqdm(final['Text'].values)):
                 filtered sentence=[]
                 #print(sent);
                 sent=cleanhtml(sent) # remove HTMl tags
                 for w in sent.split():
```

```
# we have used cleanpunc(w).split(), one more split functio
n here because consider w="abc.def", cleanpunc(w) will return "abc def"
            # if we dont use .split() function then we will be considri
ng "abc def" as a single word, but if you use .split() function we will
get "abc", "def"
            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('utf
8')
                        filtered sentence.append(s)
                        if (final['Score'].values)[i] == 1:
                            all positive words.append(s) #list of all w
ords used to describe positive reviews
                        if(final['Score'].values)[i] == 0:
                            all negative words.append(s) #list of all w
ords used to describe negative reviews reviews
        str1 = b" ".join(filtered sentence) #final string of cleaned wo
rds
        #print(";
***************
       final string.append(str1)
    ############---- storing the data into .sqlite file -----#######
#################
    final['CleanedText']=final string #adding a column of CleanedText w
hich displays the data after pre-processing of the review
    final['CleanedText']=final['CleanedText'].str.decode("utf-8")
       # store final table into an SOLLite table for future.
    conn = sqlite3.connect('final.sqlite')
    c=conn.cursor()
    conn.text factory = str
    final.to sql('Reviews', conn, schema=None, if exists='replace', \
                 index=True, index label=None, chunksize=None, dtype=No
ne)
    conn.close()
```

```
with open('positive words.pkl', 'wb') as f:
                 pickle.dump(all positive words, f)
             with open('negitive words.pkl', 'wb') as f:
                 pickle.dump(all negative words, f)
In [19]: if os.path.isfile('final.sqlite'):
             conn = sqlite3.connect('final.sqlite')
             final = pd.read sql query(""" SELECT * FROM Reviews WHERE Score !=
          3 """, conn)
             conn.close()
         else:
             print("Please the above cell")
In [23]: random positive = final[final['Score']== 'positive'].sample(n = 1000)
         random negative = final[final['Score'] == 'negative'].sample(n = 1000)
In [24]: total 2000 = pd.concat([random positive,random negative])
In [25]: #BOW
         count vect = CountVectorizer() # in scikit learn
         final counts = count vect.fit transform(total 2000['CleanedText'].value
         print("the type of count vectorizer ",type(final counts))
         the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
In [26]: print("the shape of out text BOW vectorizer", final counts.get shape())
         print("the number of unique words", final counts.get shape()[1])
         the shape of out text BOW vectorizer (2000, 6461)
         the number of unique words 6461
In [27]: from sklearn.manifold import TSNE
         from sklearn.decomposition import TruncatedSVD
         import seaborn as sn
         SC = total 2000['Score']
```

```
#Dimensionality reduction using truncated SVD
svdT = TruncatedSVD(n components=390)
bowT = svdT.fit transform(final counts)
print(bowT.shape)
print(type(bowT))
# t-distributed Stochastic Neighbor Embedding.it is a tool to visualize
high-dimensional data. This is T-SNE for BOW
model = TSNE(n components=2, random state=0)
tsne data = model.fit transform(bowT)
tsne data = np.vstack((tsne data.T,SC)).T
tsne df = pd.DataFrame(data = tsne data, columns=("Dim 1", "Dim 2", "la
bel"))
# Ploting the result of tsne
a = sn.FacetGrid(tsne df, hue="label", size=6)
a.map(plt.scatter, 'Dim 1', 'Dim 2').add legend()
plt.title("Bow tsne")
plt.show()
(2000, 390)
<class 'numpy.ndarray'>
```



In Above Bow tsne, positive and negative data points are overlapping, so unable to classify them

```
In [28]: #bi-gram, tri-gram and n-gram

#removing stop words like "not" should be avoided before building n-gra
ms

count_vect = CountVectorizer(ngram_range=(1,2)) #in scikit-learn
final_bigram_counts = count_vect.fit_transform(total_2000['CleanedText'].values)
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 (2000, 66169)
    the number of unique words including both unigrams and bigrams 66169

In [29]: #Dimensionality reduction using truncated SVD
    svdT = TruncatedSVD(n_components=390)
    bowT = svdT.fit_transform(final_bigram_counts)
    # t-distributed Stochastic Neighbor Embedding.it is a tool to visualize
    high-dimensional data. This is t-SNE for bi-gram
    model = TSNE(n_components=2, random_state=0)

    tsne_data = model.fit_transform(bowT)

    tsne_data = np.vstack((tsne_data.T,SC)).T
    tsne_df = pd.DataFrame(data = tsne_data, columns=("Dim 1", "Dim 2", "la")
```

sn.FacetGrid(tsne df, hue="label", size=6).map(plt.scatter, 'Dim 1', 'D

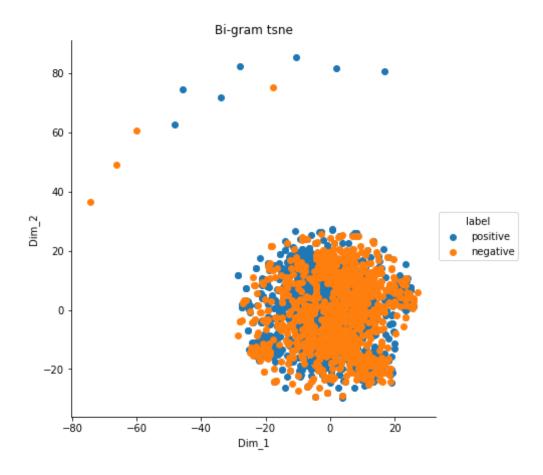
bel"))

plt.show()

Ploting the result of tsne

plt.title('Bi-gram tsne')

im 2').add legend()

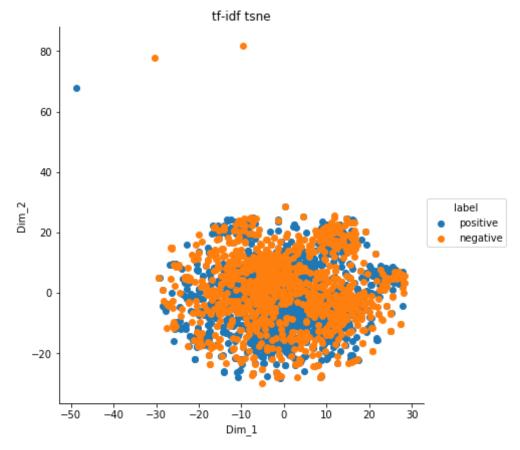


In Above bi -gram tsne, positive and negative data points are overlapping, so unable to classify them

```
In [30]: # tf-idf
    tf_idf_vect = TfidfVectorizer(ngram_range=(1,2))
    final_tf_idf = tf_idf_vect.fit_transform(total_2000['CleanedText'].valu
    es)
    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])
         the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
         the shape of out text TFIDF vectorizer (2000, 66169)
         the number of unique words including both unigrams and bigrams 66169
In [33]: features = tf idf vect.get feature names()
         print("some sample features(unique words in the corpus)", features[100:1
         101)
         some sample features(unique words in the corpus) ['accept', 'accept bac
         k', 'accept chemic', 'accept complain', 'accept even', 'accept howev',
         'accept marley', 'accept way', 'access', 'access car']
In [34]: # source: https://buhrmann.github.io/tfidf-analysis.html
         def top tfidf feats(row, features, top n=25):
             ''' Get top n tfidf values in row and return them with their corres
         ponding feature names.'''
             topn ids = np.argsort(row)[::-1][:top_n]
             top feats = [(features[i], row[i]) for i in topn ids]
             df = pd.DataFrame(top feats)
             df.columns = ['feature', 'tfidf']
             return df
         top tfidf = top tfidf feats(final tf idf[1,:].toarray()[0],features,25)
In [35]: #Dimensionality reduction using truncated SVD
         svdT = TruncatedSVD(n components=390)
         tf idf = svdT.fit transform(final bigram counts)
         model = TSNE(n components=2, random state=0)
         # t-distributed Stochastic Neighbor Embedding.it is a tool to visualize
          high-dimensional data. This is t-SNE for tf-idf
         tsne data = model.fit transform(tf idf)
         tsne data = np.vstack((tsne data.T,SC)).T
         tsne df = pd.DataFrame(np.array(tsne data), columns=("Dim 1", "Dim 2",
```

```
"label"))
# Ploting the result of tsne
sn.FacetGrid(tsne_df, hue="label", size=6).map(plt.scatter, 'Dim_1', 'D
im_2').add_legend()
plt.title("tf-idf tsne")
plt.show()
```



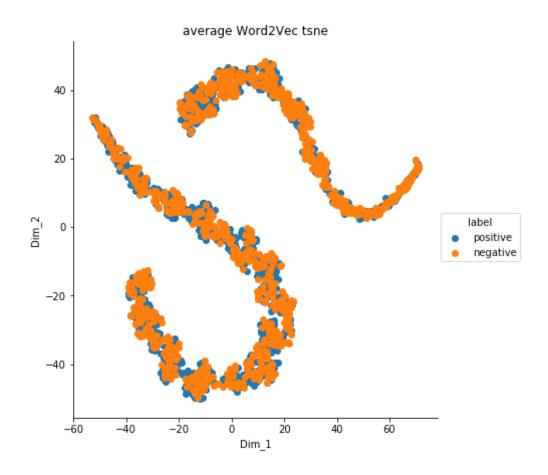
In Above tf-idf tsne, positive and negative data points are overlapping, so unable to classify them

```
In [36]: # Word2Vec
# Train your own Word2Vec model using your own text corpus
i=0
```

```
list of sent=[]
         for sent in total 2000['CleanedText'].values:
             list of sent.append(sent.split())
         print(total 2000['CleanedText'].values[0])
In [37]:
         print("**********
         print(list of sent[0])
         chocol wafer delici hard find gluten free product tast great hit mark s
         tapl pantri
         ************************
         ['chocol', 'wafer', 'delici', 'hard', 'find', 'gluten', 'free', 'produc
         t', 'tast', 'great', 'hit', 'mark', 'stapl', 'pantri']
In [38]: # min count = 5 considers only words that occured atleast 5 times
         w2v model=Word2Vec(list of sent,min count=5,size=50, workers=4)
In [39]: 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 2069
         sample words ['chocol', 'delici', 'hard', 'find', 'gluten', 'free', 'p
         roduct', 'tast', 'great', 'hit', 'mark', 'stapl', 'pantri', 'reciev',
         'yesterday', 'made', 'cold', 'glass', 'tea', 'without', 'sugar', 'oka
         y', 'also', 'think', 'use', 'much', 'powder', 'teaspoon', 'strong', 'cl
         ump', 'like', 'anoth', 'batch', 'tip', 'stir', 'alot', 'fill', 'comple
         t', 'beat', 'sweeten', 'stevia', 'fantast', 'know', 'japanes', 'set',
         'matcha', 'scoop', 'easier', 'control', 'amount']
In [40]: w2v model.wv.most similar('tasti')
Out[40]: [('lot', 0.9998493194580078),
          ('without', 0.9998478293418884),
          ('rather', 0.9998447895050049),
          ('red', 0.9998346567153931),
          ('leav', 0.9998288154602051),
```

```
('thing', 0.9998228549957275),
          ('hair', 0.9998174905776978),
          ('cook', 0.9998151063919067),
          ('regular', 0.9998137950897217),
          ('qo', 0.9998114109039307)]
In [41]: w2v model.wv.most similar('like')
Out[41]: [('tast', 0.9996771812438965),
          ('hot', 0.9996645450592041),
          ('water', 0.9996306896209717),
          ('doesnt', 0.9996281266212463),
          ('cup', 0.9996254444122314),
          ('flavor', 0.9996230006217957),
          ('sweet', 0.9996134638786316),
          ('strong', 0.9995839595794678),
          ('bitter', 0.9995784163475037),
          ('realli', 0.9995777606964111)]
In [42]: # 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 sent): # for each review/sentence
             sent vec = np.zeros(50) # as word vectors are of zero length
             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)
         print(len(sent vectors))
         print(len(sent vectors[0]))
         100%|
                                                      2000/2000 [00:02<00:00, 83
```

```
3.03it/s1
         2000
         50
In [46]: \# S = ["abc def pgr", "def def def abc", "pgr pgr def"]
         model = TfidfVectorizer()
         tf idf matrix = model.fit transform(total 2000['CleanedText'].values)
         # 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 [48]: model = TSNE(n components=2, random state=0)
         tsne data = model.fit transform(sent vectors)
         tsne data = np.vstack((tsne data.T,SC)).T
         tsne df = pd.DataFrame(np.array(tsne data), columns=("Dim_1", "Dim_2",
         "label"))
         # Ploting the result of tsne
         sn.FacetGrid(tsne df, hue="label", size=6).map(plt.scatter, 'Dim 1', 'D
         im 2').add legend()
         plt.title("average Word2Vec tsne")
         plt.show()
```

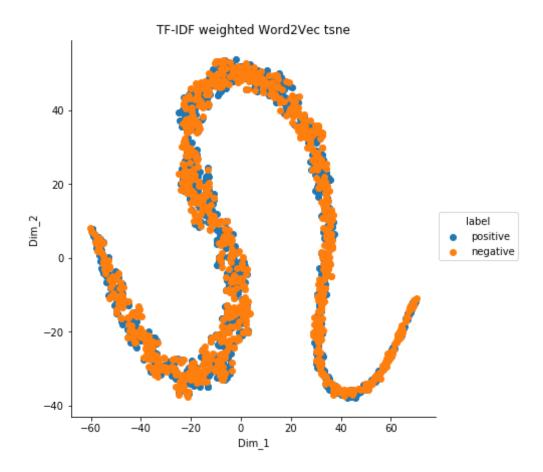


In Above average Word2Vec tsne, positive and negative data points are overlapping, so unable to classify them

```
In [51]: # TF-IDF weighted Word2Vec
    #tfidf_feats = 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 sent): # 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:
                     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 \overline{!} = 0:
                 sent vec /= weight sum
             tfidf sent vectors.append(sent vec)
             row += 1
         100%|
                                                      2000/2000 [00:03<00:00, 62
         8.06it/sl
In [53]: model = TSNE(n components=2, random state=0)
         tsne data = model.fit transform(tfidf sent vectors)
         tsne data = np.vstack((tsne data.T,SC)).T
         tsne df = pd.DataFrame(np.array(tsne data), columns=("Dim 1", "Dim 2",
         "label"))
         # Ploting the result of tsne
         sn.FacetGrid(tsne df, hue="label", size=6).map(plt.scatter, 'Dim 1', 'D
         im 2').add legend()
         plt.title("TF-IDF weighted Word2Vec tsne")
         plt.show()
```



In Above TF-IDF weighted Word2Vec tsne, positive and negative data points are overlapping, so unable to classify them

Observation :- In above all tsne plots, positive and negative datapoints are overlapping, so unable to classify them.