Phase 4 Reccomendation Modeling

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In this notebook we will be be using the CRISP-DM process to walk through using Faiss K-means algorithm and naive Bayes to predict sentiment on space travel related tweets and YouTube comments.

```
1 # !pip install faiss-gpu
 2 # !pip install vaderSentiment
 1 # from google.colab import drive
 2 # drive.mount('/content/drive')
 1 import faiss
 2 import math
 3 import pandas as pd
 4 import numpy as np
 5 from datetime import datetime
 1 import scipy
 2 from scipy.linalg import norm
 3 from scipy.sparse import csr matrix, coo matrix, hstack, vstack
 4 from imblearn.combine import SMOTEENN
 1 from textblob import TextBlob
 2 from vaderSentiment.vaderSentiment import SentimentIntensityAnalyzer
 1 from sklearn.decomposition import TruncatedSVD
 2 from sklearn.naive_bayes import GaussianNB, BernoulliNB, ComplementNB
 3 from sklearn.preprocessing import MaxAbsScaler
 5 from sklearn.feature extraction.text import TfidfVectorizer, \
                                                 CountVectorizer
 7 from sklearn.metrics import silhouette samples, confusion matrix, \
 8
                         accuracy_score, recall_score, precision_score, \
 9
                         roc_curve, auc, roc_auc_score, confusion_matrix, \
                         classification report, f1 score, r2 score
10
11 from sklearn.model selection import GridSearchCV, cross val score, \
12
                                       train test split, RandomizedSearchCV
```

¹ from mlxtend.plotting import plot confusion matrix

```
1 %matplotlib inline
2 import matplotlib.pyplot as plt
3 from matplotlib import cm
```

Functions

```
1 def get_sentiment_an(df):
 2
 3 Given a pandas dataframe, this will shuffle it & get sentiment
 4
    analysis values from TextBlob & VaderSentiment & assign columns inplace.
 5
    Will return completed dataframe & dataframe columns we'll be using for
    modeling as a coo_matrix.
 6
 7
                   Columns returned:
    'polarity' : mean of text sentiment (TextBlob)
 8
 9
     'neg': negative sentiment value (VaderSentiment)
    'pos': positive sentiment value (VaderSentiment)
10
11
    'neu': neutral sentiment value (VaderSeniment)
12
13
    # shuffling our dataframe so data is no longer sorted
    df = df.sample(frac = 1)
14
15
    df = df.reset index().rename(columns={'index':'id'})
16
    # getting rid of all unnessecary columns
17
    df = df[['favorite count', 'repost count', 'text', 'id']]
18
    # feature generation using polarization
19
    df['polarity'] = df['text'].apply(lambda x:
20
                                       TextBlob(x).sentiment.polarity)
21
    # feature generation using intensity analyzer
22
    analyzer = SentimentIntensityAnalyzer()
    df['pol'] = ''
23
24
    for i in range(len(df)):
25
      sentence = df.at[i, 'text']
26
      df.at[i, 'pol'] = analyzer.polarity_scores(sentence)
27
    for i in range(len(df)):
      for h in ['neg', 'neu', 'pos']:
28
        df.at[i, h] = float(df['pol'][i][h])
29
30
    #turning our other variables into a matrix to combine our features
31
    maxab = MaxAbsScaler()
    drop = ['text', 'pol', 'id']
32
33
    hstack = maxab.fit_transform(coo_matrix(df.drop(labels = drop, axis=1)))
34
    return df, hstack
 1 def word transform(df):
 2
    Pass in a pandas dataframe and this will return a coo matrix
   featuring the TFIDF vectorization of your text, with a maximum
 4
 5
    of 100 features of single words and 100 features of bigrams.
    11 11 11
 6
 7
    sing = TfidfVectorizer(max_features=100).fit_transform(df['text'])
```

```
bi = TfidfVectorizer(ngram_range = (2, 2),
 8
 9
                          max features=100).fit transform(df['text'])
10
    vect = coo matrix(np.append(bi.toarray(), sing.toarray(), axis=1))
11
    return vect
 1 def svd_pca(vect, haystack):
    11 11 11
 2
    Pass in your coo_matrix of your word vectors & the scaled
 3
    features from your dataframe this will perform Principal Component
 4
 5
    Analysis in order to keep your most relevant 100 features, calculated
 6
     in chunks to avoid RAM issues. It will return a numpy
 7
     array containing all of the data for your model.
 8
 9
    svd = TruncatedSVD(n components=95, n iter=250000,
                        random_state=0, algorithm='arpack').fit(vect)
10
11
    # splitting it so we never keep it all in RAM at once as an array
12
    splits = []
13
    shape = vect.shape[1]
    tbs = vect
14
15
    # since you cant directly index a matrix the easiest way is train_test
16
    # chose 50000 since my computer can handle it
17
    while shape > 50000:
18
         shape -= 50000
19
        # first, run through the whole matrix
20
         split1, split2 = train_test_split(tbs, random_state=0,
21
                                           train size=shape)
22
        #all of our leftover rows (not in the 50k) are in split2
23
        tbs = split2
24
        # put the chunk of our matrix into a list to iterate over later
25
         splits.append(split1)
26
    # append the remainder (whatever was left < 50k) to the end of our list
27
    splits.append(tbs)
28
    # start a new matrix by fit transforming our first chunk
29
    init = coo matrix(svd.fit transform(splits[0]))
30
    # start a loop that will fit transform and stack matricies together
31
    for item in splits[1:]:
32
        # fit transform will return it as a dense array
         init = vstack([init, coo matrix(svd.fit transform(item))])
33
34
    # combine our text matrix & our repost/favorite matrix
35
    return np.append(haystack.toarray(),
36
                      init.toarray(), axis=1).astype(np.float32)
 1 def vect and stck(dataframe, first run):
    Pass in a DataFrame & this function will preprocess and return your
 3
    dataframe & all of the columns nessecary to model as an array
 4
    The first step is to get sentiment analysis scores,
 5
    which is only nessecary the first time you run this, otherwise
 6
 7
    they can be turned to False to save time,
 8
    sentiment analysis is followed by TDIDF vectorizaton on
```

```
9
    single words & bigrams, and lastly it ends with PCA to reduce
10
    our dimensions to a reasonable amount. Make drop clust True to
    return a copy of your dataframe with no cluster column before
11
12
    rerunning your model on it.
13
14
    if first run:
15
      # getting sentiment analysis columns the first time
      dataframe, haystack = get sentiment an(dataframe)
16
17
    else:
18
    # keeping only of the columns we need
      clmn = ['neg', 'neu', 'pos', 'polarity',
19
20
               'repost_count', 'favorite_count']
21
      haystack = coo matrix(dataframe[clmn])
22
    # word vectorizing
    vect = word transform(dataframe)
23
24
    # return our matrix with PCA done to reduce
25
    # dimensionality to 100 features as was reccomended
    # for PCA in the documentation
26
27
    return svd_pca(vect, haystack), dataframe
 1 def plt_elbow(matrix, df, display):
    11 11 11
 2
 3
    Plots your clusters from k 1-15 and returns the best one as an int,
 4
    if display is true it will display the elbow plot, if not it will
 5
    just return your value.
 6
 7
    max = 15 if display else 5
 8
    K = range(1, max+1)
 9
    # start numpy arrays to store results in
    inrt = np.zeros(max)
10
11
    diff = np.zeros(max)
12
    diff2 = np.zeros(max)
13
    diff3 = np.zeros(max)
    for k in K:
14
15
      kmeans, predictions, not k = clusterizer(matrix, df, False, k=k)
16
      inrt[k - 1] = kmeans.obj[-1]
17
      # first difference
18
      if k > 1:
19
           diff[k-1] = inrt[k-1] - inrt[k-2]
20
      # second difference
21
      elif k > 2:
22
           diff2[k-1] = diff[k-1] - diff[k-2]
23
      # third difference
24
      elif k > 3:
25
           diff3[k - 1] = diff2[k - 1] - diff2[k - 2]
26
    # use differences & numpy argmin to determine best cluster
27
    elbow = np.argmin(diff3[3:]) + 3
28
    if display:
29
      print(f'Elbow {str(elbow)}')
      plt.plot(K, inertias, 'b*-')
30
31
      plt.plot(K[elbow-1], inertias[elbow-1], marker='o',
```

```
32
                markersize=12, markeredgewidth=2, markeredgecolor='r')
      plt.ylabel('Inertia')
33
      plt.xlabel('K')
34
35
      plt.show()
36
    return int(elbow)
 1 def chunked samples(dataframe, stack, fit predict, drop):
 2
 3
    Given your dataframe, feature array/matrix & predictions
 4
    Computes Silhouette Scores using sklearn and chunking
    the data into sets of 75000 and returns a deep copy of
 5
    your dataframe with the silhouette scores in the column
 6
    'sil' Scores will not be perfect as this function chunks them, and
 7
    therefore cannot get complete pairwise distances
 8
 9
    but without chunking silhouette scores cannot be run on
    any computer or service we can find and it will be pretty close.
10
    Use drop = True to calculate the z-score of your silhouette scores
11
12
    and get rid of the bottom 10%, assuming they're likely to be outliers.
    11\ 11\ 11
13
14
    start = 0
    clust = 75000
15
    # since this is about the most my computer can handle we'll
17
    # stick to a cap of 75,000
    for i in range(math.ceil(stack.shape[0]/75000)):
18
19
      # gathering our chunks
      df = dataframe[start:start+clust].reset index(drop=True).copy()
20
21
      fp = np.ravel(fit predict[start:start+clust][0:])
22
      # making sure our dtypes are correct
23
      if type(stack) != np.ndarray:
24
        hstk = stack.toarray()[start:start+clust][0:]
25
      else:
26
        hstk = stack[start:start+clust][0:]
27
      # getting our silhouette comments
      df['sil'] = silhouette samples(hstk, fp)
28
29
      # we want to return our dataframe at the end so
      # on our first ieration we'll start with a chunk
30
31
      # and append to it every future iteration
32
      head = df if i == 0 else head.append(df)
      if drop:
33
        # if drop = True, this will calculate and drop the
34
        # lowest ~10th percentile of our silhouette scores
35
        # assuming they're likely to be outliers
36
37
        mean = head['sil'].mean()
        std = head['sil'].std()
38
39
        for j in head.index:
          head.at[j, 'silstd'] = (head.at[j, 'sil'] - mean)/std
40
        head = head.loc[head['silstd'] > -1.28].drop('silstd', 1)
41
42
      head.reset index(drop=True, inplace=True)
43
       start += 75000
44
    return head
```

```
1 def quick silhouette(dataframe, stack, predict):
 2
 3
    Pass in your dataframe, your feature array/matrix,
 4
    & your predictions, and this will use chunked samples
 5
    to get your score & try to quickly form a matplotlib visual
 6
    Scores won't be 100% accurate as you cannot get pairwise distances
 7
    with chunking, but they use a sample size of 75,000 so they should
    be close, and the decrease in accuracy is well worth the 6-7 hours
    in speed it saves
 9
    11 11 11
10
11
    #first we calculate our silhouette scores
    dataframe = chunked samples(dataframe, stack, predict, False)
12
    # convert our predictions and their scores to numpy arrays
13
14
    predictions = dataframe.cluster.to numpy()
15
    silhouette vals = dataframe.sil.to numpy()
16
    # get our labels
17
    labels = np.unique(predictions)
18
    # start plotting
19
    y_ax_lower, y_ax_upper = 0, 0
20
    yticks = []
21
    # go through our labels and assign our predictions & values to them
22
    for i, c in enumerate(labels):
23
      c silhouette vals = silhouette vals[predictions == c]
24
      # sort to keeo our clusters together
25
      c silhouette vals.sort()
26
      # change plot size & ticks accordingly
      y ax upper += len(c silhouette vals)
27
      yticks.append((y_ax_lower + y_ax_upper) / 2.)
28
29
      # get colors & plot!
30
      color = cm.jet(float(i) / labels.shape[0])
31
      plt.barh(range(y_ax_lower, y_ax_upper), c_silhouette_vals,
32
                 height=1.0, color=color)
      y ax lower += len(c silhouette vals)
33
34
    # find our average and visualize it
35
    plt.axvline(np.mean(silhouette vals), color="red", linestyle="--")
36
    # fix up our labels and ticks
37
    plt.yticks(yticks, labels + 1)
38
    plt.ylabel('Cluster')
39
    plt.xlabel('Silhouette Coefficient')
40
    plt.show()
 1 def get_imb_clstr(dataframe):
 2
    Pass in your dataframe and get the cluster with the most values in it
   returned as an interger
 4
    11 11 11
 5
    clstr = dataframe['cluster'].value_counts()
 6
 7
    # largest cluster gets returned at the first index
 8
   clstr = int(clstr.to frame().reset index()['index'][0])
    return clstr
```

```
1 def reset clusters(new predict, dataframe):
 2
    Pass in your new predictions and your dataframe
 3
    and it will renumber all of your clusters.
 4
 5
    Be sure to use this before making visuals, if you don't
 6
    clusters that have been reassigned will just appear to be missing.
    Edits your dataframe inplace. This is intentionally slower in
 7
    the hopes of not crashing my ram by iterating over each line one by one
 8
 9
10
    # get a list of our starting clusters
11
    clusters = [int(x) for x in np.unique(new predict)]
12
    # generate a list of new clusters
    _range = list(range(len(_clusters)))
13
14
    to rename = {}
15
    # generate a dictionary with key value pairs of new : old clusters
16
    for num in range:
      to rename[ clusters[num]] = num + 1
17
18
    # go through the dataframe and edit our clusters
19
    for i in range(len(dataframe)):
20
      dataframe.at[i, 'cluster'] = to rename[dataframe.at[i, 'cluster']]
 1 def add_new_clst(df, dataframe, k, num_clust):
 2
    Pass in your new predictions and your dataframe
 3
    and it will reassign the cluster you modeled to its new
 4
 5
    cluster assignments. Edits dataframe inplace.
 6
 7
    # when generating new clusters after your first modeling
    # you'll need to assign them to new unique cluster numbers
 8
 9
    for i in df.index:
      # use the id's we assigned earlier to match up our clusters
10
      id = df.at[i, 'id']
11
      index = dataframe.loc[dataframe['id'] == _id].index[0]
12
      # make sure we always get unique clusters, and that numbers being
13
      # reassigned to the same cluster always turn out the same
14
15
      n cluster = (df.at[i, 'cluster'] + 2) * 10 + num clust*2
16
      # reassign our clusters
17
      dataframe.at[index, 'cluster'] = n cluster
 1 def clusterizer(matrix, dataframe, calc k, k=3, display elbow=False):
 3
    Pass in your feature array/matrix and your dataframe
    If you would like to have k be calculated for you using the
 4
    elbow method with a maximum number of 5 clusters, do so by
    setting calc k = True, otherwise, k will be 3
 6
    11 11 11
 7
 8
    if calc k:
 9
      if display elbow:
        k = nlt elhow(matrix dataframe True)
10
```

```
11
      else:
12
        k = plt elbow(matrix, dataframe, False)
13
    # making sure our dtypes always line up
    if type(matrix) != np.ndarray:
14
15
     matrix = matrix.toarray()
    # we dont want our clusters to be imbalanced, so we want them
16
17
    # to be roughly equal to the number of points/ number of clusters
18
    shape = matrix.shape[0]
    # start our k-means
19
    kmeans = faiss.Kmeans(d = matrix.shape[1], k = k, nredo = 100,
20
                           update index = True, seed = 42, verbose=True,
21
22
                           max points per centroid = math.ceil(shape/k),
23
                           niter = 20)
24
    # train our k means
25
    kmeans.train(matrix)
    # get our predictions
26
27
    predict = kmeans.index.search(matrix, 1)[1]
28
    # assign them inplace to our dataframe
29
    dataframe['cluster'] = predict
30
    return kmeans, predict, k
 1 def get_slice(df, cluster):
    1.1.1
 3
   Pass in your dataframe and return just the values predicted to be in the
   largest cluster.
 4
    1.1.1
 5
   return df.loc[df['cluster'] == cluster].reset_index(drop=True)
 1 def showconfusionmatrix(y_t, y_hat_t, title):
 2
      Plots confusion matrix for provided y_train, y_hat_train
 3
      OR y test, y hat test using mlxtend
 4
 5
 6
      # assign parameters to dictionary so line wont exceed 78 char
 7
      p = {conf mat:confusion matrix(y t, y hat t), colorbar:True,
            show_absolute:True, show_normed:True, cmap:'jet'}
 8
      # gather and plot confusion matrix
 9
10
      fig, ax = plot_confusion_matrix(**p)
11
      # set title and axis names
12
      ax.set title(f'{title} Model')
13
      ax.set_ylabel('Actual Data')
      ax.set xlabel('Predicted Data')
14
15
      plt.show()
 1 def printscores(y_test, y_hat_test, X_test, y_train,
                   y_hat_train, X_train, model):
       """Uses sklearn.metrics to compute testing and training scores
 3
 4
      Results include: Accuracy, Precision, Recall, F1, Residual Counts,
 5
      Residuals, CV Accuracy With Standard Deviation & R^2"""
```

K - pic Cidow(macriin, dacarrame, rrac)

+0

```
6
      print('
 7
       print('Training Accuracy Score: ' +
             str(round(accuracy score(y train, y hat train)* 100, 4)))
 8
       train_precision_scores = []
 9
      for item in precision_score(y_test, y_hat_test, average=None):
10
11
        train precision scores.append(round(item, 2))
12
       print('Training Precision Scores: ' + str(train_precision_scores))
      train recall scores = []
13
      for item in recall_score(y_test, y_hat_test, average=None):
14
15
        train_recall_scores.append(round(item, 2))
       print('Training Recall Scores: ' + str(train recall scores))
16
17
      train_f1_scores = []
18
       for item in f1_score(y_test, y_hat_test, average=None):
19
        train_f1_scores.append(round(item, 2))
       print('Training F1 Scores: ' + str(train f1 scores))
20
21
      print('__
22
      trainresiduals = np.abs(y_train - y_hat_train)
      print('Training residual counts:')
23
      print(str(pd.Series(trainresiduals).value_counts())[:-29])
24
25
      print('
      print('Testing Accuracy Score: ' +
26
27
             str(round(accuracy_score(y_test, y_hat_test) * 100, 4)))
28
      test precision scores = []
29
       for item in precision_score(y_test, y_hat_test, average=None):
30
        test_precision_scores.append(round(item, 2))
       print('Testing Precision Scores: ' + str(test_precision_scores))
31
32
      test recall scores = []
33
      for item in recall_score(y_test, y_hat_test, average=None):
        test_recall_scores.append(round(item, 2))
34
35
       print('Testing Recall Scores: ' + str(test_recall_scores))
      test_f1_scores = []
36
37
      for item in f1_score(y_test, y_hat_test, average=None):
        test_f1_scores.append(round(item, 2))
38
39
      print('Testing F1 Scores: ' + str(test_f1_scores))
40
       print('_
41
      testresiduals = np.abs(y_test - y_hat_test)
42
      print('Testing residual counts:')
      print(str(pd.Series(testresiduals).value_counts())[:-29])
43
44
      print('_
      r2 scoretst = r2 score(y test, y hat test)
45
46
      print(f'Testing R2 Score: {round(r2_scoretst, 4)}')
47
      r2_scoretrn = r2_score(y_train, y_hat_train)
       print(f'Training R2 Score: {round(r2 scoretrn, 4)}')
48
49
      print('
50
      print('Cross validated model accuracy:')
       scores = cross_val_score(model, X_test, y_test, cv=10)
51
52
      mean = round(scores.mean(), 4)
       std = round(scores.std())
53
54
      print(f'Testing: {mean} with a standard deviation of {std}')
      scores = cross val score(model, X train, y train, cv=10)
55
56
      mean = round(scores.mean(), 4)
57
       std = round(scores.std())
```

```
print(f'Training: {mean} with a standard deviation of {std}')
58
59
1 def printreports(y_test, y_hat_test, y_train, y_hat_train):
2
      Provided all input & predicted y values, prints classification
3
      report for training and testing data right next to each other"""
4
5
      print('
                                                               ')
                  Testing Report:')
      print('
6
      report1 = classification_report(y_test, y_hat_test)
7
      print(report1)
9
      print('____
      print(' Training Report:')
10
    report2 = classification_report(y_train, y_hat_train)
11
12
      print(report2)
      print('____
13
```

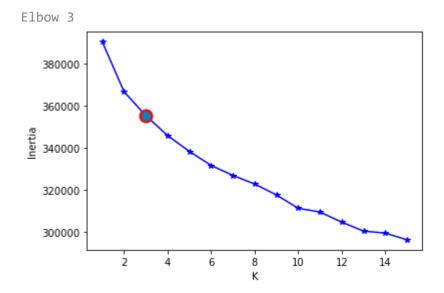
Unsupervised Model Setup

First, we begin by opening up our data and doing some preprocessing, our data has already been cleaned and tokenized and lemmatized, so here we mostly focus on feature generation and dimensionality reduction

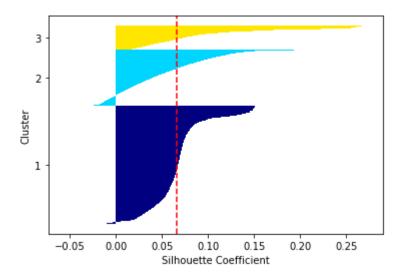
First Unsupervised Model

Model Evaluation

Looking at our initial model, our clusters aren't particuarly well defined, and even with multiple attempts at what k should be, they're not showing any substantial improvement



1 # initial silhouette scores very low
2 quick_silhouette(first_model, matrix, predict)



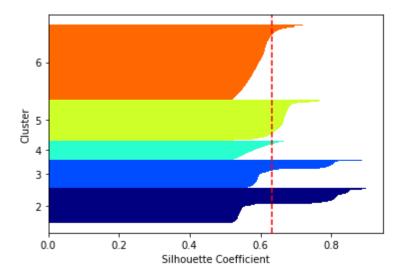
Unsupervised Model Refinement

1 # start our model & make our initial predictions

```
# ni.obbing onfitter.2 and 2001111108 ont. nafa
 Э
    df = chunked_samples(df, matrix, predict, True)
 6
    df = df.sample(frac=1).reset index(drop=True)
    # processing our data without outliers
 8
 9
    matrix, df = vect and stck(df, False)
10
    # re-modeling our data with less outliers
    bttr_mn, bttr_prd, k = clusterizer(matrix, df, False, 3, False)
11
12
    # slicing our dataframe to get disproportionate cluster
13
    cl = get_imb_clstr(df)
    cut matrix, cut df = vect and stck(get slice(df, cl), False)
14
15
    # plugging our new k value into a new model optimized for our cluster
16
    clst_mn, clst_prd, k = clusterizer(cut_matrix, cut_df, True, 3, False)
17
    # updating our original datframe with our new clusters clusters
    add_new_clst(cut_df, df, k, num_clust)
18
19
    # updating our predictions
20
    predict = df['cluster'].to numpy()
21
    # updating our largest cluster
22
    cl = get imb clstr(df)
23
    # keeping a count of clusters
24
    num clust += k-1
```

Model Evaluation

```
1 # grab our final clusters & plot our silhouettes
2 predict = df['cluster'].to_numpy()
3 reset_clusters(predict, df)
4 quick silhouette(df, matrix, predict)
```



```
1 # calculating how long our modeling took & how much data was lost
2 end = [len(df), datetime.now()]
3 time = end[1]-start[1]
4 print(f' {round(end[0]/start[0], 4)}% Unclassifiable')
5 print(f' Model took {round(time.total_seconds()/60, 2)} min to complete')
```

```
0.5903% Unclassifiable
Model took 126.75 min to complete
```

Our model showed substantial improvement after iterating through & reclustering, losing minimal amounts of data. With our mean silhouette scores looking good and in a reasonable range, we move on to taking a look at visulizing our clusters and descerning the differences between them, which you can find in our notebook /wordclouds.ipynb

Bayesian Modeling

We chose bayesian modeling because it is said to work especially well for natural language processing, 2 of the methods available from Sklearn were not compatible with our data, but the other 2 are tested & refined below.

Testing models

There are our baseline models for making predictions off of our data

```
1 # the baseline model is actually incredibly close to our results
2 # while tuning it we discovered a few variables that did have considerable
3 # effect, so here is one to help showcase the difference
4 gau_model = GaussianNB(var_smoothing=1)
5 y_hat_test = gau_model.fit(X_train, y_train).predict(X_test)
6 y_hat_train = gau_model.predict(X_train)

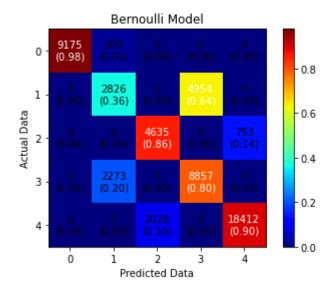
1 showconfusionmatrix(y_test, y_hat_test, 'Gaussian')
2 # certainly too good to be true
```

```
1.0
          9384
          (1.00)
                                            0.8
                7782
(1.00)
       1
    Actual Data
                                           0.6
                      5388
                      (1.00)
                                           0.4
                            11130
                            (1.00)
                                           0.2
                                  20334
                                  (0.99)
1 print('
                          Gaussian Model')
2 print('
3 printscores(y_test, y_hat_test, X_test, y_train, y_hat_train, X_train, gau_model)
                     Gaussian Model
    Training Accuracy Score: 99.9006
    Training Precision Scores: [1.0, 1.0, 0.98, 1.0, 1.0]
    Training Recall Scores: [1.0, 1.0, 1.0, 1.0, 0.99]
    Training F1 Scores: [1.0, 1.0, 0.99, 1.0, 1.0]
    Training residual counts:
         30
    Testing Accuracy Score: 99.8042
    Testing Precision Scores: [1.0, 1.0, 0.98, 1.0, 1.0]
    Testing Recall Scores: [1.0, 1.0, 1.0, 1.0, 0.99]
    Testing F1 Scores: [1.0, 1.0, 0.99, 1.0, 1.0]
    Testing residual counts:
    Testing R2 Score: 0.9966
    Training R2 Score: 0.998
    Cross validated model accuracy:
    Testing: 0.7858 with a standard deviation of 0.0
    Training: 0.999 with a standard deviation of 0.0
```

As you can see from our scores, this model is way too good to be true and is not producing a consistent output despite its high scores

```
1 bern_model = BernoulliNB()
2 y_hat_test = bern_model.fit(X_train, y_train).predict(X_test)
3 y_hat_train = bern_model.predict(X_train)
```

Gaussian Model



- 1 print(' Bernoulli Model')
 2 print(' _____')
- 3 printreports(y_test, y_hat_test, y_train, y_hat_train)

Bernoulli Model

	Testing Re	port:		
	precision	recall	f1-score	support
1	1.00	0.98	0.99	9384
2	0.53	0.36	0.43	7782
3	0.70	0.86	0.77	5388
4	0.64	0.80	0.71	11130
5	0.96	0.90	0.93	20440
accuracy			0.81	54124
macro avg	0.77	0.78	0.77	54124
weighted avg	0.81	0.81	0.81	54124

		Report:	Training	
support	f1-score	recall	precision	
61186	0.99	0.98	1.00	1
61186	0.46	0.37	0.62	2
60348	0.89	0.87	0.91	3
61185	0.66	0.80	0.56	4
58953	0.89	0.91	0.88	5
302858	0.79			accuracy
302858	0.78	0.79	0.79	macro avg
302858	0.78	0.79	0.79	weighted avg

```
1 print('
                         Bernoulli Model')
2 print('
3 printscores(y_test, y_hat_test, X_test, y_train, y_hat_train, X_train, bern_model)
                    Bernoulli Model
\Box
    Training Accuracy Score: 78.5124
    Training Precision Scores: [1.0, 0.53, 0.7, 0.64, 0.96]
    Training Recall Scores: [0.98, 0.36, 0.86, 0.8, 0.9]
    Training F1 Scores: [0.99, 0.43, 0.77, 0.71, 0.93]
    Training residual counts:
        237781
    2
        6
    Testing Accuracy Score: 81.1193
    Testing Precision Scores: [1.0, 0.53, 0.7, 0.64, 0.96]
    Testing Recall Scores: [0.98, 0.36, 0.86, 0.8, 0.9]
    Testing F1 Scores: [0.99, 0.43, 0.77, 0.71, 0.93]
    Testing residual counts:
        43905
    2
    Testing R2 Score: 0.6812
    Training R2 Score: 0.5753
    Cross validated model accuracy:
    Testing: 0.8185 with a standard deviation of 0.0
    Training: 0.7848 with a standard deviation of 0.0
```

This model is only slightly better than random chance overall, and in some clusters, it's even worse! Certainly not the route we want to be going.

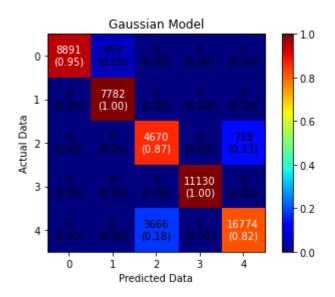
Final Model

Here, with a slight parameter tune, we see our final model come into realization with solid, consistent results.

```
1 final_model = GaussianNB(var_smoothing= 1e-200)
2 y_hat_test = final_model.fit(X_train, y_train).predict(X_test)
3 y_hat_train = final_model.predict(X_train)
```

Evaluation

```
1 showconfusionmatrix(y_test, y_hat_test, 'Gaussian')
```



1 print(' Final Gaussian Model')
2 print(' _____')

3 printreports(y_test, y_hat_test, y_train, y_hat_train)

Final Gaussian Model

	T			
	Testing F			
	precision	recall	f1-score	support
1	1.00	0.95	0.97	9384
2	0.94	1.00	0.97	7782
3	0.56			5388
4	1.00			
5	0.96			
accuracy			0.91	54124
macro avg	0.89	0.93	0.90	
weighted avg	0.93			
	Training	Report:		
	precision	recall	f1-score	support
1	1.00	0.95	0.97	61186
2	0.95	1.00	0.98	61186
3	0.84	0.86	0.85	60348
4	1.00	1.00	1.00	61185
5	0.86	0.83	0.84	58953
accuracy			0.93	302858
macro avg	0.93	0.93	0.93	302858
weighted avg	0.93	0.93	0.93	302858

1 print(' Final Gaussian Model')

```
2 print('
3 printscores(y_test, y_hat_test, X_test,
             y train, y hat train, X train, final model)
                 Final Gaussian Model
   Training Accuracy Score: 93.0278
   Training Precision Scores: [1.0, 0.94, 0.56, 1.0, 0.96]
   Training Recall Scores: [0.95, 1.0, 0.87, 1.0, 0.82]
   Training F1 Scores: [0.97, 0.97, 0.68, 1.0, 0.88]
   Training residual counts:
        281742
   2
         1
   Testing Accuracy Score: 90.9892
   Testing Precision Scores: [1.0, 0.94, 0.56, 1.0, 0.96]
   Testing Recall Scores: [0.95, 1.0, 0.87, 1.0, 0.82]
   Testing F1 Scores: [0.97, 0.97, 0.68, 1.0, 0.88]
   Testing residual counts:
        49247
   2
   Testing R2 Score: 0.8572
   Training R2 Score: 0.875
   Cross validated model accuracy:
   Testing: 0.9145 with a standard deviation of 0.0
   Training: 0.93 with a standard deviation of 0.0
```

In conclusion,

Even though we failed to find **strong** sentiment between our clusters, our model itself found clear enough differences between them to make meaningful and replicable predictions and the knowledge we came away from this with has given us great inspiration to make buisness reccomendations that will help your company move forward in the growing space hospitality industry.

Our reccomendations are:

- 1. Leverage use of the word 'Space' Space is the most commonly found word within the discourse, with that in mind, tagging along a positive sentiment will continue to build a positive arena around space and space travel & extend your reach to the largest audience.
- 2. Evoke emotion and drive the conversation. Customers think with their hearts, and by utilizing words like "love", which are some of the most popular non-science words we found, you are likely to create a positive sentiment around your product. That positive emotional response adds tremendous value to your marketing capabilities.

3. Partner with the best Institutions like NASA and SpaceX not only have scientific headway, but they have built reputations. If you want to get people talking about you, working with buisnesses they already support can help extend your reach, as even with the variation between our clusters, the big names in them stay rather consistent.

We look forward to working with you and appreciate your time. Thank you.