```
In [65]: # Exercise Apply Naive Bayes on Amazon reviews dataset
         %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
         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 sklearn.decomposition import TruncatedSVD
         from sklearn.preprocessing import StandardScaler
         from collections import Counter
         from sklearn.metrics import accuracy score
         from sklearn import cross validation
```

```
import re
# Tutorial about Python regular expressions: https://pymotw.com/2/re/
import string
from nltk.corpus import stopwords
from nltk.stem import PorterStemmer
from nltk.stem.wordnet import WordNetLemmatizer

from gensim.models import Word2Vec
from gensim.models import KeyedVectors
import pickle

from tqdm import tqdm
import os
```

```
In [66]: # using the SQLite Table to read data.
         con = sqlite3.connect(r'G:\machine learning\NAIVE BAYES\database3.sqlit
         e')
         #filtering only positive and negative reviews i.e.
         # not taking into consideration those reviews with Score=3
         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.</pre>
         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 (525814, 10)

Out[66]:

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfulnes
0	1	B001E4KFG0	A3SGXH7AUHU8GW	delmartian	1	1
1	2	B00813GRG4	A1D87F6ZCVE5NK	dli pa	0	0
2	3	B000LQOCH0	ABXLMWJIXXAIN	Natalia Corres "Natalia Corres"	1	1

Exploratory Data Analysis

[7.1.2] 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 [67]: display= pd.read_sql_query("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND UserId="AR5J8UI46CURR"
    ORDER BY ProductID
    """, con)
    display.head()
```

Out[67]:

	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

	ld	ProductId	UserId	ProfileName	HelpfulnessNumerator	Helpfuln
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 [68]: #Sorting data according to ProductId in ascending order
    sorted_data=filtered_data.sort_values('ProductId', axis=0, ascending=Tr
    ue, inplace=False, kind='quicksort', na_position='last')
```

```
In [69]: #Deduplication of entries
  final=sorted_data.drop_duplicates(subset={"UserId","ProfileName","Time"
    ,"Text"}, keep='first', inplace=False)
  final.shape
```

Out[69]: (364173, 10)

```
In [70]: #Checking to see how much % of data still remains
  (final['Id'].size*1.0)/(filtered_data['Id'].size*1.0)*100
```

Out[70]: 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 [71]: display= pd.read_sql_query("""
    SELECT *
    FROM Reviews
    WHERE Score != 3 AND Id=44737 OR Id=64422
    ORDER BY ProductID
    """, con)
    display.head()
```

Out[71]:

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

	ld	ProductId	Userld	ProfileName	HelpfulnessNumerator	Helpfuln
1	44737	B001EQ55RW	A2V0I904FH7ABY	Ram	3	2

In [72]: final=final[final.HelpfulnessNumerator<=final.HelpfulnessDenominator]</pre>

In [73]: #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[73]: 1 307061 0 57110

Name: Score, dtype: int64

7.2.3 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:-

- 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 [74]: # 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.

| '>

| '><

```
In [75]: stop = set(stopwords.words('english')) #set of stopwords
sno = nltk.stem.SnowballStemmer('english') #initialising the snowball s
temmer

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
print(stop)
print('**********************************
print(sno.stem('tasty'))
```

{'do', 've', 'don', 'when', 'and', 'there', "haven't", 'ours', 'shoul d', "mightn't", 'her', 'where', 'between', 'hasn', 'doesn', 'a', 'must n', 'does', 'through', 'you', 'their', 'is', 'them', 'who', 'yours', 'm', 'am', 'ma', 'that', 'ourselves', "mustn't", 'his', 'or', 'too', 't', 'at', 'hers', 'if', 'aren', 'more', 'few', 'needn', "won't", 'abou t', 'shouldn', "doesn't", 'on', 'for', 'against', 'yourself', 'again', 'mightn', 'weren', 'not', 'how', 'such', 'yourselves', 'out', 'our', 'a s', 'itself', "didn't", 'being', 'no', 'by', 'theirs', "couldn't", 'l l', 'myself', 'themselves', 'didn', 'most', 'i', 'from', 'why', "would n't", 'both', 'then', 'own', 'my', 'this', 'o', 'the', 'nor', 'whom', 'it', "you'd", 'below', 'over', 'an', 'above', 'before', 'himself', 've ry', 'just', "shouldn't", "isn't", 'are', 'until', 'any', 'with', 'wha t', "shan't", "that'll", 'she', 'be', 'each', 'isn', "you've", 'same', 'was', 'will', 'having', 'because', 'were', 'y', "hasn't", "needn't", 'we', "aren't", 're', 'but', 'can', 'under', 'only', 'now', 'had', "i t's", 'some', 'ain', "should've", 'which', 'couldn', 'so', 'he', 'in', 'here', 'after', 'during', 'all', "don't", 'they', 'wouldn', 'up', 'hav e', 'down', 'these', 'those', "wasn't", 'of', 'hadn', "weren't", 'to', 'once', 'd', 'its', 'doing', 'wasn', "you're", 'your', 'been', 'haven', "hadn't", 'into', "she's", 'has', 'shan', 'him', 's', 'further', 'herse lf', 'did', 'off', 'other', "you'll", 'than', 'won', 'while', 'me'} *********** tasti

```
In [76]: #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('finall.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 SQLLite table for future.
    conn = sqlite3.connect('final1.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 [77]:

if os.path.isfile('finall.sqlite'):
    conn = sqlite3.connect('finall.sqlite')
    final = pd.read_sql_query(""" SELECT * FROM Reviews WHERE Score !=
    3 """, conn)
    conn.close()
else:
    print("Please the above cell")
```

randomly generate data and sort in ascending order

```
In [107]: #random_sample = final.sample(n = 6000)
    #random_sample.shape
    #random_sample = random_sample.sort_values('Time', axis=0, ascending=Tr
    ue, inplace=False, kind='quicksort', na_position='last')
    sorted_sample = final.sort_values('Time', axis=0, ascending=True, inpla
    ce=False, kind='quicksort', na_position='last')
    sample_60000 = sorted_sample.iloc[0:100000]
    final.shape
    y = sample_60000['Score']
In [108]: # sort the data in 60:20:20 ratio
    x_train_size = int(len(sample_60000)*.60)
    y_size = int(len(y)*.60)
```

```
# split into Train and Test sets
x train = sample 60000[0:x train size]
x test = sample 60000[x train size:len(sample 60000)]
print("total data",len(sample 60000))
print("x train data",len(x train))
#print("x test data",len(x test))
y train = y[0:y size]
y \text{ test} = y[y \text{ size:len}(y)]
print("total output data",len(y))
print("total y train data",len(y train))
#print("total y test data",len(y test))
x \text{ tr size} = int(len(x \text{ test})*.50)
y tr size = int(len(y test)*.50)
x cv = x test[0:x tr size]
x ts = x test[x tr size:len(x test)]
#print("total data",x tr size)
print("x cv data",len(x cv))
print("x ts data",len(x ts))
y cv = y[0:y tr size]
y ts = y[y tr size:len(y test)]
#print("total data",y tr size)
print("y cv data",len(y cv))
print("y ts data",len(y ts))
total data 100000
x train data 60000
total output data 100000
total y train data 60000
x cv data 20000
x ts data 20000
```

```
y_cv data 20000
y_ts data 20000
```

[7.2.2] Bag of Words (BoW)

```
In [109]: #BoW
          count vect = CountVectorizer(ngram range=(1, 2), max features=2000, min d
          f=10) #in scikit-learn
          x tr final counts = count vect.fit transform(x train['CleanedText'].val
          ues)
          x cv final counts = count vect.transform(x cv['CleanedText'].values)
          x ts final counts = count vect.transform(x ts['CleanedText'].values)
          print("the type of count vectorizer ",type(x tr final counts))
          print("the shape of out text BOW vectorizer ",x tr final counts.get sha
          pe())
          print("the number of unique words ", x tr final counts.get shape()[1])
          print("the type of count vectorizer ",type(x cv final counts))
          print("the shape of out text BOW vectorizer ",x cv final counts.get sha
          pe())
          print("the number of unique words ", x cv final counts.get shape()[1])
          print("the type of count vectorizer ",type(x ts final counts))
          print("the shape of out text BOW vectorizer", x ts final counts.get sha
          pe())
          print("the number of unique words ", x ts final counts.get shape()[1])
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
          the shape of out text BOW vectorizer (60000, 2000)
          the number of unique words 2000
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
          the shape of out text BOW vectorizer (20000, 2000)
          the number of unique words 2000
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
```

```
the shape of out text BOW vectorizer (20000, 2000)
          the number of unique words 2000
In [110]: from sklearn.metrics import fl_score
          from sklearn.naive bayes import MultinomialNB
          alpha values = np.arange(14)
          acc = np.empty(len(alpha values))
          error = np.empty(len(alpha values))
          lst = np.array([1000,500,100,50,10,5,1,.1,.5,.01,.05,.001,.005,0.0001])
          ap = len(lst)
          for i in range(ap):
              mnb = MultinomialNB(alpha = lst[i])
              mnb.fit(x tr final counts, y train)
              # predict the response on the crossvalidation train
              pred = mnb.predict(x cv final counts)
              # evaluate CV accuracy
              acc[i] = f1 score(y cv, pred, average='macro') * float(100)
                  #print('\nCV accuracy for k = %d is %d%%' % (i, acc))
              \#error[i] = 100-acc[i]
            # optimal k = int(min(error))
            # print('\nThe optimal number of neighbors is %d.' % optimal k)
              #Generate plot
          d = acc.max()
          i = np.where(acc == d)
          i = i[0][0]
          best alpha = float(lst[i])
          print("Best alpha is:-",best alpha)
```

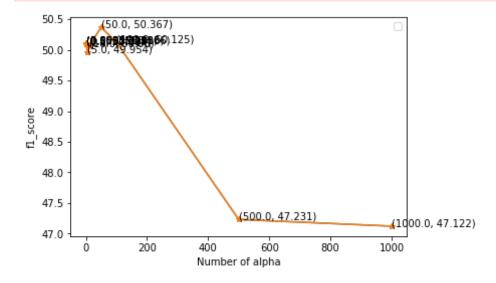
```
# plt.title('k-NN Varying number of neighbors')
plt.plot(lst, acc, marker = '*')

for xy in zip(lst, np.round(acc,3)):
    plt.annotate('(%s, %s)' % xy, xy=xy, textcoords='data')

plt.plot(lst, acc, marker = '*')
#plt.plot(neighbors, train_accuracy, label='Training accuracy')
plt.legend()
plt.xlabel('Number of alpha')
plt.ylabel('fl_score')
plt.show()
```

Best alpha is: - 50.0

No handles with labels found to put in legend.



```
# fitting the model
mnb.fit(x_tr_final_counts, y_train)

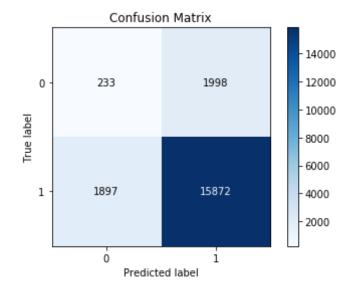
# predict the response
pred = mnb.predict(x_ts_final_counts)

# evaluate accuracy
fl_score_bowT = fl_score(y_ts, pred, average='macro') * float(100)
fl_score_bowT_alpha = best_alpha
print('\nThe fl_score of the Naive Bayes classifier of best_alpha = %f
is %f%' % (best_alpha, fl_score_bowT))
```

The fl_score of the Naive Bayes classifier of best_alpha = 50.000000 is 49.878292%

```
In [112]: import scikitplot.metrics as skplt
skplt.plot_confusion_matrix(y_ts ,pred)
```

Out[112]: <matplotlib.axes._subplots.AxesSubplot at 0x6d90f2add8>



In [113]: #classification report

from sklearn.metrics import classification_report print(classification_report(y_ts, pred))

	precision	recall	T1-score	support
Θ	0.11	0.10	0.11	2231
1	0.89	0.89	0.89	17769
avg / total	0.80	0.81	0.80	20000

Terminology

true positives (TP): We predicted +ve review, and review is also +ve. true negatives (TN): We predicted -ve, and review is also -ve. false positives (FP): We predicted +ve, but the review is not actually +ve.(Also known as a "Type I error.") false negatives (FN): We predicted -ve, but the review is actually +ve.(Also known as a "Type II error.")

False Positive rate --> when it is actually -ve, how often does it predicted +ve = fp/actual-ve = 1998/2231 = .89

```
In [114]: # FPR for bowt bowt_FPR = .89
```

Top 10 features

```
In [115]: # To get all the features name
    features = count_vect.get_feature_names()
    print("some sample features(unique words in the corpus)",features[100:1
    10])

some sample features(unique words in the corpus) ['avail', 'avail amazo
    n', 'averag', 'avoid', 'aw', 'awar', 'away', 'awesom', 'awhil', 'babi']

In [116]: # To count feature for each class while fitting the model
```

```
# Number of samples encountered for each (class, feature) during fittin
          feat count = mnb.feature count
          feat count.shape
Out[116]: (2, 2000)
In [117]: # Number of samples encountered for each class during fitting
          mnb.class count
Out[117]: array([ 6853., 53147.])
In [118]: # Empirical log probability of features given a class(i.e. P(x_i|y))
          log prob = mnb.feature log prob
          log prob
Out[118]: array([[-7.45222902, -8.62574262, -7.48897856, ..., -8.69473549,
                  -8.1982986 , -7.924296631,
                 [-7.04423544, -8.59841724, -7.00470933, \ldots, -8.08182791,
                  -7.30519568, -9.0243210111)
In [119]: feature prob = pd.DataFrame(log prob, columns = features)
          feature prob tr = feature prob.T
          feature prob tr.shape
Out[119]: (2000, 2)
In [120]: # To show top 10 feature from both class
          # Feature Importance
          print("Top 10 Negative Features:-\n", feature prob tr[0].sort values(asc
          ending = False)[0:10])
          print("\n\n Top 10 Positive Features:-\n", feature prob tr[1].sort value
          s(ascending = False)[0:10])
          Top 10 Negative Features:-
                     -4.430046
           tast
```

```
like
         -4.527517
product -4.624184
         -4.920334
one
flavor
         -5.029801
would
         -5.092056
tri
         -5.104296
         -5.204743
good
         -5.378603
tea
buy
         -5.385758
Name: 0, dtype: float64
 Top 10 Positive Features:-
 like
           -4.443833
tast
         -4.465797
tea
         -4.584119
         -4.615069
good
flavor
         -4.642516
         -4.647451
great
love
         -4.729386
        -4.771474
use
         -4.806507
one
product -4.841258
Name: 1, dtype: float64
```

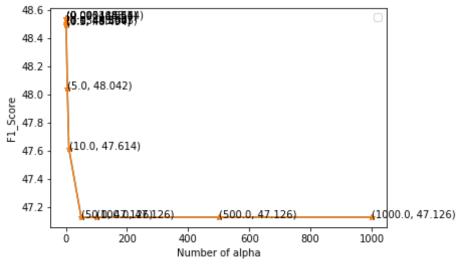
[7.2.5] TF-IDF

```
print("the number of unique words including both unigrams and bigrams "
          , x tr final counts.get shape()[1])
          print("the type of count vectorizer ",type(x_cv_final_counts))
          print("the shape of out text TFIDF vectorizer ",x cv final counts.get s
          hape())
          print("the number of unique words including both unigrams and bigrams "
          , x cv final counts.get shape()[1])
          print("the type of count vectorizer ",type(x ts final counts))
          print("the shape of out text TFIDF vectorizer ",x ts final counts.get s
          hape())
          print("the number of unique words including both unigrams and bigrams "
          , x ts final counts.get shape()[1])
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
          the shape of out text TFIDF vectorizer (60000, 2000)
          the number of unique words including both unigrams and bigrams 2000
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
          the shape of out text TFIDF vectorizer (20000, 2000)
          the number of unique words including both unigrams and bigrams 2000
          the type of count vectorizer <class 'scipy.sparse.csr.csr matrix'>
          the shape of out text TFIDF vectorizer (20000, 2000)
          the number of unique words including both unigrams and bigrams 2000
In [122]: from sklearn.metrics import f1 score
          from sklearn.naive bayes import MultinomialNB
          alpha values = np.arange(14)
          acc = np.empty(len(alpha values))
          error = np.empty(len(alpha values))
          lst = np.array([1000,500,100,50,10,5,1,.1,.5,.01,.05,.001,.005,0.0001])
          ap = len(lst)
          for i in range(ap):
              mnb = MultinomialNB(alpha = lst[i])
```

```
mnb.fit(x tr final counts, y train)
    # predict the response on the crossvalidation train
    pred = mnb.predict(x cv final counts)
    # evaluate CV accuracy
    acc[i] = f1_score(y_cv, pred, average='macro') * float(100)
        \#print('\nCV \ accuracy \ for \ k = %d \ is \ %d%' \ % \ (i, acc))
    #error[i] = 100-acc[i]
 # optimal k = int(min(error))
 # print('\nThe optimal number of neighbors is %d.' % optimal k)
    #Generate plot
d = acc.max()
i = np.where(acc == d)
i = i[0][0]
best alpha = float(lst[i])
print("Best alpha is:-",best alpha)
 # plt.title('k-NN Varying number of neighbors')
plt.plot(lst, acc, marker = '*')
for xy in zip(lst, np.round(acc,3)):
    plt.annotate('(%s, %s)' % xy, xy=xy, textcoords='data')
plt.plot(lst, acc, marker = '*')
#plt.plot(neighbors, train accuracy, label='Training accuracy')
plt.legend()
plt.xlabel('Number of alpha')
plt.ylabel('F1 Score')
plt.show()
```

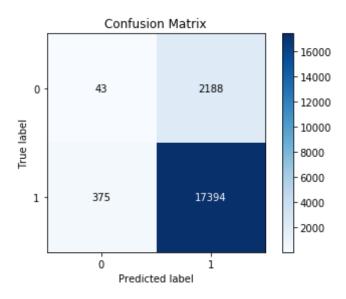
Best alpha is: - 0.01

No handles with labels found to put in legend.



```
In [123]:
                                ======= Naive Bayes = best alpha =========
          mnb = MultinomialNB(alpha=best_alpha)
          # fitting the model
          mnb.fit(x_tr_final_counts, y_train)
          # predict the response
          pred = mnb.predict(x_ts_final_counts)
          # evaluate f1 score
          f1 score tfidf = f1 score(y ts, pred, average='macro') * float(100)
          f1 score tfidf alpha = best alpha
          print('\nThe f1 score of the Naive Bayes classifier of best alpha = %f
           is %f%%' % (best alpha, f1 score tfidf))
          The f1 score of the Naive Bayes classifier of best alpha = 0.010000 is
          48.192288%
In [125]: skplt.plot confusion matrix(y ts ,pred)
```

Out[125]: <matplotlib.axes._subplots.AxesSubplot at 0x6e0bfd0208>



In [126]: #classification report
 print(classification_report(y_ts, pred))

support	f1-score	recall	precision	
2231 17769	0.03 0.93	0.02 0.98	0.10 0.89	0 1
20000	0.83	0.87	0.80	avg / total

Terminology

true positives (TP): We predicted +ve review, and review is also +ve. true negatives (TN): We predicted -ve, and review is also -ve. false positives (FP): We predicted +ve, but the review is not actually +ve.(Also known as a "Type I error.") false negatives (FN): We predicted -ve, but the review is actually +ve.(Also known as a "Type II error.")

False Positive rate --> when it is actually -ve, how often does it predicted +ve = fp/actual-ve = 2188/2231 = .98

```
In [127]: #FPR for tfidf
tfidf_FPR = .98
```

Top 10 features

```
In [128]: # To get all the features name
          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) ['avail', 'avail amazo
          n', 'averag', 'avoid', 'aw', 'awar', 'away', 'awesom', 'awhil', 'babi']
In [129]: # To count feature for each class while fitting the model
          # Number of samples encountered for each (class, feature) during fittin
          q
          feat count = mnb.feature count
          feat count.shape
Out[129]: (2, 2000)
In [130]: # Number of samples encountered for each class during fitting
          mnb.class count
Out[130]: array([ 6853., 53147.])
In [131]: # Empirical log probability of features given a class(i.e. P(x i|y))
          log_prob = mnb.feature_log_prob_
          log_prob
```

```
Out[131]: array([[ -7.37946468, -9.81426237, -7.38675871, ..., -10.20811146,
                  -8.45413408, -7.688395141,
                 [-6.88155419, -8.16472337, -6.80145897, \ldots, -7.66013923,
                   -6.97293912, -9.0533472711)
In [132]: feature prob = pd.DataFrame(log prob, columns = features)
          feature prob tr = feature prob.T
          feature prob tr.shape
Out[132]: (2000, 2)
In [133]: # To show top 10 feature from both class
          # Feature Importance
          print("Top 10 Negative Features:-\n", feature prob tr[0].sort values(asc
          ending = False)[0:10])
          print("\n\n Top 10 Positive Features:-\n", feature prob tr[1].sort value
          s(ascending = False)[0:10])
          Top 10 Negative Features:-
           tast
                    -4.680602
          product -4.774609
          like
                   -4.801611
          would
                   -5.123808
                   -5.153619
          one
          flavor
                  -5.215313
                   -5.276224
          order
          tri
                  -5.304676
                  -5.308975
          buy
          box
                   -5.419335
          Name: 0, dtype: float64
           Top 10 Positive Features:-
           tea
                    -4.941942
                   -4.953177
          great
          tast
                   -5.029164
          love
                   -5.029358
                   -5.035356
          good
          like
                   -5.061389
```

flavor -5.143643 product -5.182316 use -5.241207 one -5.324948 Name: 1, dtype: float64

```
In [134]: from prettytable import PrettyTable
x = PrettyTable()
x.field_names = ["Vectorizer", "hyperparameter(alpha)", "F1_Score", "FPR"]
x.add_row(["BOW",f1_score_bowT_alpha,f1_score_bowT,bowt_FPR])
x.add_row(["TF-IDF",f1_score_tfidf_alpha,f1_score_tfidf,tfidf_FPR])
print(x)
```

Vectorizer	+ hyperparameter(alpha) +	F1_Score	FPR
BOW TF-IDF	50.0	49.87829207532883 48.19228835440566	0.89

as per the above table, we will consider bow vectorizer for classification of positive and negative points because FPR of BOW(.89) is less than TF-IDF(.98) and alpha of BOW(50) is greater than TFIDF(.01) and it is well fit. it is neither underfit nor overfit.