Entendendo a base de dados.

O conjunto de dados fornecido contém dados de tweets nas alças de twitter de várias companhias aéreas.

Ele contém um total de 12 colunas, das quais uma coluna especifica o sentimento do tweet. Todas as outras colunas fornecem várias informações relacionadas ao que foi o tweet, de onde foi postado, quando foi postado, é retuitado; etc.

Data Description

Description of columns of the dataset is given below -

```
tweet id -- Id of the tweet
```

airline sentiment -- Sentiment of the tweet (Target variable)

airline sentiment confidence -- Confidence with which the given sentiment was determined

negativereason confidence -- Confidence with which the negative reason of tweet was predicted

name -- Name of the person who tweeted

retweet count -- Number of retweets

text -- Text of the tweet whose sentiment has to be predicted

tweet created -- Time at which the tweet was created

tweet location -- Location from where the tweet was posted

user_timezone -- Time zone from where the tweet was posted

negativereason -- Reason for which user posted a negative tweet

airline -- Airline for which the tweet was posted

In [1]:

```
import numpy as np
import pandas as pd
import re
import nltk
import matplotlib.pyplot as plt
%matplotlib inline
```

In [2]:

```
data_source_url = 'Tweets.csv'
airline_tweets = pd.read_csv(data_source_url)
```

Precisamos Fazer um pre-processamento desse texto, para isso vamos tirar pontos, virgulas, underlines e etc.

In [3]:

```
features = airline_tweets.iloc[:, 10].values#Pegando os textos
labels = airline_tweets.iloc[:, 1].values#classes
```

In [4]:

```
processed_features = []

for sentence in range(0, len(features)):
    # Remove todos os caracteres especiais
    processed_feature = re.sub(r'\W', ' ', str(features[sentence]))

# remova todos os caracteres únicos
    processed_feature= re.sub(r'\s+[a-zA-Z]\s+', ' ', processed_feature)

# Remova caracteres únicos desde o início
    processed_feature = re.sub(r'\^[a-zA-Z]\s+', ' ', processed_feature)

# substitui multiplos espaços
    processed_feature = re.sub(r'\s+', ' ', processed_feature, flags=re.I)

# Removing prefixed 'b'
    processed_feature = re.sub(r'\b\s+', '', processed_feature)

# Converting to caixa baixa
    processed_feature = processed_feature.lower()

processed_features.append(processed_feature)
```

In [5]:

```
#Doc1 = "I like to play football"
#Doc2 = "It is a good game"
#Doc3 = "I prefer football over rugby"

#feature vector

#Vocab = [I, like, to, play, football, it, is, a, good, game, prefer, over, rugb y]
#[1, 1, 1, 1, 0, 0, 0, 0, 0, 0, 0]
```

In [6]:

```
#nltk.download("stopwords")
```

In [7]:

```
from nltk.corpus import stopwords

from sklearn.feature_extraction.text import TfidfVectorizer

vectorizer = TfidfVectorizer(max_features=2500, min_df=7, max_df=0.8, stop_words = stopwords.words('english'))

processed_features = vectorizer.fit_transform(processed_features).toarray()
```

max_dffloat or int, default=1.0

When building the vocabulary ignore terms that have a document frequency strictly higher than the given threshold (corpus-specific stop words). If float in range [0.0, 1.0], the parameter represents a proportion of documents, integer absolute counts. This parameter is ignored if vocabulary is not None.

min dffloat or int, default=1

When building the vocabulary ignore terms that have a document frequency strictly lower than the given threshold. This value is also called cut-off in the literature. If float in range of [0.0, 1.0], the parameter represents a proportion of documents, integer absolute counts. This parameter is ignored if vocabulary is not None.

max features int, default=None

If not None, build a vocabulary that only consider the top max_features ordered by term frequency across the corpus. This parameter is ignored if vocabulary is not None.

In [8]:

```
# separando o modelo em treino e text
from sklearn.metrics import accuracy_score
from sklearn.metrics import confusion_matrix
from sklearn.metrics import accuracy_score
from sklearn.metrics import classification_report
```

In [9]:

```
from sklearn.model_selection import train_test_split

# Nessa situação, vamos separara train, test and validation.
X_train, X_test, y_train, y_test = train_test_split(processed_features, labels, test_size=0.2, random_state=0)
```

In [10]:

```
from sklearn.linear_model import LogisticRegression
from sklearn.neighbors import KNeighborsClassifier
from sklearn.svm import SVC, LinearSVC, NuSVC
from sklearn.tree import DecisionTreeClassifier
from sklearn.ensemble import RandomForestClassifier, AdaBoostClassifier, Gradien
tBoostingClassifier
from sklearn.naive_bayes import GaussianNB
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
from sklearn.discriminant_analysis import QuadraticDiscriminantAnalysis
from sklearn.metrics import accuracy_score
```

In [11]:

```
from sklearn.metrics import confusion_matrix
from sklearn.metrics import accuracy_score
from sklearn.metrics import classification_report
```

In [12]:

```
def plot confusion matrix(cm, classes,
                          normalize=False,
                          title='Confusion matrix',
                          cmap=plt.cm.Blues):
    plt.imshow(cm, interpolation='nearest', cmap=cmap)
    plt.title(title)
    plt.colorbar()
    tick marks = np.arange(len(classes))
    plt.xticks(tick_marks, classes, rotation=45)
    plt.yticks(tick marks, classes)
    if normalize:
        cm = cm.astype('float') / cm.sum(axis=1)[:, np.newaxis]
    thresh = cm.max() / 2.
    for i, j in itertools.product(range(cm.shape[0]), range(cm.shape[1])):
        plt.text(j, i, cm[i, j],
                 horizontalalignment="center",
                 color="white" if cm[i, j] > thresh else "black")
    plt.tight_layout()
    plt.ylabel('True label')
    plt.xlabel('Predicted label')
```

In [13]:

```
## k-means
from sklearn.cluster import KMeans
kmeans = KMeans(n_clusters=3, random_state=0).fit(X_train)
```

In [14]:

```
print( doc )
from time import time
import numpy as np
import matplotlib.pyplot as plt
from sklearn import metrics
from sklearn.cluster import KMeans
from sklearn.datasets import load digits
from sklearn.decomposition import PCA
from sklearn.preprocessing import scale
np.random.seed(42)
data = scale(X train)
n samples, n features = data.shape
n digits = len(np.unique(y train))
labels = y train
sample size = 300
print("n digits: %d, \t n samples %d, \t n features %d"
      % (n digits, n samples, n features))
print(82 * ' ')
print('init\t\ttime\tinertia\thomo\tcompl\tv-meas\tARI\tAMI\tsilhouette')
def bench k means(estimator, name, data):
    t0 = time()
    estimator.fit(data)
    print('%-9s\t%.2fs\t%i\t%.3f\t%.3f\t%.3f\t%.3f\t%.3f\t%.3f\
          % (name, (time() - t0), estimator.inertia ,
             metrics.homogeneity score(labels, estimator.labels ),
             metrics.completeness score(labels, estimator.labels ),
             metrics.v measure score(labels, estimator.labels ),
             metrics.adjusted_rand_score(labels, estimator.labels_),
             metrics.adjusted mutual info score(labels, estimator.labels),
             metrics.silhouette score(data, estimator.labels ,
                                      metric='euclidean',
                                      sample size=sample size)))
bench_k_means(KMeans(init='k-means++', n_clusters=3, n init=10),
              name="k-means++", data=data)
bench k means(KMeans(init='random', n clusters=3, n init=10),
              name="random", data=data)
pca = PCA(n components=n digits).fit(data)
bench_k_means(KMeans(init=pca.components_, n_clusters=3, n_init=1),
              name="PCA-based",
              data=data)
```

init silhouette		time	inertia homo	compl	v-meas	ARI	AMI
k-means++		8.93s	26904994	0.008	0.016	0.011	-0.0
19	0.011	0.002		0.000	0.020	0.011	
random		10.02s	26910058	0.067	0.107	0.083	-0.0
57	0.082	-0.025	222222				
PCA-based		2.93s	26901659	0.107	0.091	0.099	0.07
4	0.099	-0.004					

In []: