```
!pip install fuzzywuzzy

Requirement already satisfied: fuzzywuzzy in
/usr/local/lib/python3.10/dist-packages (0.18.0)
```

Imports

```
import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
import nltk
import re
import resource
from nltk.stem import PorterStemmer
from bs4 import BeautifulSoup
from fuzzywuzzy import fuzz
nltk.download('stopwords')
nltk.download('punkt')
from nltk.corpus import stopwords
from nltk import word tokenize, ngrams
import warnings
color = sns.color palette()
stopwords = set(stopwords.words('english'))
%matplotlib inline
warnings.filterwarnings('ignore')
pd.options.mode.chained assignment = None
[nltk data] Downloading package stopwords to /root/nltk data...
[nltk data]
              Package stopwords is already up-to-date!
[nltk data] Downloading package punkt to /root/nltk data...
[nltk data]
              Package punkt is already up-to-date!
```

Data reading

```
df = pd.read_csv('/content/quora_duplicate_questions.tsv', sep='\t')
```

Data Understanding

```
print("Number of data points:",df.shape[0])
print("Number of features:",df.shape[1])

Number of data points: 404290
Number of features: 6

df.head()
{"type":"dataframe","variable_name":"df"}
```

```
df.tail()
{"summary":"{\n \"name\": \"df\",\n \"rows\": 5,\n \"fields\": [\n
{\n \"column\": \"id\",\n \"properties\": {\n
\"dtype\": \"number\",\n \"std\": 1,\n \"min\": 404285,\
n \"max\": 404289,\n \"num_unique_values\": 5,\n
\"std\": 225059,\n \"min\": 18840,\n \"max\": 537932,\n
\"num_unique_values\": 5,\n \"samples\": [\n 18840
537932,\n 537928\n ],\n \"semantic_type\":
\"\",\n \"description\": \"\"\n }\n },\n {\n
                                             18840,\n
\"column\": \"qid2\",\n \"properties\": {\n \"dtype\":
[\n \"Do you believe there is life after death?\",\n
\"What is like to have sex with cousin?\",\n \"What is one
\"Is it true that there is life after death?\",\n
\"What is it like to have sex with your cousin?\",\n
this coin?\"\n ],\n \"semantic type\": \"\",\n
\"\",\n \"description\": \"\"n }\n }\n ]\
n}","type":"dataframe"}
```

##EDA

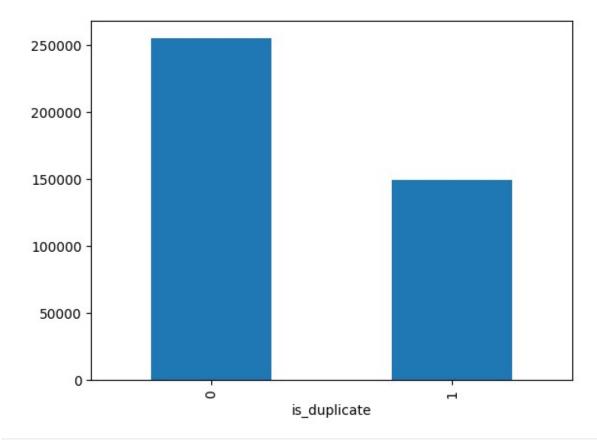
```
len(df)
404290
```

Dataset with question pairs of 404287 size

Target Variable Distribution

```
df['is_duplicate'].value_counts()
```

```
is_duplicate
0    255027
1    149263
Name: count, dtype: int64
df['is_duplicate'].value_counts().plot(kind='bar')
<Axes: xlabel='is_duplicate'>
```



```
len(df[df['is_duplicate'] == 0])/len(df) * 100
63.08021469737069
len(df[df['is_duplicate'] == 1])/len(df) * 100
36.9197853026293
```

We have 63% of non-similar question pairs and 37% of similar question pairs from the calculations above

This indicates that possibly our dataset is imbalanced.

Unique questions details

```
print(len(df[df.isnull().any(axis=1)]))
df = df.fillna('')
print(len(df[df.isnull().any(axis=1)]))
3
0
questions combined = np.concatenate([df['question1'],
df['question2']])
full questions df = pd.DataFrame(questions combined,
columns=['questions'])
full questions df['questions'].nunique()
537361
question wise value counts =
full questions df['questions'].value counts()
print(np.sum(question wise value counts > 1))
print(round(np.sum(question wise value counts >
1)/len(set(questions combined)) * 100, 2))
111873
20.82
print(np.sum(question wise value counts == 1))
print(round(np.sum(question wise value counts ==
1)/len(set(questions combined)) * 100, 2))
425488
79.18
```

So there are total 537361 unique questions in our dataset.

And there are 111872 questions repeated more than once which is 20.82% of unique questions.

And 425488 questions appeared only once means 79.18% of total unique questions

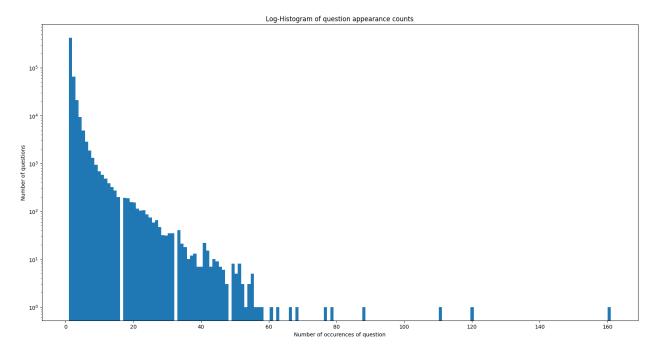
```
print(question_wise_value_counts.index[0])
question_wise_value_counts[0]
What are the best ways to lose weight?
161
```

Most common question: What are the best ways to lose weight?

It appeared 161 times

Each question occurences count

```
plt.figure(figsize=(20, 10))
plt.hist(question_wise_value_counts, bins=170)
plt.yscale('log')
plt.title('Log-Histogram of question appearance counts')
plt.xlabel('Number of occurences of question')
plt.ylabel('Number of questions')
Text(0, 0.5, 'Number of questions')
```



From the above histogram graph we can see that most of the questions repeated maxium of 50 times. Other questions those appeared more than 60 times are very less. As we know the most repeated question appeared to be 161 times.

Word distribution of questions

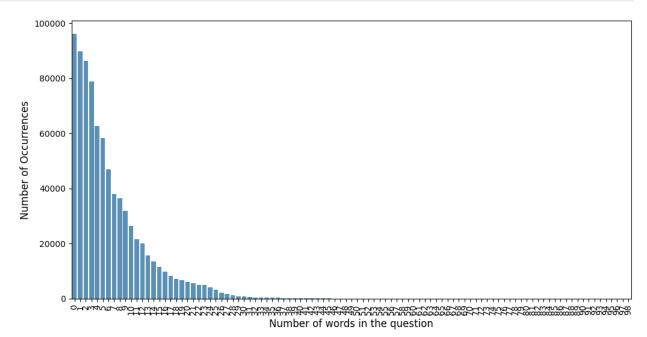
How words are distributed in question pairs. This feature allows us to understand word level distribution of the questions.

```
full_questions_df['number_of_words'] =
full_questions_df["questions"].str.split().apply(len)

cnt_srs = full_questions_df['number_of_words'].value_counts()

plt.figure(figsize=(12,6))
sns.barplot(cnt_srs.values, alpha=0.8, color=color[0])
plt.ylabel('Number of Occurrences', fontsize=12)
plt.xlabel('Number of words in the question', fontsize=12)
```

```
plt.xticks(rotation='vertical')
plt.show()
```



Character distribution

How character legths varies in question pairs. This feature allows us to understand character level distribution of the questions.

```
full_questions_df['number_of_chars'] =
full_questions_df["questions"].apply(len)

cnt_srs = full_questions_df['number_of_chars'].value_counts()

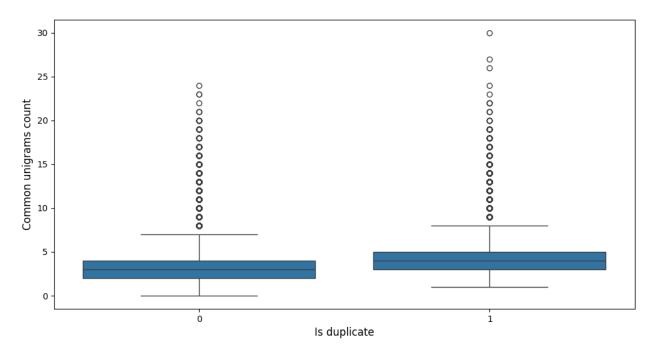
plt.figure(figsize=(50,8))
sns.barplot(cnt_srs.values, alpha=0.8, color=color[3])
plt.ylabel('Number of Occurrences', fontsize=12)
plt.xlabel('Number of characters in the question', fontsize=12)
plt.xticks(rotation='vertical')
plt.show()
```



Unigrams

By understanding the unigrams present in our question pairs we can understand common unigrams present in our questions.

```
def unigrams(question):
    return [word for word in word tokenize(question.lower()) if word
not in stopwords]
def common unigrams(row data):
    return len(
set(row data["unigrams ques1"]).intersection(set(row data["unigrams qu
es2"])) )
def common unigram ratio(row data):
    return float(row data["unigrams common count"]) / max(len(
set(row_data["unigrams_ques1"]).union(set(row_data["unigrams_ques2"]))
),1)
df["unigrams ques1"] = df['question1'].apply(lambda x:
unigrams(str(x)))
df["unigrams_ques2"] = df['question2'].apply(lambda x:
unigrams(str(x)))
df["unigrams common count"] = df.apply(lambda row:
common unigrams(row),axis=1)
df["unigrams common ratio"] = df.apply(lambda row:
common unigram ratio(row), axis=1)
plt.figure(figsize=(12,6))
sns.boxplot(x="is duplicate", y="unigrams common count", data=df)
plt.xlabel('Is duplicate', fontsize=12)
plt.ylabel('Common unigrams count', fontsize=12)
plt.show()
```



We see there is some count of common unigrams present in our question and we have some significant change in count among two target classes which will be useful for the modelling.

Data Preprocessing

Missing values

```
#Checking whether there are any rows with null values
null_rows = df[df.isnull().any(axis=1)]
null_rows

{"repr_error":"Out of range float values are not JSON compliant:
nan", "type":"dataframe", "variable_name":"null_rows"}

# Rows with null values are removed
print(df.shape)
df = df[~df.isnull().any(axis=1)]
print(df.shape)

(404290, 10)
(404290, 10)
```

Data Cleaning

Text data cleaning needs to be done to make sure we deal with proper data. Different NLP stratergies listed below are followed

```
def clean_sentence(text):
   if pd.isnull(text):
      return ''
```

```
if type(text) != str or text=='':
         return ''
    ps = PorterStemmer()
    text = re.sub("\'s", " ", text)
    text = re.sub("i'm", "i am", text, flags=re.IGNORECASE)
    text = text.replace("can't", "can not")
    text = re.sub("\'re", " are ", text)
    text = text.replace(" whats ", " what is ")
    text = re.sub("\'ve", " have ", text)
text = re.sub("n't", " not ", text)
text = text.replace("b\.g\.", " bg ")
    text = re.sub("\'d", " would ", text)
    text = text.replace("\'ll", " will ")
    text = re.sub("i'm", "i am", text, flags=re.IGNORECASE)
text = text.replace("e\.g\.", " eg ")
    text = re.sub("(the[\s]+|The[\s]+)?U\.S\.A\.", "America ", text,
flags=re.IGNORECASE)
    text = re.sub("(\d+)(kK)", " \g<1>000 ", text)
    text = re.sub("\(s\)", " ", text, flags=re.IGNORECASE)
    text = re.sub("[c-fC-F]\:\/", " disk ", text)
    text = re.sub('(? <= [0-9]) \setminus, (? = [0-9])', "", text)
    text = text.replace('\$', " dollar ")
text = text.replace('\%', " percent ")
    text = text.replace('\&', " and ")
    text = ' '.join([word for word in text.split(" ") if word not in
stopwords]).lower()
    #text = ' '.join([ps.stem(word) for word in text])
df["question1"] = df["question1"].fillna("").apply(clean sentence)
df["question2"] = df["question2"].fillna("").apply(clean sentence)
```

Feature Engineering

Basic Textual Features

```
def get_unique_words(sentence):
    sent_split = sentence.split(" ")
    sent_split_processed = [word.lower().strip() for word in sent_split]
    return set(sent_split_processed)

def word_common_details(row):
    words1 = get_unique_words(row['question1'])
    words2 = get_unique_words(row['question2'])
    return 1.0 * len(words1 & words2)
```

```
def word count details(row):
    words1 = get unique words(row['question1'])
    words2 = get_unique words(row['question2'])
    return 1.0 * (len(words1) + len(words2))
def word share data(row):
    words1 = get unique words(row['question1'])
    words2 = get unique words(row['question2'])
    return 1.0 * len(words1 & words2)/(len(words1) + len(words2))
df['q1len'] = df['question1'].str.len()
df['q2len'] = df['question2'].str.len()
df['diff len'] = df.q1len - df.q2len
df['freq qid1'] = df.groupby('qid1')['qid1'].transform('count')
df['freq gid2'] = df.groupby('gid2')['gid2'].transform('count')
df['freq_q1+q2'] = df['freq_qid1']+df['freq_qid2']
df['freq_q1-q2'] = abs(df['freq_qid1']-df['freq_qid2'])
df['q1 n words'] = df['question1'].apply(lambda row: len(row.split('
')))
df['q2 n words'] = df['question2'].apply(lambda row: len(row.split('
')))
df['len char q1'] = df.question1.apply(lambda x:
len(''.join(set(str(x)))))
df['len char q2'] = df.question2.apply(lambda x:
len(''.join(set(str(x)))))
df['len word q1'] = df.question1.apply(lambda x: len(str(x).split()))
df['len word q2'] = df.question2.apply(lambda x: len(str(x).split()))
df['word Common'] = df.apply(word common details, axis=1)
df['word Total'] = df.apply(word count details, axis=1)
df['word share'] = df.apply(word share data, axis=1)
```

Similarity Features

To get semantic features we used 'word2vec-google-news-300' model to get the vector representation of the questions first then find the similarity score using the different distance measures

```
import gensim.downloader as api
from scipy.spatial.distance import cosine, cityblock, jaccard,
canberra, euclidean, minkowski, braycurtis

model = api.load('word2vec-google-news-300')
```

```
def get sent vector repr(sent):
    tokens = str(sent).lower()
    tokens = word tokenize(tokens)
    tokens = [w for w in tokens if w.isalpha()]
    sentence vector = []
    for w in tokens:
        try:
            sentence vector.append(model[w])
        except:
            continue
    sentence vector = np.array(sentence vector)
    vector = sentence vector.sum(axis=0)
    return vector / np.sqrt((vector ** 2).sum())
from tgdm.notebook import tgdm
vectors question1 = np.zeros((df.shape[0], 300))
for index, question in enumerate(tqdm(df['question1'].values)):
    vectors question1[index, :] = get sent vector repr(question)
vectors question2 = np.zeros((df.shape[0], 300))
for index, question in enumerate(tqdm(df['question2'].values)):
    vectors question2[index, :] = get sent vector repr(question)
{"model_id": "c2d0a9bb021d4120a1d2f9975f479387", "version major": 2, "vers
ion minor":0}
{"model id": "3d9eaf22191846e49a4e95c25b5c432d", "version major": 2, "vers
ion minor":0}
df['cosine distance'] = [cosine(x, y) for (x, y) in
zip(np.nan to num(vectors question1),
np.nan to num(vectors question2))]
df['cityblock\ distance'] = [cityblock(x, y)\ for\ (x, y)\ in
zip(np.nan to num(vectors question1),
np.nan to num(vectors question2))]
df['jaccard distance'] = [jaccard(x, y) for (x, y) in
zip(np.nan to num(vectors question1),
np.nan to num(vectors question2))]
df['canberra distance'] = [canberra(x, y) for (x, y) in
zip(np.nan to num(vectors question1),
np.nan to num(vectors guestion2))]
df['euclidean\ distance'] = [euclidean(x, y)\ for\ (x, y)\ in
zip(np.nan_to_num(vectors_question1),
np.nan to num(vectors_question2))]
df['minkowski distance'] = [minkowski(x, y, 3) for (x, y) in
zip(np.nan to num(vectors question1),
np.nan to num(vectors question2))]
df['braycurtis distance'] = [braycurtis(x, y) for (x, y) in
```

```
zip(np.nan_to_num(vectors_question1),
np.nan_to_num(vectors_question2))]
```

Substring level features

```
df['fuzz gratio'] = df.apply(lambda x:
fuzz.QRatio(str(x['question1']), str(x['question2'])), axis=1)
df['fuzz partial ratio'] = df.apply(lambda x:
fuzz.partial ratio(str(x['question1']), str(x['question2'])), axis=1)
# df['fuzz WRatio'] = df.apply(lambda x:
fuzz.WRatio(str(x['question1']), str(x['question2'])), axis=1)
# df['fuzz partial token set ratio'] = df.apply(lambda x:
fuzz.partial token set ratio(str(x['question1']),
str(x['question2'])), axis=1)
# df['fuzz partial token sort ratio'] = df.apply(lambda x:
fuzz.partial token sort ratio(str(x['question1']),
str(x['question2'])), axis=1)
# df['fuzz token set ratio'] = df.apply(lambda x:
fuzz.token set ratio(str(x['question1']), str(x['question2'])),
axis=1)
# df['fuzz token sort ratio'] = df.apply(lambda x:
fuzz.token sort ratio(str(x['question1']), str(x['question2'])),
axis=1)
df.head()
{"type": "dataframe", "variable name": "df"}
df.columns
Index(['id', 'qid1', 'qid2', 'question1', 'question2', 'is duplicate',
        'unigrams_ques1', 'unigrams_ques2', 'unigrams_common_count',
        'unigrams common ratio', 'qlen', 'q2len', 'diff len',
'freq qid1',
        'freq qid2', 'freq q1+q2', 'freq q1-q2', 'q1 n words',
'a2 n words',
       'len_char_q1', 'len_char_q2', 'len_word_q1', 'len_word_q2', 'word_Common', 'word_Total', 'word_share', 'cosine_distance', 'cityblock_distance', 'jaccard_distance', 'canberra_distance',
        'euclidean_distance', 'minkowski_distance',
'braycurtis distance',
        'fuzz gratio'],
      dtype='object')
```

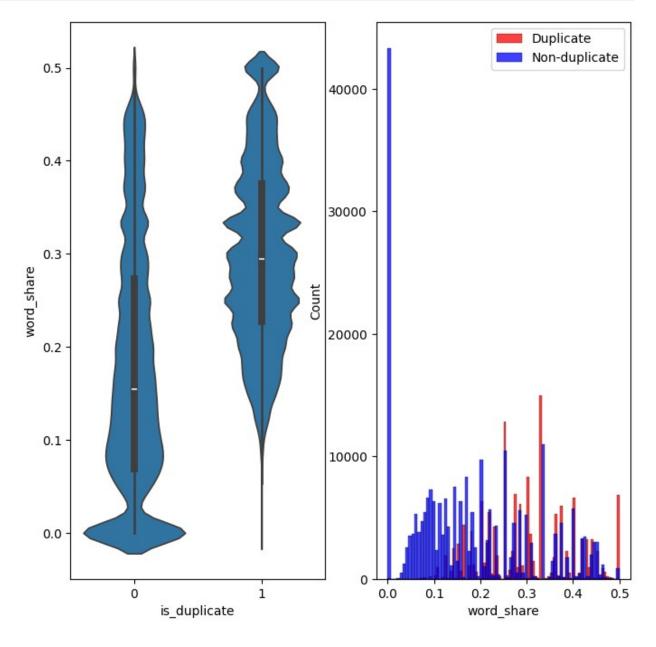
Plots understand features extracted

shared words plot and distribution

```
plt.figure(figsize=(8, 8))
```

```
plt.subplot(1,2,1)
sns.violinplot(x = 'is_duplicate', y = 'word_share', data = df[0:])

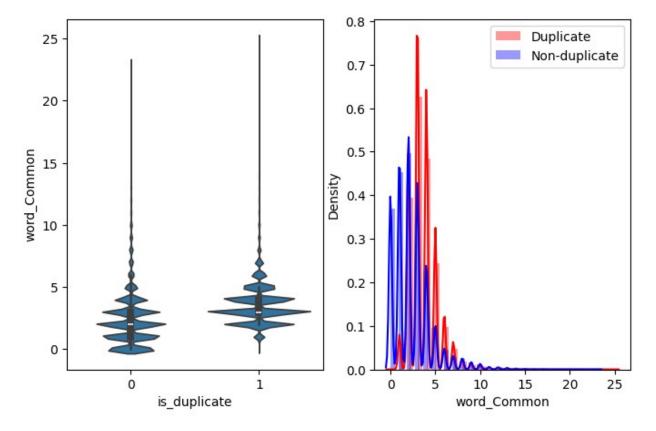
plt.subplot(1,2,2)
sns.histplot(df[df['is_duplicate'] == 1.0]['word_share'][0:] , label =
"Duplicate", color = 'red')
sns.histplot(df[df['is_duplicate'] == 0.0]['word_share'][0:] , label =
"Non-duplicate" , color = 'blue' )
plt.legend()
plt.show();
```



common words plot & distribution

```
plt.figure(figsize=(8,5))
plt.subplot(1,2,1)
sns.violinplot(x = 'is_duplicate', y = 'word_Common', data = df[0:])

plt.subplot(1,2,2)
sns.distplot(df[df['is_duplicate'] == 1.0]['word_Common'][0:] , label
= "Duplicate", color = 'red')
sns.distplot(df[df['is_duplicate'] == 0.0]['word_Common'][0:] , label
= "Non-duplicate" , color = 'blue' )
plt.legend()
plt.show();
```



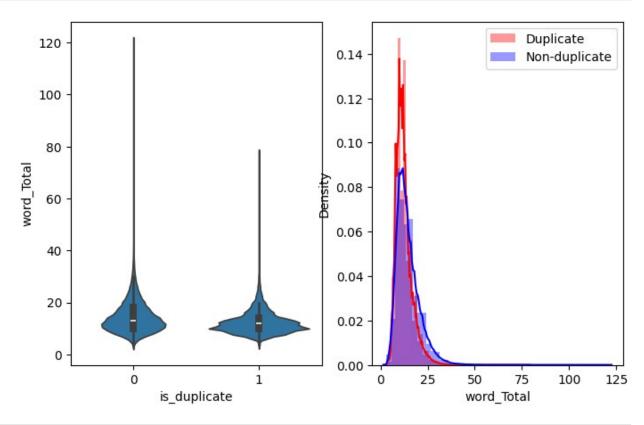
total words plots and distribution

```
plt.figure(figsize=(8, 5))

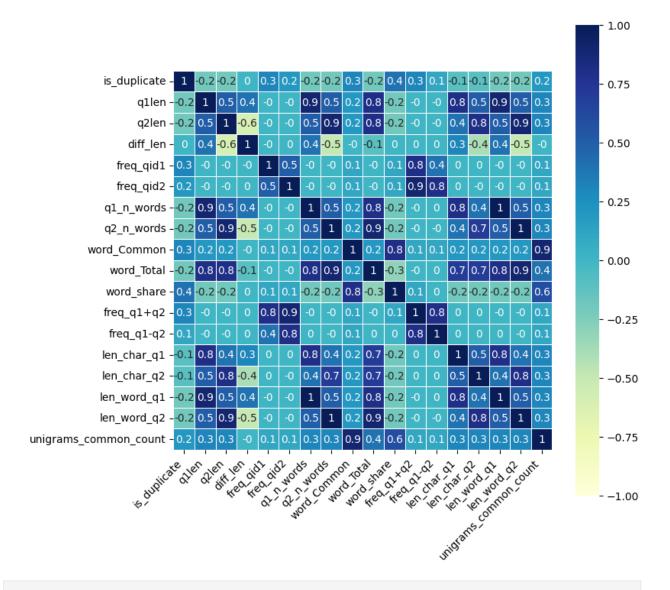
plt.subplot(1,2,1)
sns.violinplot(x = 'is_duplicate', y = 'word_Total', data = df[0:])

plt.subplot(1,2,2)
sns.distplot(df[df['is_duplicate'] == 1.0]['word_Total'][0:] , label =
"Duplicate", color = 'red')
sns.distplot(df[df['is_duplicate'] == 0.0]['word_Total'][0:] , label =
```

```
"Non-duplicate" , color = 'blue' )
plt.legend()
plt.show();
```



```
corr columns = ['is duplicate', 'q1len', 'q2len', 'diff len',
'freq qid1',
                  'freq qid2', 'q1 n words', 'q2 n words',
'word_Common', 'word_Total',
'word_share', 'freq_q1+q2', 'freq_q1-q2',
'len_char_q1', 'len_char_q2', 'len_word_q1', 'len_word_q2',
'unigrams_common_count']
correlation data = df[corr columns].corr()
fig, ax = plt.subplots(figsize=(8,8))
ax = sns.heatmap(
    round(correlation data,1),
    vmin=-1, vmax=1, center=0, annot=True, linewidths=.5,
    cmap="YlGnBu",
    square=True
ax.set xticklabels(
    ax.get xticklabels(),
    rotation=45,
    horizontalalignment='right'
);
```



Model Building

```
df_original = df.copy()

# Removing features those are not useful and non-numeric. And also
braycurtis_distance as it has all null values

df.drop(['qid1','qid2','question1','question2', 'unigrams_ques1',
    'unigrams_ques2', 'braycurtis_distance'], axis=1, inplace=True)

df.dropna(axis = 0,inplace = True)

y_true = df['is_duplicate']
y_true = list(map(int, y_true.values))
df.drop(['id','is_duplicate'], axis=1, inplace=True)
```

```
cols = list(df.columns)
for i in cols:
    df[i] = df[i].apply(pd.to_numeric)
```

Test-train split

```
from sklearn.model selection import train test split
X_train,X_test, y_train, y_test = train_test_split(df, y_true,
stratify=y true, test size=0.2)
from collections import Counter
print("-"*10, "Output variable distribution in train data", "*"*10)
train distr = Counter(y train)
train len = len(y train)
print("Class 0: ",round(int(train distr[0])*100/train len,2),"Class 1:
', round(int(train_distr[1])*100/train_len,2))
print("-"*10, "output variable distribution in validation data",
"*"*10)
test distr = Counter(y_test)
test len = len(y_test)
print("Class 1: ",round(int(test distr[0])*100/test len,2), "Class 1:
",round(int(test distr[1])*100/test len,2))
----- Output variable distribution in train data ********
Class 0: 63.08 Class 1: 36.92
----- output variable distribution in validation data *******
Class 1: 63.08 Class 1: 36.92
```

Logistic Regression

SGD is a optimization method SGDClassifier is a linear classifier with SGD.

Logistic Regression == SGDClassifier(class_weight='balanced', alpha=i, penalty='l2', loss='log', random_state=42)

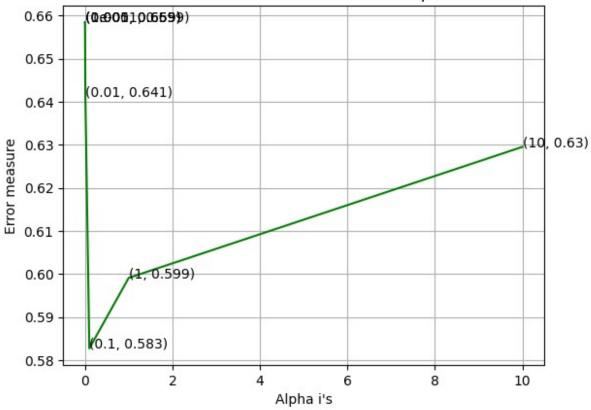
```
from sklearn.linear_model import SGDClassifier
from sklearn.calibration import CalibratedClassifierCV
from sklearn.metrics import confusion_matrix
from sklearn.metrics import accuracy_score, log_loss

alpha_values = [10 ** x for x in range(-5, 2)]

log_error_values=[]
for i in alpha_values:
    clf = SGDClassifier(alpha=i, penalty='l2', loss='log', random_state=42)
    clf.fit(X_train, y_train)
    sig_clf = CalibratedClassifierCV(clf, method="sigmoid")
```

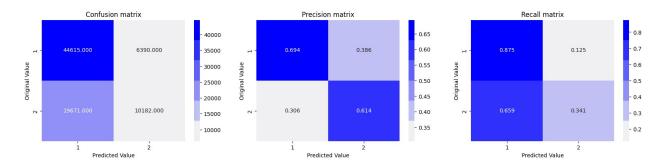
```
sig clf.fit(X train, y train)
    predict y = sig clf.predict proba(X test)
    log error values.append(log_loss(y_test, predict_y,
labels=clf.classes , eps=1e-15))
    print('For values of alpha = ', i, "The log loss
is:",log_loss(y_test, predict_y, labels=clf.classes_, eps=1e-15))
fig, ax = plt.subplots()
ax.plot(alpha_values, log_error_values,c='g')
for i, txt in enumerate(np.round(log error values,3)):
    ax.annotate((alpha values[i],np.round(txt,3)),
(alpha values[i],log error values[i]))
plt.grid()
plt.title("Cross Validation Error for each alpha")
plt.xlabel("Alpha i's")
plt.ylabel("Error measure")
plt.show()
best alpha = np.argmin(log error values)
clf = SGDClassifier(alpha=alpha_values[best_alpha], penalty='l2',
loss='log', random state=42)
clf.fit(X_train, y_train)
sig clf = CalibratedClassifierCV(clf, method="sigmoid")
sig clf.fit(X train, y train)
For values of alpha = 1e-05 The log loss is: 0.6585300338196388
For values of alpha = 0.0001 The log loss is: 0.6585300338196388
For values of alpha = 0.001 The log loss is: 0.6585300338196388
For values of alpha = 0.01 The log loss is: 0.6412728467929882
For values of alpha = 0.1 The log loss is: 0.5827239969320069
For values of alpha = 1 The log loss is: 0.5990993513923978
For values of alpha = 10 The log loss is: 0.6295331603050178
```





```
CalibratedClassifierCV(estimator=SGDClassifier(alpha=0.1, loss='log',
                                               random state=42))
print("Best value of alpha - ", alpha_values[best_alpha])
predict y = sig clf.predict proba(X train)
print("The train log loss is:",log loss(y train, predict y,
labels=clf.classes_, eps=le-15))
predict y = sig clf.predict proba(X test)
print("The test log loss is:",log loss(y test, predict y,
labels=clf.classes_, eps=1e-15))
predicted y =np.argmax(predict y,axis=1)
Best value of alpha - 0.1
The train log loss is: 0.5837994604444074
The test log loss is: 0.5827239969320069
from sklearn.metrics import confusion matrix, classification report,
accuracy score
print("Accuracy Score - ", accuracy_score(y test, predicted y))
Accuracy Score - 0.6776942293897944
print("Classification Report")
print(classification report(y test, predicted y))
```

```
Classification Report
                           recall f1-score
              precision
                                              support
                   0.69
                             0.87
                                       0.77
                                                 51005
           1
                   0.61
                             0.34
                                       0.44
                                                 29853
                                                 80858
    accuracy
                                       0.68
                   0.65
                             0.61
                                       0.61
                                                 80858
   macro avq
weighted avg
                   0.66
                             0.68
                                       0.65
                                                 80858
def confusion matrix plot(test values, prediction values):
    cf matrix = confusion matrix(test values, prediction values)
    precision details =(((cf matrix.T)/(cf matrix.sum(axis=1))).T)
    recall details = (cf matrix/cf matrix.sum(axis=0))
    plt.figure(figsize=(20,4))
    labels = [1,2]
    cmap=sns.light palette("blue")
    plt.subplot(1, 3, 1)
    sns.heatmap(cf matrix, annot=True, cmap=cmap, fmt=".3f",
xticklabels=labels, yticklabels=labels)
    plt.xlabel('Predicted Value')
    plt.ylabel('Original Value')
    plt.title("Confusion matrix")
    plt.subplot(1, 3, 2)
    sns.heatmap(recall details, annot=True, cmap=cmap, fmt=".3f",
xticklabels=labels, yticklabels=labels)
    plt.xlabel('Predicted Value')
    plt.vlabel('Original Value')
    plt.title("Precision matrix")
    plt.subplot(1, 3, 3)
    sns.heatmap(precision details, annot=True, cmap=cmap, fmt=".3f",
xticklabels=labels, yticklabels=labels)
    plt.xlabel('Predicted Value')
    plt.ylabel('Original Value')
    plt.title("Recall matrix")
    plt.show()
confusion_matrix_plot(y_test, predicted_y)
```

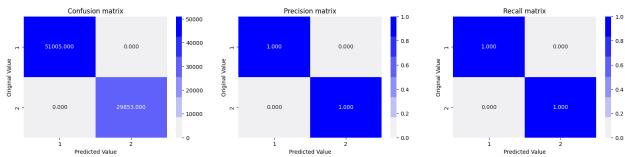


SVM

```
clf = SGDClassifier(alpha=alpha values[best alpha], penalty='l1',
loss='hinge', random state=42)
clf.fit(X train, y train)
sig clf = CalibratedClassifierCV(clf, method="sigmoid")
sig clf.fit(X train, y train)
CalibratedClassifierCV(estimator=SGDClassifier(alpha=0.1,
penalty='l1',
                                                 random state=42))
predict y = sig clf.predict proba(X test)
predicted y =np.argmax(predict y,axis=1)
print(accuracy_score(y_test, predicted_y))
print(classification_report(y_test, predicted_y))
0.6307972000296816
              precision
                            recall f1-score
                                               support
                                        0.77
           0
                   0.63
                              1.00
                                                 51005
           1
                   0.00
                              0.00
                                        0.00
                                                 29853
                                        0.63
                                                 80858
    accuracy
                   0.32
                              0.50
                                        0.39
                                                 80858
   macro avg
weighted avg
                   0.40
                              0.63
                                        0.49
                                                 80858
```

XGBoost

```
feature types=None,
              gamma=None, grow policy=None, importance type=None,
              interaction constraints=None, learning rate=None,
max bin=None,
              max cat threshold=None, max cat to onehot=None,
              max delta step=None, max depth=5, max leaves=None,
              min child weight=None, missing=nan,
monotone constraints=None,
              multi strategy=None, n estimators=300, n jobs=-1,
              num parallel tree=None, random state=25, ...)
y pred test=clf.predict proba(X test)
y pred train=clf.predict proba(X train)
log loss train = log loss(y train, y pred train, eps=le-15)
log_loss_test=log_loss(y_test,y_pred_test,eps=le-15)
print('Train log loss = ',log loss train,' Test log loss =
,log loss test)
predicted y=np.argmax(y pred test,axis=1)
confusion matrix plot(y test,predicted y)
Train log loss = 6.091918669980021e-06 Test log loss =
6.091016389706224e-06
```



```
print("Classification Report")
print(classification report(y test, predicted y))
Classification Report
              precision
                            recall
                                    f1-score
                                                support
           0
                    1.00
                              1.00
                                         1.00
                                                  51005
           1
                    1.00
                              1.00
                                         1.00
                                                  29853
    accuracy
                                         1.00
                                                  80858
                                         1.00
                                                  80858
   macro avq
                    1.00
                              1.00
                              1.00
                                         1.00
                                                  80858
weighted avg
                    1.00
print("Accuracy Score - ", accuracy_score(y_test, predicted_y))
Accuracy Score - 1.0
```

Seems XGBoosting is overfitting the data

These traditional models are giving moderate results so we can try deep learning models to get better results. Due to resource and time constraints couldn't those.