Getting familiar with the dataset

```
import pandas as pd
file_path = 'Project1-ClassificationDataset.csv'

#selected_columns = ["full_text", "summary", "keywords", "publish_date", "authors", "url", "leaf_label", "root_label"]

df = pd.read_csv(file_path)

num_rows, num_columns = df.shape

print(f"Number of rows (samples): {num_rows}")
print(f"Number of columns (features): {num_columns}")

Number of rows (samples): 3476
```

Question 1:

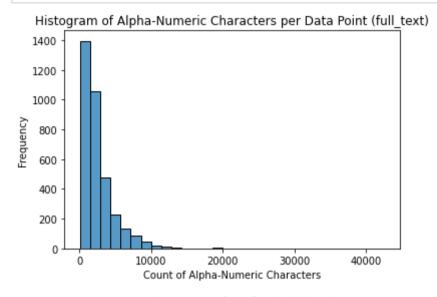
Number of columns (features): 8

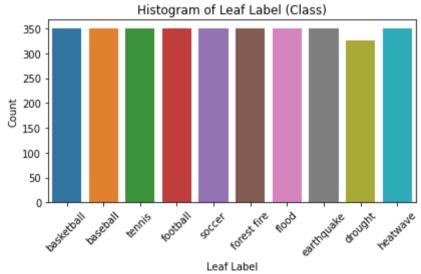
1. Overview: Number of rows (samples): 3476, Number of columns (features): 8

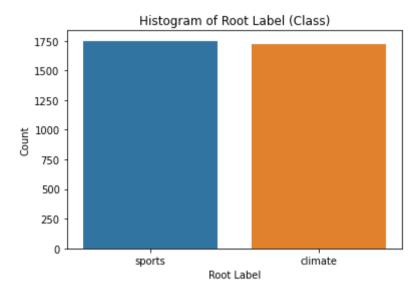
```
In [ ]:
            df.head()
Out[ ]:
                       full text
                                         summary
                                                             keywords
                                                                            publish date
                                                                                                authors
                                                                                                                                                        url leaf label root label
                'Personalize Your 'Personalize Your
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```

```
full text
                                                                                                                                   url leaf label root label
                                                     keywords
                                                                  publish date
                                                                                  authors
                                   summary
                 'The Golden
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In [ ]:
          import matplotlib.pyplot as plt
          import seaborn as sns
          # (a) Histogram for the total number of alpha-numeric characters per data point in the feature 'full text'
          df['alpha numeric count'] = df['full text'].apply(lambda x: sum(c.isalnum() for c in x))
          #plt.figure(figsize=(10, 6))
          sns.histplot(df['alpha numeric count'], bins=30, kde=False)
          plt.title('Histogram of Alpha-Numeric Characters per Data Point (full text)')
          plt.xlabel('Count of Alpha-Numeric Characters')
          plt.ylabel('Frequency')
          plt.show()
          # (b) Histogram for the column 'leaf label' (class)
          #plt.figure(figsize=(10, 6))
          sns.countplot(x='leaf label', data=df)
          plt.title('Histogram of Leaf Label (Class)')
          plt.xlabel('Leaf Label')
          plt.xticks(rotation = 45)
          plt.ylabel('Count')
          plt.tight layout()
          plt.show()
          # (c) Histogram for the column 'root label' (class)
          #plt.figure(figsize=(10, 6))
          sns.countplot(x='root label', data=df)
          plt.title('Histogram of Root Label (Class)')
```

```
plt.xlabel('Root Label')
plt.ylabel('Count')
plt.show()
```







2. Histograms:

- (1). The first histograms shows the frequency of the length of text in the full-text column. The peak on the histogram is in the very left, indicates that this dataset mostly contain texts with very short lengths. As the text length increase, the frequency decrease dramatically, and becomes very small when text length reach to 10,000. This trends suggests that shorter texts are more common in this dataset.
- (2). The second histograms shows the frequency of each class in the "leaf-label" column. This dataset have 10 leaf labels, and the frequency of each label is almost similar, approximately 350 datapoints. The exception is the drought label, which has slightly fewer dataset than the other categories.
- (3). From the third histograms, this dataset only contains two root labels: sports and climate. Each root label have approximately 1750 datapoints.

Binary Classification

1. Splitting the entire dataset into training and testing data

```
import numpy as np
import random
np.random.seed(42)
random.seed(42)
```

```
from sklearn.model_selection import train_test_split

train, test = train_test_split(df[["full_text", "root_label"]], test_size=0.2, random_state=42)

print(f"Number of Training Samples: {train.shape[0]}")
print(f"Number of Testing Samples: {test.shape[0]}")
print(f"Training Samples Shape: {train.shape}")
print(f"Test Samples Shape: {test.shape}")
```

Number of Training Samples: 2780 Number of Testing Samples: 696 Training Samples Shape: (2780, 2) Test Samples Shape: (696, 2)

Question 2:

- Number of Training Samples: 2780
- Number of Testing Samples: 696

2. Feature extraction

```
In [ ]:
         import re
         import nltk
         from sklearn.feature_extraction.text import TfidfVectorizer
         from nltk.corpus import stopwords
         from nltk.stem import WordNetLemmatizer
         from nltk import pos tag
         from sklearn.feature extraction.text import TfidfVectorizer
         from sklearn.feature_extraction.text import CountVectorizer
         from sklearn.feature extraction.text import TfidfTransformer
         from string import punctuation
         from sklearn.feature_extraction import text
         nltk.download('punkt')
         nltk.download('stopwords')
         nltk.download('wordnet')
         nltk.download('averaged_perceptron_tagger')
```

```
[nltk_data] Downloading package punkt to
[nltk_data] C:\Users\Michael_Mi\AppData\Roaming\nltk_data...
```

```
[nltk data]
                      Package punkt is already up-to-date!
        [nltk data] Downloading package stopwords to
        [nltk data]
                        C:\Users\Michael Mi\AppData\Roaming\nltk data...
        [nltk data]
                      Package stopwords is already up-to-date!
        [nltk data] Downloading package wordnet to
                        C:\Users\Michael Mi\AppData\Roaming\nltk data...
        [nltk data]
                      Package wordnet is already up-to-date!
        [nltk data]
        [nltk data] Downloading package averaged perceptron tagger to
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                      Package averaged perceptron tagger is already up-to-
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Out[]: True
In [ ]:
         import re
         import nltk
         from sklearn.feature extraction.text import TfidfVectorizer
         from nltk.corpus import stopwords
         from nltk.stem import WordNetLemmatizer
         from nltk import pos_tag
         from sklearn.feature extraction.text import TfidfVectorizer
         from sklearn.feature extraction.text import CountVectorizer
         from sklearn.feature extraction.text import TfidfTransformer
         from string import punctuation
         from sklearn.feature extraction import text
         nltk.download('punkt')
         nltk.download('stopwords')
         nltk.download('wordnet')
         nltk.download('averaged perceptron tagger')
         nltk.download('omw-1.4')
         def clean(text):
             text = re.sub(r'^https?:\\/.*[\r\n]*', '', text, flags=re.MULTILINE)
             texter = re.sub(r"<br />", " ", text)
             texter = re.sub(r""", "\"", texter)
             texter = re.sub(''', "\"", texter)
             texter = re.sub('\n', " ", texter)
             texter = re.sub(' u '," you ", texter)
             texter = re.sub('`',"", texter)
             texter = re.sub(' +', ' ', texter)
             texter = re.sub(r"(!)\1+", r"!", texter)
             texter = re.sub(r"(\?)\1+", r"?", texter)
             texter = re.sub('&', 'and', texter)
```

```
texter = re.sub('\r', ' ',texter)
    clean = re.compile('<.*?>')
   texter = texter.encode('ascii', 'ignore').decode('ascii')
   texter = re.sub(clean, '', texter)
    if texter == "":
        texter = ""
    return texter
def is number(s):
    return re.match(r'-?\d+\.?\d*', s) is not None
def penn2morphy(penntag):
    """ Converts Penn Treebank tags to WordNet. """
    morphy_tag = {'NN':'n', 'JJ':'a',
                  'VB':'v'. 'RB':'r'}
    try:
        return morphy tag[penntag[:2]]
    except:
        return 'n'
def lemmatize sent(list word):
    # Text input is string, returns array of lowercased strings(words).
    return [wnl.lemmatize(word.lower(), pos=penn2morphy(tag))
            for word, tag in pos_tag(list_word)]
def stem rmv punc(doc): # this should have been at the sentence-level because the pos-tag performs best at sentence-level
    return (word for word in lemmatize sent(analyzer(doc)) if word not in combined stopwords and not is number(word))
train['cleaned text'] = train['full text'].apply(clean)
test['cleaned text'] = test['full text'].apply(clean)
stop words skt = text.ENGLISH STOP WORDS
stop words en = stopwords.words('english')
combined stopwords = set.union(set(stop words en),set(punctuation),set(stop words skt))
wnl = nltk.wordnet.WordNetLemmatizer()
analyzer = CountVectorizer().build analyzer()
count_vectorizer = CountVectorizer(min_df=3, stop_words='english', analyzer=stem_rmv_punc)
tfidf = TfidfTransformer()
X train count = count vectorizer.fit transform(train['cleaned text'])
X_test_count = count_vectorizer.transform(test['cleaned_text'])
```

```
X train tfidf = tfidf.fit transform(X train count)
         X test tfidf = tfidf.transform(X test count)
         [nltk data] Downloading package punkt to
         [nltk data]
                        C:\Users\Michael Mi\AppData\Roaming\nltk data...
         [nltk data]
                      Package punkt is already up-to-date!
         [nltk data] Downloading package stopwords to
         [nltk data]
                        C:\Users\Michael Mi\AppData\Roaming\nltk data...
         [nltk data]
                      Package stopwords is already up-to-date!
        [nltk data] Downloading package wordnet to
         [nltk data]
                        C:\Users\Michael Mi\AppData\Roaming\nltk data...
         [nltk data]
                      Package wordnet is already up-to-date!
         [nltk data] Downloading package averaged perceptron tagger to
         [nltk data]
                        C:\Users\Michael Mi\AppData\Roaming\nltk data...
         [nltk data]
                      Package averaged perceptron tagger is already up-to-
         [nltk data]
                          date!
         [nltk data] Downloading package omw-1.4 to
         [nltk data]
                         C:\Users\Michael Mi\AppData\Roaming\nltk data...
         [nltk data]
                      Package omw-1.4 is already up-to-date!
In [ ]:
         print(f"Shape of TF-IDF-processed train matrix: {X train tfidf.shape}")
         print(f"Shape of TF-IDF-processed test matrix: {X test tfidf.shape}")
        Shape of TF-IDF-processed train matrix: (2780, 13173)
        Shape of TF-IDF-processed test matrix: (696, 13173)
       Question 3:
        answer
```

3a.

3a.1 Lemmatization:

Pros:

- a. The root word is an actual word
- b. Using a dictionary to analysis word and can convery word into root form, have high accuracy and can maintain the original context

cons:

- a. Requires a detailed large dictionary to convert word
- b. Needs more computational resource
- c. Requires the word's correct part of speech to accurately lemmatize it

Effect on size: Lemmatization decreases the size of dictionary by convert words into root form

3a.2 Stemming

Pros:

- a. Simple implementation and running faster than lemmatization
- b. Only removes the end portions of a word to try to find its stem

cons:

a. Do not consider the context and can convert to wrong word, so the accuracy is low

Effect on size: Stemming decreases the size of dictionary, but might not decrease as much as Lemmatization.

3b.

Effect of min df on TF-IDF Matrix: A higher min_df means fewer words are included, filtering out some less common words, but some important subtle information may be lost. A lower min_df means that words that appear less often are allowed to be added, which is beneficial to capturing some uncommon subtle information, but it may contain noise, thus affecting the performance of the model.In short, the choice of min_df is very important

3c. Text Preprocessing Order:

The numbers and punctuations should be removed before Lemmatizing, since they are not contribute to the dictionary.

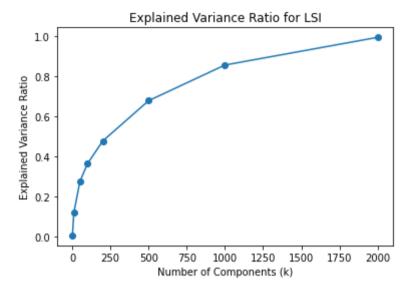
The stopwords should be removed after Lemmatizing, since Lemmatizer requires full sentence as input, including stopwords.

3d. TF-IDF Matrices Shape: The shape of the train matrix is (2780, 13173) and the shape of the test matrix is (696, 13173); both have the same number of rows as the result of Question 2.

3. Dimensionality Reduction

• 4a

```
In [ ]:
         from sklearn.decomposition import TruncatedSVD
         import matplotlib.pyplot as plt
         k values = [1, 10, 50, 100, 200, 500, 1000, 2000]
         explained_variances = []
         # for k in k values:
              svd = TruncatedSVD(n components=k)
              svd.fit(X train tfidf) # X train tfidf
               explained variances.append(svd.explained variance ratio .sum())
         svd list = []
         for i in range(len(k values)):
             # svd = TruncatedSVD(n components=k)
             svd list.append(TruncatedSVD(n components=k values[i]))
             svd list[i].fit(X train tfidf) # X train tfidf
             explained variances.append(svd list[i].explained variance ratio .sum())
         plt.plot(k_values, explained_variances, marker='o')
         plt.xlabel('Number of Components (k)')
         plt.ylabel('Explained Variance Ratio')
         plt.title('Explained Variance Ratio for LSI')
         plt.show()
```



Questation 4

4a-answer.

Explained Variance Ratio Plot for LSI

The var ratio plot for LSI indicates the extend of how the total var in TF_IDF is retained as increasing number of k, and overall, as k increases, the explained var ratio also increases, which means that more information is captured.

Concavity in the context of dimensionality reduction implies a trade-off between the retained elements and the explained variance. Increasing the value of K initially leads to a significant rise in explained variance; however, as K further increases, the growth in explained variance gradually plateaus. Hence, while a larger K corresponds to a greater explained variance, the incremental gain diminishes with higher K values. Moreover, higher K values result in prolonged algorithm runtime. Enhanced explained variance signifies a preservation of more data as K increases, yet in the case of dimensionality reduction for categorized datasets, retaining more data is crucial for maintaining higher test accuracy.

• 4b

```
from sklearn.decomposition import NMF
from numpy.linalg import norm
from sklearn.metrics import mean_squared_error
```

```
k = 50

lsi_model = TruncatedSVD(n_components=k)
train_lsi = lsi_model.inverse_transform(lsi_model.fit_transform(X_train_tfidf))
# Lsi_mse = norm(np.subtract(X_train_tfidf.toarray(), train_lsi), 'fro')**2
lsi_mse = mean_squared_error(X_train_tfidf.toarray(), train_lsi)

nmf_model = NMF(n_components=k, init='random', max_iter=300)
W = nmf_model.fit_transform(X_train_tfidf)
H = nmf_model.components_
train_nmf = nmf_model.inverse_transform(W)
# nmf_mse = norm(np.subtract(X_train_tfidf.toarray(), train_nmf), 'fro')**2
nmf_mse = mean_squared_error(X_train_tfidf.toarray(), train_nmf)
print('LSI_MSE:', lsi_mse)
print('NMF_MSE:', nmf_mse)
```

LSI MSE: 5.317909141726312e-05 NMF MSE: 5.409425986967508e-05

4b-answer.

The NMF results in a higher residual MSE error, with $\|X-WH\|_{2F} = 5.409425986967508e-05$, compared to $\|X-Uk\Sigma kVTk\|_{2F} = 5.317909141726312e-05$. This increased error is attributed to the constraints imposed by NMF on matrices W and H, limiting them to non-negative values. In contrast, LSI does not enforce such restrictions. The truncated SVD used by LSI incorporates negative elements, providing a less constrained approximation of X. This allows for a more flexible representation in reduced dimensionality, contributing to the contrast in error rates between NMF and LSI.

4. Classification Algorithms

Question 5

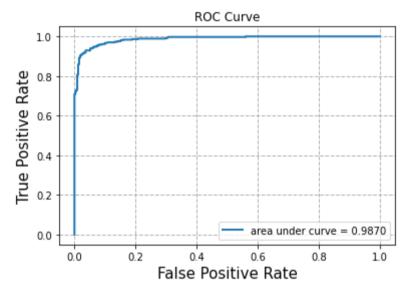
• 5a

```
from sklearn import svm
from sklearn import metrics
import seaborn as sn
from sklearn.preprocessing import LabelEncoder
from sklearn.decomposition import TruncatedSVD
```

```
# Apply LSI on data
svd_components = 50
truncated_svd = TruncatedSVD(n_components=svd_components, random_state=42)
train_truncated = truncated_svd.fit_transform(X_train_tfidf)
test_truncated = truncated_svd.transform(X_test_tfidf)
```

```
In [ ]:
         from sklearn.svm import SVC
         import matplotlib.pyplot as plt
         def train evaluate SVM(train data, train labels, test data, test labels, C value):
             # Train SVM
             svm model = SVC(kernel='linear', C=C value, random state=42)
             svm model.fit(train data, train labels)
             # PLot ROC curve
             prob estimates = svm model.decision_function(test_data)
             fpr, tpr, = metrics.roc curve(test labels, prob estimates)
             roc auc = metrics.auc(fpr, tpr)
             plt.plot(fpr, tpr, lw=2, label='area under curve = %0.4f' % roc auc)
             plt.grid(color='0.7', linestyle='--', linewidth=1)
             plt.title('ROC Curve')
             plt.xlabel('False Positive Rate', fontsize=15)
             plt.ylabel('True Positive Rate', fontsize=15)
             plt.legend(loc="lower right")
             plt.show()
             # Print SVM metrics
             print(f"C = {C value}")
             predictions = svm model.predict(test data)
             print('Accuracy:', metrics.accuracy_score(test_labels, predictions))
             print('Recall:', metrics.recall score(test labels, predictions))
             print('Precision:', metrics.precision score(test labels, predictions, zero division=1))
             print('F-1 Score:', metrics.f1 score(test labels, predictions))
             # Plot confusion matrix
             confusion matrix = metrics.confusion matrix(test labels, predictions)
             confusion display = metrics.ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=np.unique(test labels))
             confusion display.plot(cmap=plt.cm.Blues, xticks rotation='vertical')
             plt.title('Confusion Matrix')
```

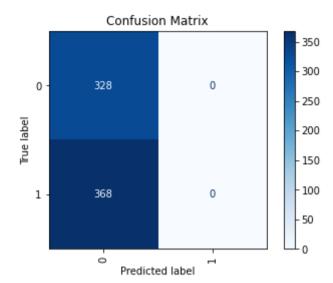
```
In []:
# Transform categories into numbers (Climate=0, Sports=1)
label_encoder = LabelEncoder()
train['binary_root_label'] = label_encoder.fit_transform(train['root_label'])
test['binary_root_label'] = label_encoder.fit_transform(test['root_label'])
categories = ['climate', 'sports']
train_evaluate_SVM(train_truncated, train['binary_root_label'], test_truncated, test['binary_root_label'], 0.0001)
```



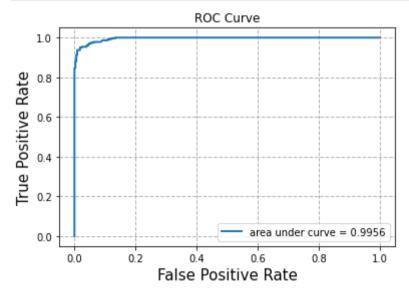
C = 0.0001

Accuracy: 0.47126436781609193

Recall: 0.0 Precision: 1.0 F-1 Score: 0.0

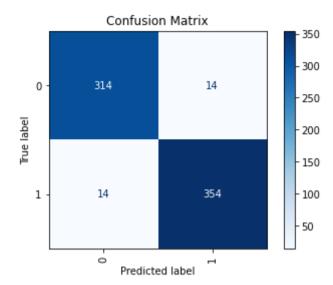


In []: train_evaluate_SVM(train_truncated, train['binary_root_label'], test_truncated, test['binary_root_label'], 1000)

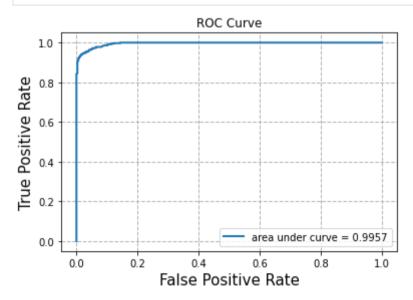


C = 1000

Accuracy: 0.9597701149425287 Recall: 0.9619565217391305 Precision: 0.9619565217391305 F-1 Score: 0.9619565217391305

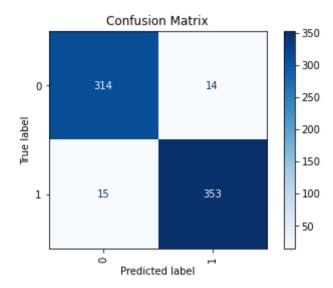


In []: train_evaluate_SVM(train_truncated, train['binary_root_label'], test_truncated, test['binary_root_label'], 100000)



C = 100000

Accuracy: 0.9583333333333334 Recall: 0.9592391304347826 Precision: 0.9618528610354223 F-1 Score: 0.9605442176870749



5a-answer:

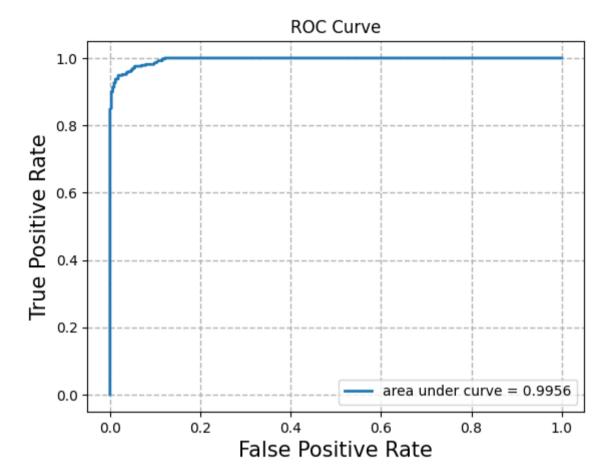
- The SVM with a γ value of 1000 (hard margin) showcased superior performance, demonstrating higher accuracy, F-1 score, precision, and ROC area under the curve in comparison to the soft margin SVM. Also when The SVM with a γ value of 100000, the performance this similiar to γ =1000, a little bit worse than γ =1000.
- Gamma acts as a hyperparameter that governs SVM error during the training process. A low γ value, like 0.0001, leads to minimal errors in class separation, resulting in overfitting to the training data. This overfitting translates to low accuracy when validating on testing data, as indicated by the confusion matrix, which exposes misclassifications. The model accurately identified sports articles but inaccurately categorized all climate articles as sports, revealing a flawed classification. This also shows that when gamma is too small, the model cannot well distinguish the differences between the two categories in the two-classification task. It will be easier to summarize one category into another, and it will be more difficult to identify the two categories with different feature.
- Although the ROC curve for the soft margin SVM implies high separability, this metric alone does not necessarily align with accurate classification. A model with strong separability may still exhibit poor performance in terms of classification accuracy, underscoring the importance of considering multiple metrics for a thorough model evaluation.

• 5b

```
In [ ]:
         from sklearn.model selection import GridSearchCV
         # Optimize SVM parameters with a linear kernel
         c values = [0.001, 0.01, 0.1, 1, 10, 100, 1000, 10000, 100000, 1000000]
         param grid = {'C': [0.001, 0.01, 0.1, 1, 10, 100, 1000, 10000, 100000, 1000000],
                       'kernel': ['linear']}
         svm model = SVC(random state=42)
         grid search cv = GridSearchCV(svm model, param grid, cv=5, refit=True, scoring='accuracy', n jobs=-1)
         grid search cv.fit(train truncated, train['binary root label'])
         predicted labels = grid search cv.predict(test truncated)
         # Display cross-validation accuracy scores for different C values
         for c in c values:
             print(f'C Parameter: {c}, Cross-Validation Accuracy: {grid search cv.cv results ["mean test score"][c values.index(c)]}')
         # Output the best C parameter
         print("\nOptimal C Parameter: ", grid search cv.best params ['C'])
         # Evaluate SVM performance using the best C parameter
         train evaluate SVM(train truncated, train['binary_root_label'], test_truncated, test['binary_root_label'], grid_search_cv.best_par
        C Parameter: 0.001, Cross-Validation Accuracy: 0.5028776978417266
        C Parameter: 0.01, Cross-Validation Accuracy: 0.7848920863309352
        C Parameter: 0.1, Cross-Validation Accuracy: 0.9384892086330936
        C Parameter: 1, Cross-Validation Accuracy: 0.9507194244604318
        C Parameter: 10, Cross-Validation Accuracy: 0.9546762589928057
        C Parameter: 100, Cross-Validation Accuracy: 0.9550359712230214
        C Parameter: 1000, Cross-Validation Accuracy: 0.9532374100719425
```

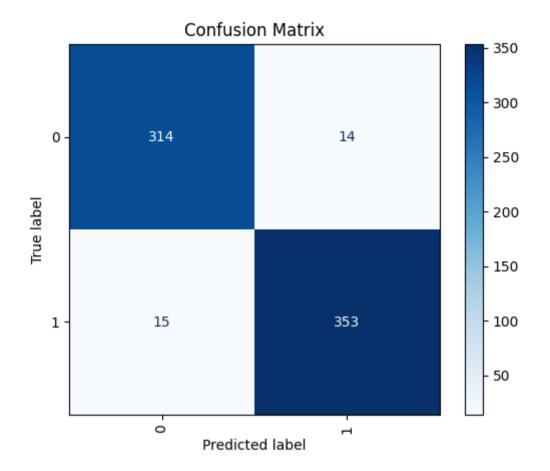
C Parameter: 10000, Cross-Validation Accuracy: 0.9532374100719425 C Parameter: 100000, Cross-Validation Accuracy: 0.9528776978417266 C Parameter: 1000000, Cross-Validation Accuracy: 0.9528776978417266

Optimal C Parameter: 100



C = 100

Accuracy: 0.9583333333333334 Recall: 0.9592391304347826 Precision: 0.9618528610354223 F-1 Score: 0.9605442176870749



5b-answer:

The best Gamma is 100

the result of this parameter is:

Accuracy: 0.9583333333333334

Recall: 0.9592391304347826

Precision: 0.9618528610354223

F-1 Score: 0.9605442176870749

Question 6

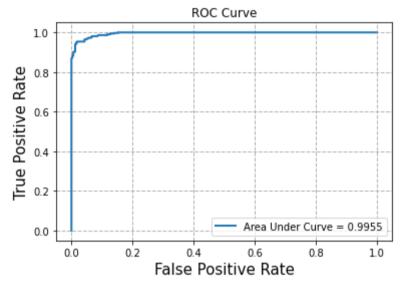
• 6a.

```
In [ ]:
         from sklearn.linear model import LogisticRegression
         from sklearn.model selection import GridSearchCV
         from sklearn import metrics
         from sklearn.metrics import ConfusionMatrixDisplay
         import matplotlib.pyplot as plt
         def plot roc curve(probabilities, test labels):
             fpr, tpr, = metrics.roc curve(test labels, probabilities)
             roc auc = metrics.auc(fpr, tpr)
             plt.plot(fpr, tpr, lw=2, label='Area Under Curve = %0.4f' % roc auc)
             plt.grid(color='0.7', linestyle='--', linewidth=1)
             plt.title('ROC Curve')
             plt.xlabel('False Positive Rate', fontsize=15)
             plt.ylabel('True Positive Rate', fontsize=15)
             plt.legend(loc="lower right")
             plt.show()
         def print classification metrics(predictions, test labels, label names):
             print('Accuracy: ', metrics.accuracy score(test labels, predictions))
             print('Recall: ', metrics.recall_score(test_labels, predictions))
             print('Precision: ', metrics.precision score(test labels, predictions))
             print('F-1 Score: ', metrics.f1 score(test labels, predictions))
         def plot_confusion_matrix(confusion_matrix, label_names):
             display = ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=label names)
             display.plot(cmap=plt.cm.Blues, xticks rotation='vertical')
             plt.title('Confusion Matrix')
         def train and evaluate logistic regression(regularization, regularization strength, train data, train labels, test data, test labe
             logistic model = LogisticRegression(penalty=regularization, C=regularization strength, random state=42, solver='saga', max ite
             logistic model.fit(train data, train labels)
             # PLot ROC curve
             probabilities = logistic_model.decision_function(test_data)
             plot roc curve(probabilities, test labels)
             # Print classification metrics
```

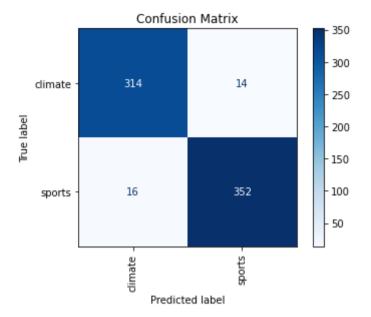
```
predictions = logistic_model.predict(test_data)
    print_classification_metrics(predictions, test_labels, label_names)

# Plot confusion matrix
    confusion_matrix = metrics.confusion_matrix(test_labels, predictions)
    plot_confusion_matrix(confusion_matrix, label_names)

# Train Logistic Regression with no regularization
train_and_evaluate_logistic_regression('none', 1.0, train_truncated, train['binary_root_label'], test_truncated, test['binary_root_label']
```



Accuracy: 0.9568965517241379
Recall: 0.9565217391304348
Precision: 0.9617486338797814
F-1 Score: 0.9591280653950953



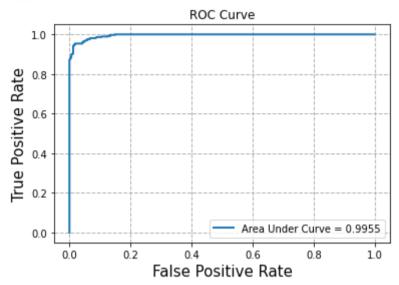
C: 10, Cross Validation accuracy scores: 0.9535971223021582
C: 100, Cross Validation accuracy scores: 0.9543165467625899

6b.1

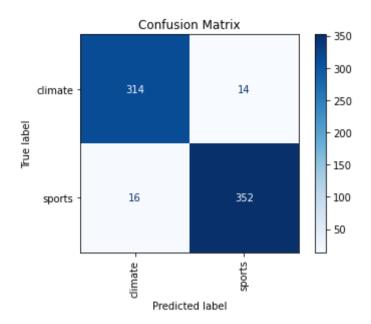
```
In [ ]:
        # Find best C value for L1 Regularization
        param grid = \{'C': [10**(-5), 0.0001, 0.001, 0.01, 1, 10, 100, 1000, 10000, 100000]\}
        grid 11 = GridSearchCV(LogisticRegression(penalty='11', random state=42, solver='saga', max iter=100000), param grid, cv=5, refit=
        grid l1.fit(train truncated, train['binary root label'])
        print("L1 Regularization Results")
        for i in range(len(c values)):
            print(f'C: {c values[i]}, Cross Validation accuracy scores: {grid l1.cv results ["mean test score"][i]}')
        print("\nBest C Value: ", grid l1.best params ['C'])
        train and evaluate logistic regression('l1', grid l1.best params ['C'], train truncated, train['binary root label'], test truncate
        L1 Regularization Results
       C: 1e-05, Cross Validation accuracy scores: 0.5014388489208633
       C: 0.0001, Cross Validation accuracy scores: 0.5014388489208633
       C: 0.001, Cross Validation accuracy scores: 0.5014388489208633
       C: 0.01, Cross Validation accuracy scores: 0.5014388489208633
       C: 0.1, Cross Validation accuracy scores: 0.9262589928057554
       C: 1, Cross Validation accuracy scores: 0.9496402877697842
```

C: 1000, Cross Validation accuracy scores: 0.9532374100719423
C: 10000, Cross Validation accuracy scores: 0.9532374100719423
C: 100000, Cross Validation accuracy scores: 0.9532374100719423

Best C Value: 100



Accuracy: 0.9568965517241379
Recall: 0.9565217391304348
Precision: 0.9617486338797814
F-1 Score: 0.9591280653950953



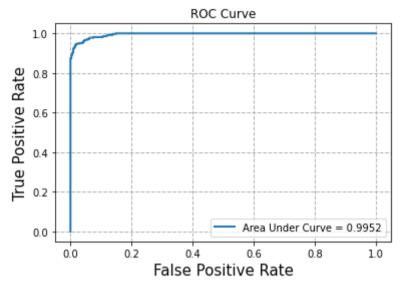
```
In []:
# Find best C value for L2 Regularization
grid_12 = GridSearchCV(LogisticRegression(penalty='12', random_state=42, solver='saga', max_iter=100000), param_grid, cv=5, refit=
grid_12.fit(train_truncated, train['binary_root_label'])

print("L2 Regularization Results")
for i in range(len(c_values)):
    print(f'C: {c_values[i]}, Cross Validation accuracy scores: {grid_12.cv_results_["mean_test_score"][i]}')

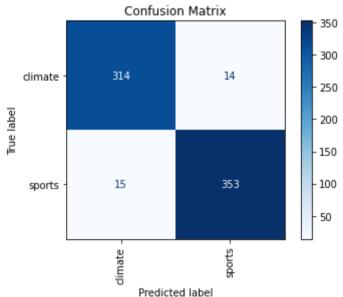
print("\nBest C Value: ", grid_12.best_params_['C'])
train_and_evaluate_logistic_regression('12', grid_12.best_params_['C'], train_truncated, train['binary_root_label'], test_truncate
```

```
L2 Regularization Results
C: 1e-05, Cross Validation accuracy scores: 0.5028776978417266
C: 0.0001, Cross Validation accuracy scores: 0.5028776978417266
C: 0.001, Cross Validation accuracy scores: 0.8201438848920862
C: 0.01, Cross Validation accuracy scores: 0.94136690647482
C: 0.1, Cross Validation accuracy scores: 0.94136690647482
C: 1, Cross Validation accuracy scores: 0.9496402877697842
C: 10, Cross Validation accuracy scores: 0.9546762589928057
C: 100, Cross Validation accuracy scores: 0.9553956834532373
C: 1000, Cross Validation accuracy scores: 0.9543165467625899
C: 10000, Cross Validation accuracy scores: 0.9532374100719423
C: 100000, Cross Validation accuracy scores: 0.9532374100719423
```

Best C Value: 100



Accuracy: 0.9583333333333334 Recall: 0.9592391304347826 Precision: 0.9618528610354223 F-1 Score: 0.9605442176870749



• Shown below are the best accuracy, recall, precision, and F-1 scores achieved by regression models with different regularizations. The best regularization strength for L1 and L2 regularization were found to be 100 and 100, respectively. Examination of the table reveals that the logistic regressions, optimized with their best C parameters, exhibited nearly identical performance metrics for No Regularization and L1. Moreover, both underperformd the L2 regularization by a slight margin.

Regularization Type	Accuracy	Recall	Precision	F-1 Score
No Regularization	0.9568965517241379	0.9565217391304348	0.9617486338797814	0.9591280653950953
L1	0.9568965517241379	0.9565217391304348	0.9617486338797814	0.9591280653950953
L2	0.95833333333333334	0.9592391304347826	0.9618528610354223	0.9605442176870749

6b.3

• How does the regularization parameter affect the test error?

Regularization is mainly used to prevent model overfitting. The use of regularization parameters is mainly to make the model less complex, thereby reducing the risk of overfitting and improving performance. At the same time, when the regularization parameters are too large, The model becomes too simple and cannot capture the key information in the data well, which may increase the test error.

• How are the learnt coefficients affected?

L1 regularization mainly produces a sparse model, in which many coefficients become 0, and the model will only use a part of the features, which means that such regularization helps in feature selection. L2 regularization can mainly solve the problem when the correlation of features is too high or the noise is large, it helps to control the complexity of the model, thereby improving the performance of the model.

• Why might one be interested in each type of regularization?

When the data has many irrelevant features, people can use L1 regularization to help simplify the model and feature selection. When the correlation of the data features is high or the nosie is too large,L2 regularization will help control the complexity of the model, thereby improving the performance of the model. People can apply different regularization methods according to different needs to improve the performance of the model. 6b.4

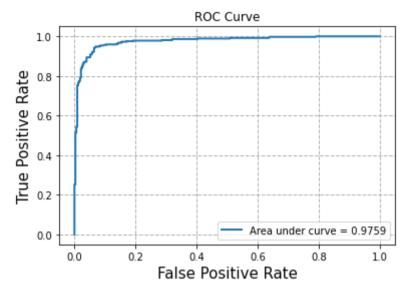
• The linear SVM utilizes a hyperplane as the decision boundary, aiming to maximize the distance between different classes in the data. Specifically, it seeks to find a linear decision boundary that optimally separates these classes. On the other hand, logistic regression adjusts the decision boundary to maximize the likelihood estimation. Unlike the SVM, logistic regression focuses on the probability of a data point belonging to a

specific class rather than the distance between classes and the boundary. Their performance diverges due to their distinct approaches in determining the decision boundary. Moreover, SVM, by emphasizing the separation between classes, exhibits lower susceptibility to overfitting compared to logistic regression. The performance contrast is further accentuated by the probabilistic nature of logistic regression and the more deterministic nature of SVM. While they often demonstrate similar performance in comparable tasks, the disparity becomes statistically significant when applied to markedly different tasks. SVM may be preferred for tasks like image classification or text recognition, whereas logistic regression might be more suitable for scenarios where class separation is challenging.

Question 7

```
In [ ]:
         from sklearn.naive baves import GaussianNB
         # Train Gaussian Naive Bayes classifier, print metrics, and plot ROC and confusion matrix
         def train gaussian nb(train data, train labels, test data, test labels, label names):
             gnb classifier = GaussianNB()
             gnb classifier.fit(train data, train labels)
             # PLot ROC curve
             probabilities = gnb classifier.predict_proba(test_data)
             fpr, tpr, _ = metrics.roc_curve(test_labels, probabilities[:, 1])
             roc auc = metrics.auc(fpr, tpr)
             plt.plot(fpr, tpr, lw=2, label='Area under curve = %0.4f' % roc auc)
             plt.grid(color='0.7', linestyle='--', linewidth=1)
             plt.title('ROC Curve')
             plt.xlabel('False Positive Rate', fontsize=15)
             plt.ylabel('True Positive Rate', fontsize=15)
             plt.legend(loc="lower right")
             plt.show()
             # Print classification metrics
             predictions = gnb_classifier.predict(test data)
             print('Accuracy: ', metrics.accuracy_score(test_labels, predictions))
             print('Recall: ', metrics.recall score(test labels, predictions))
             print('Precision: ', metrics.precision score(test labels, predictions))
             print('F-1 Score: ', metrics.f1 score(test labels, predictions))
             # Plot confusion matrix
             confusion matrix = metrics.confusion matrix(test labels, predictions)
             display = metrics.ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=label names)
             display.plot(cmap=plt.cm.Blues, xticks rotation='vertical')
             plt.title('Confusion Matrix')
```

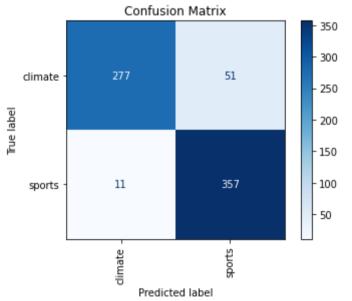
```
In [ ]: train_gaussian_nb(train_truncated, train['binary_root_label'], test_truncated, test['binary_root_label'], categories)
```



Accuracy: 0.9109195402298851 Recall: 0.970108695652174

Precision: 0.875

F-1 Score: 0.9201030927835052



Grid Search of Parameters

Question 8

```
In [ ]:
         from nltk.stem import PorterStemmer
         stemmer = PorterStemmer()
         def stem tokenize(text):
             return [stemmer.stem(token) for token in nltk.word_tokenize(text)]
In [ ]:
         import time
         from sklearn.pipeline import Pipeline
         full time = time.time()
         from shutil import rmtree
         pipe = Pipeline([
             ('vect', CountVectorizer()),
             ('tfidf', TfidfTransformer()),
             ('dim red', TruncatedSVD()),
             ('classifier', SVC())
         1)
         param_grid = [
             {
                 # feature extraction
                 'vect min df': [3, 5],
                 'vect__tokenizer': [None], # default lemmatization
                 'vect__analyzer': [stem_rmv_punc], # if Lemmatization
                 'vect__stop_words':['english'],
                 # dimensionality Reduction
                 'dim_red': [TruncatedSVD()],
                 'dim_red__n_components': [5, 30, 80],
                 # Classifier
                 'classifier': [LogisticRegression(penalty='l1', max_iter=100000, solver='saga',C = 100), LogisticRegression(penalty='l2',
                             SVC(C=100), GaussianNB()]
             },
                 # feature extraction
                 'vect min df': [3, 5],
                 'vect__tokenizer': [stem_tokenize], # if stemming
```

```
'vect analyzer': ['word'], # default analyzer
    'vect stop words':['english'],
    # dimensionality Reduction
    'dim red': [TruncatedSVD()].
    'dim red n components': [5, 30, 80],
    # Classifier
    'classifier': [LogisticRegression(penalty='l1', max iter=100000, solver='saga', C = 100), LogisticRegression(penalty='l2',
                SVC(C=100), GaussianNB()]
},
    # feature extraction
    'vect min df': [3, 5],
    'vect tokenizer': [None], # both None and stem tokenize
    'vect analyzer': [stem rmv punc], # both stem rmv punc and default analyzer
    'vect stop words': ['english'],
    # dimensionality reduction
    'dim red': [NMF(init='random', random state=0)],
    'dim_red__n_components': [5, 30, 80],
    'dim red init': ['random'], # NMF-specific parameters
    'dim red random state': [0],
    # Classifier
    'classifier': [LogisticRegression(penalty='l1', max iter=100000, solver='saga', C=100),
                   LogisticRegression(penalty='12', max iter=100000, solver='saga', C=100),
                   SVC(C=100), GaussianNB()]
},
# Grid for NMF
    # feature extraction
    'vect min df': [3, 5],
    'vect tokenizer': [stem tokenize],
    'vect analyzer': ['word'],
    'vect stop words': ['english'],
    # dimensionality reduction
    'dim_red': [NMF(init='random', random state=0)],
    'dim red n components': [5, 30, 80],
    'dim red init': ['random'], # NMF-specific parameters
    'dim red random state': [0],
    # Classifier
    'classifier': [LogisticRegression(penalty='l1', max iter=100000, solver='saga', C=100),
                   LogisticRegression(penalty='12', max_iter=100000, solver='saga', C=100),
                   SVC(C=100), GaussianNB()]
```

```
# GridSearchCV
          search = GridSearchCV(pipe,param_grid, n_jobs=8, cv=5,scoring='accuracy')
          search.fit(train['cleaned text'], train['binary root label'])
          print(f"--- Total time {time.time() - full time} seconds ---")
         --- Total time 1581.886403799057 seconds ---
In [ ]:
          results df = pd.DataFrame(search.cv results )
          top 5 = results df.sort values(by='mean test score', ascending=False).head(5)
In [ ]:
          with open('output.txt', 'w') as f:
              f.write(results df.to string())
In [ ]:
          top 5
Out[ ]:
             mean fit time std fit time mean score time std score time
                                                                             param classifier
                                                                                                           param dim red param dim red n components
         16
                 23.596305
                             0.254585
                                              5.920521
                                                            0.129643
                                                                                 SVC(C=100) TruncatedSVD(n components=80)
                                                                                                                                                   80
         17
                                                                                 SVC(C=100) TruncatedSVD(n components=80)
                 23.697774
                              0.128029
                                              5.863749
                                                            0.156670
                                                                                                                                                   80
         41
                 13.679450
                              0.230603
                                              3.456381
                                                            0.065684
                                                                                 SVC(C=100)
                                                                                                            TruncatedSVD()
                                                                                                                                                   80
                                                                     LogisticRegression(C=100, TruncatedSVD(n_components=80)
         11
                 23.833757
                              0.152688
                                              5.925853
                                                                                                                                                   80
                                                                        max iter=100000, sol...
                                                                                                         NMF(init='random',
         64
                 32.886108
                                              6.536150
                                                            0.268975
                                                                                 SVC(C=100)
                                                                                                                                                   80
                              1.981341
                                                                                                           random state=0)
        5 rows × 22 columns
```

```
In [ ]:
         top 5['params'].index
Out[]: Int64Index([16, 17, 41, 11, 64], dtvpe='int64')
In [ ]:
         for i in top 5['params'].index:
             print('-' * 20)
             print(top 5['params'][i])
        {'classifier': SVC(C=100), 'dim red': TruncatedSVD(n components=80), 'dim red n components': 80, 'vect analyzer': <function stem
        rmv punc at 0x000002A8C8FCFE18>, 'vect min df': 3, 'vect stop words': 'english', 'vect tokenizer': None}
        {'classifier': SVC(C=100), 'dim red': TruncatedSVD(n components=80), 'dim red n components': 80, 'vect analyzer': <function stem
        rmv punc at 0x000002A8C8FCFE18>, 'vect min df': 5, 'vect stop words': 'english', 'vect tokenizer': None}
        {'classifier': SVC(C=100), 'dim red': TruncatedSVD(), 'dim red n components': 80, 'vect analyzer': 'word', 'vect min df': 5, 've
        ct stop words': 'english', 'vect tokenizer': <function stem tokenize at 0x000002A991501D08>}
        {'classifier': LogisticRegression(C=100, max iter=100000, solver='saga'), 'dim red': TruncatedSVD(n components=80), 'dim red n com
        ponents': 80, 'vect analyzer': <function stem rmv punc at 0x000002A8C8FCFE18>, 'vect min df': 5, 'vect stop words': 'english',
        'vect tokenizer': None}
        {'classifier': SVC(C=100), 'dim red': NMF(init='random', random state=0), 'dim red init': 'random', 'dim red n components': 80,
        'dim red random state': 0, 'vect analyzer': <function stem rmv punc at 0x000002A8C8FCFE18>, 'vect min df': 3, 'vect stop word
        s': 'english', 'vect tokenizer': None}
In [ ]:
         for index, row in top 5.iterrows():
             # Setup the pipeline with the best parameters
             pipe.set params(**row['params'])
             # Fit the pipeline on the entire training set
             pipe.fit(train['cleaned text'], train['binary root label'])
             # Evaluate on the test set
             prediction bestpip = pipe.predict(test['cleaned_text'])
             print('-' * 20)
             print('Accuracy: ', metrics.accuracy score(test['binary root label'], prediction bestpip))
             print('Recall: ', metrics.recall score(test['binary root label'], prediction bestpip))
             print('Precision: ', metrics.precision score(test['binary root label'], prediction bestpip))
             print('F-1 Score: ', metrics.f1 score(test['binary root label'], prediction bestpip))
```

Accuracy: 0.9770114942528736 Recall: 0.9755434782608695 Precision: 0.9808743169398907 F-1 Score: 0.9782016348773842

Accuracy: 0.9626436781609196 Recall: 0.9592391304347826 Precision: 0.9697802197802198 F-1 Score: 0.9644808743169399

Accuracy: 0.9669540229885057 Recall: 0.9619565217391305 Precision: 0.9752066115702479 F-1 Score: 0.9685362517099864

Accuracy: 0.9741379310344828 Recall: 0.9646739130434783 Precision: 0.9861111111111112 F-1 Score: 0.9752747252747254

Q8-answer

Top 5 Combinations

NO.1: 'classifier': SVC(C=100), 'dim_red': TruncatedSVD(n_components=80)-LSI, 'dim_redn_components': 80, Lemmatization, 'vectmin_df': 3'

• **Accuracy:** 0.9770114942528736

• **Recall:** 0.9755434782608695

• Precision: 0.9808743169398907

• **F-1 Score:** 0.9782016348773842

No.2: 'classifier': SVC(C=100), 'dim_red': TruncatedSVD(n_components=80)-LSI, 'dim_redn_components': 80, Lemmatization, 'vectmin_df': 5

• **Accuracy:** 0.9683908045977011

• Recall: 0.9592391304347826

• **Precision:** 0.980555555555555

• **F-1 Score:** 0.9697802197802197

No.3: {'classifier': SVC(C=100), 'dim red': TruncatedSVD()-LSI, 'dim redn components': 80, Stemming, 'vectmin df': 5}

• **Accuracy:** 0.9626436781609196

• **Recall:** 0.9592391304347826

• **Precision:** 0.9697802197802198

• **F-1 Score:** 0.9644808743169399

No.4: {'classifier': LogisticRegression(penalty='l2',C=100, max_iter=100000, solver='saga'), 'dim_red': TruncatedSVD(n_components=80)-LSI, 'dim_redn_components': 80, Lemmatization, 'vectmin_df': 5}

• **Accuracy:** 0.9669540229885057

• **Recall:** 0.9619565217391305

• **Precision:** 0.9752066115702479

• **F-1 Score:** 0.9685362517099864

No.5:{'classifier': SVC(C=100), 'dim_red': NMF(init='random', random_state=0), 'dim_redn_components':80, Lemmatization, 'vectmin_df': 3}

• **Accuracy:** 0.9741379310344828

• **Recall:** 0.9646739130434783

• **Precision:** 0.9861111111111112

• **F-1 Score:** 0.9752747252747254

All result of Gridsearchcv shown in output.txt

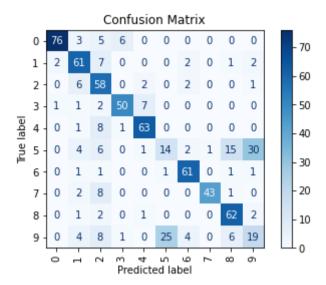
Multiclass Classification

Questation 9

```
X train tfidf mult = tfidf.fit transform(count vectorizer.fit transform(train mult['cleaned text']))
         X test tfidf mult = tfidf.transform(count vectorizer.transform(test mult['cleaned text']))
         map row to class = {
             0: "basketball", 1: "baseball", 2: "tennis",
             3: "football", 4: "soccer", 5: "forest fire",
             6: "flood", 7: "earthquake", 8: "drought", 9: "heatwave"
         class to num = {v: k for k, v in map row to class.items()}
         test mult['binary root label'] = [class to num[label] for label in test mult['leaf label']]
         train mult['binary root label'] = [class to num[label] for label in train mult['leaf label']]
         svd components = 80
         truncated svd mult = TruncatedSVD(n components=svd components, random state=42)
         train truncated mult = truncated svd.fit transform(X train tfidf mult)
         test truncated mult = truncated svd.transform(X test tfidf mult)
         # mult classifer
         from sklearn.naive_bayes import GaussianNB
         gnb classifier = GaussianNB()
         gnb classifier.fit(train truncated mult, train mult['binary root label'])
         predictions = gnb classifier.predict(test truncated mult)
In [ ]:
         confusion matrix = metrics.confusion matrix(test mult['binary root label'] , predictions)
         confusion display = metrics.ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=np.unique(test mult['binary r
         confusion display.plot(cmap=plt.cm.Blues, xticks rotation='vertical')
```

```
plt.title('Confusion Matrix')
```

Out[]: Text(0.5, 1.0, 'Confusion Matrix')



```
print('Accuracy: ', metrics.accuracy_score(test_mult['binary_root_label'], predictions))
print('Recall: ', metrics.recall_score(test_mult['binary_root_label'], predictions, average='macro'))
print('Precision: ', metrics.precision_score(test_mult['binary_root_label'], predictions, average='macro'))
print('F-1 Score: ', metrics.f1_score(test_mult['binary_root_label'], predictions, average='macro'))
```

Accuracy: 0.728448275862069 Recall: 0.7288709655591503 Precision: 0.7206829496758594 F-1 Score: 0.7163369334734864

```
print('Precision: ', metrics.precision score(test labels, predictions, average='weighted'))
   print('F-1 Score: ', metrics.f1 score(test labels, predictions, average='weighted'))
   # Plot confusion matrix
   confusion matrix = metrics.confusion matrix(test labels, predictions)
   confusion display = metrics.ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=np.unique(test labels))
   confusion display.plot(cmap=plt.cm.Blues, xticks rotation='vertical')
   plt.title('Confusion Matrix')
def train evaluate SVM ovr(train data, train labels, test data, test labels, C value):
    # Train SVM
   # clf ovr = OneVsRestClassifier(SVC(C=C value, class weight='balanced'))
   clf ovr = OneVsRestClassifier(SVC(C=C value))
   clf ovr.fit(train data, train labels)
   # Print SVM metrics
   print(f"C = {C value}")
   predictions = clf_ovr.predict(test_data)
   print('Accuracy: ', metrics.accuracy score(test labels, predictions))
   print('Recall: ', metrics.recall score(test labels, predictions, average='weighted'))
   print('Precision: ', metrics.precision score(test labels, predictions, average='weighted'))
   print('F-1 Score: ', metrics.f1_score(test_labels, predictions, average='weighted'))
   # Plot confusion matrix
   confusion matrix = metrics.confusion matrix(test labels, predictions)
   confusion display = metrics.ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=np.unique(test labels))
   confusion display.plot(cmap=plt.cm.Blues, xticks rotation='vertical')
   plt.title('Confusion Matrix')
```

```
def train_evaluate_SVM_ovo_solveci(train_data, train_labels, test_data, test_labels, C_value):
    # Train SVM
    clf_ovo = OneVsOneClassifier(SVC(C=C_value, class_weight='balanced'))
    clf_ovo.fit(train_data, train_labels)

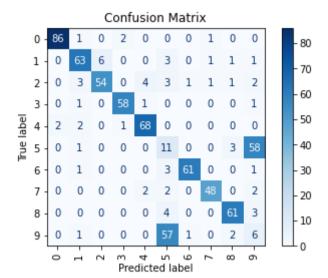
# Print SVM metrics
    print(f"C = {C_value}")
    predictions = clf_ovo.predict(test_data)
    print('Accuracy: ', metrics.accuracy_score(test_labels, predictions))
    print('Recall: ', metrics.recall_score(test_labels, predictions, average='weighted'))
    print('Precision: ', metrics.precision_score(test_labels, predictions, average='weighted'))
    print('F-1 Score: ', metrics.f1_score(test_labels, predictions, average='weighted'))

# Plot confusion matrix
```

```
confusion matrix = metrics.confusion matrix(test labels, predictions)
    confusion display = metrics.ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=np.unique(test labels))
    confusion display.plot(cmap=plt.cm.Blues, xticks rotation='vertical')
   plt.title('Confusion Matrix')
def train evaluate SVM ovr solveci(train data, train labels, test data, test labels, C value):
    clf ovr = OneVsRestClassifier(SVC(C=C value, class weight='balanced'))
    clf ovr.fit(train data, train labels)
    # Print SVM metrics
    print(f"C = {C value}")
   predictions = clf ovr.predict(test data)
   print('Accuracy: ', metrics.accuracy score(test labels, predictions))
   print('Recall: ', metrics.recall score(test labels, predictions, average='weighted'))
    print('Precision: ', metrics.precision score(test labels, predictions, average='weighted'))
    print('F-1 Score: ', metrics.f1 score(test labels, predictions, average='weighted'))
    # Plot confusion matrix
    confusion matrix = metrics.confusion matrix(test labels, predictions)
    confusion display = metrics.ConfusionMatrixDisplay(confusion matrix=confusion matrix, display labels=np.unique(test labels))
    confusion_display.plot(cmap=plt.cm.Blues, xticks_rotation='vertical')
   plt.title('Confusion Matrix')
```

result of one vs one train_evaluate_SVM_ovo_solveci(train_truncated_mult, train_mult['binary_root_label'], test_truncated_mult, test_mult['binary_root_

C = 100
Accuracy: 0.7413793103448276
Recall: 0.7413793103448276
Precision: 0.7612107213543277
F-1 Score: 0.7506511481390324



In []: #

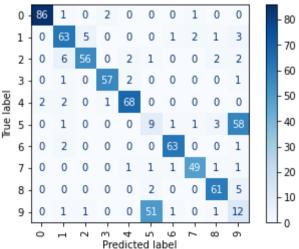
result of one vs rest

train_evaluate_SVM_ovr_solveci(train_truncated_mult, train_mult['binary_root_label'], test_truncated_mult, test_mult['binary_root_

C = 100

Accuracy: 0.7528735632183908 Recall: 0.7528735632183908 Precision: 0.7609817618889433 F-1 Score: 0.7563951590157423

Confusion Matrix



The result that after merge 9 to 5

```
In [ ]:
         # Solve class imbalance issue
         train mult['changed lable'] = 0
         test mult['changed lable'] = 0
         train mult['changed lable'] = train mult['binary root label'].replace(9, 5)
         test_mult['changed_lable'] = test_mult['binary_root_label'].replace(9, 5)
In [ ]:
         # result of one vs one(solved class imbalance issue)
         train evaluate SVM ovo(train truncated mult, train mult['changed lable'], test truncated mult, test mult['changed lable'], 100)
         C = 100
        Accuracy: 0.9051724137931034
         Recall: 0.9051724137931034
         Precision: 0.9066101760370868
        F-1 Score: 0.9049299901511274
                     Confusion Matrix
                                                - 120
                                                - 100
                                                - 80
                                                 60
                                                 40
                                                 - 20
                       Predicted label
In [ ]:
         # result of one vs rest(solved class imbalance issue)
         train_evaluate_SVM_ovr(train_truncated_mult, train_mult['changed_lable'], test_truncated_mult, test_mult['changed_lable'], 100)
         C = 100
```

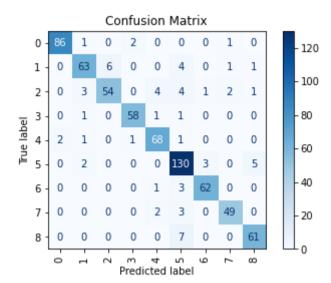
Accuracy: 0.9137931034482759 Recall: 0.9137931034482759 Precision: 0.9149731148680501 F-1 Score: 0.9138572946484304 Confusion Matrix 2 0 1 0 0 - 120 1 -0 - 100 0 6 57 0 - 80 60 40 - 20 Predicted label

The result that after merge 9 to 5-solve imbalance

In []: train_evaluate_SVM_ovo_solveci(train_truncated_mult, train_mult['changed_lable'], test_truncated_mult, test_mult['changed_lable'],

C = 100

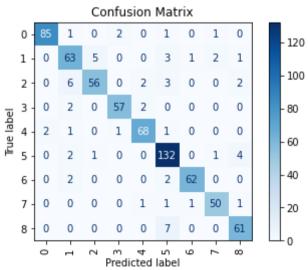
Accuracy: 0.9066091954022989
Recall: 0.9066091954022989
Precision: 0.907756332429278
F-1 Score: 0.9062940513107396



In []: train_evaluate_SVM_ovr_solveci(train_truncated_mult, train_mult['changed_lable'], test_truncated_mult, test_mult['changed_lable'],

C = 100

Accuracy: 0.9109195402298851 Recall: 0.9109195402298851 Precision: 0.9120986975869851 F-1 Score: 0.9109655717406929



Q9-answer:

1 How to resolve class imbalance?

The imbalance issue can be addressed using classs_weight='balanced' parameter in initiation the One vs rest SVM classier. The assigned weight inversely proportional to class frequency, and address the issue.

2 Observation on confusion matrix:

Yes, there are visible blocks along major diagonal of the matrix. The blocks is indicating instances that classier made right predictions for each class. A strong diagonal means that classier is doing well. But for label 9 and label 5, the squares here are not very obvious. On the contrary, it can be clearly seen in the confusion matrix. Whether it is Na ive Bayes classification or SVM classification, the model will incorrectly classify label 5 into label 9, or label 9 is incorrectly classified as label 5, which means that the two labels are very similar.

3 suggest a subset of label:

I merge label 9(heatwave) to label 5(forest fire), and re computer the result, the result is better than the previous one, Accuracy, Recall, Precison adn F-1 Score is increased about 0.17. result is shown above.

4 class imbalance impact performance?

Yes, the class imbalance is impacting the performance after merging class. For example, when heatwave and forest fire into Natural disaster, the addressed class imbalance by using different classes weight during the training process, and then the result is that accuracy is changed.

Word Embedding

Q10

Question 10 text answer:

10a: GLOVE is transined on ratio to capture the information about relations among words. Ratio is helpful to emphasis the inportance of word co-occurance, and this mades the result vector to have more information than the probablity itself.

10b: Certainly, they would produce identical embedding vectors for both scenarios. Although GLOVE is trained by considering the global context of each word, during inference, it generates embeddings by solely focusing on the individual tokens.

10c: The value of the difference between woman vs man, and wife vs hisband will be similar as they are different in gender, but the wife vs orange will be very different as they are totally different words.

• ||GLoVE["woman"] - GLoVE["man"]||2 ≈ ||GLoVE["wife"] - GLoVE["husband"]||2 < ||GLoVE["wife"] - GLoVE["orange"]||2

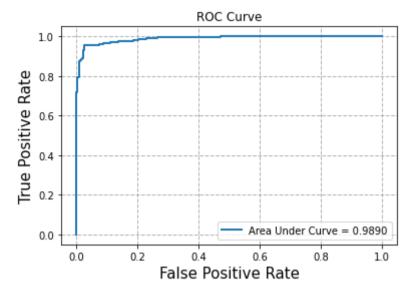
10d: GLOVE is trained using the full words, thus it is better to use lemmatization because this can retain the base form of words, and more suitable for GLOVE.

Q11

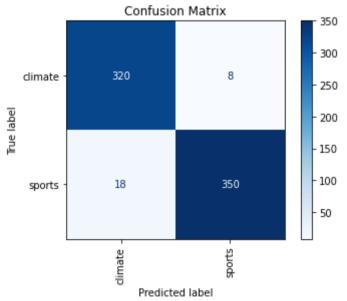
```
In [ ]:
         embeddings dict = {}
         dimension of glove = 300
         with open("glove.6B/glove.6B.300d.txt", 'rb') as f: # if 'r' fails with unicode error, please use 'rb'
             for line in f:
                 values = line.split()
                 word = values[0]
                 vector = np.asarray(values[1:], "float32")
                 embeddings dict[word] = vector
In [ ]:
         from sklearn.preprocessing import LabelEncoder
         import numpy as np
         import random
         import nltk
         from nltk import word tokenize
         import ast
         # Seed random generator
         np.random.seed(42)
         random.seed(42)
         # Split data into testing and training and clean it
         train set, test set = train test split(df[["root label", "keywords", "full text"]], test size=0.2)
         train_set['full_text'] = train_set['full_text'].apply(clean)
         test_set['full_text'] = test_set['full_text'].apply(clean)
         # Transform categories into number (Climate=0, Sports=1)
         label encoder = LabelEncoder()
         train set['binary root label'] = label encoder.fit transform(train set['root label'])
```

```
test set['binary root label'] = label encoder.fit transform(test set['root label'])
# Do the feature engineering process
def key to vec(keyword, embeddings dict):
    average embeddings = []
   for i in keyword.index:
        keyword embeddings = []
       for word in re.sub(r'[^A-Za-z0-9]+', ' ', keyword[i]).split():
            byte word = word.encode('utf-8')
            if byte word in embeddings dict:
                keyword embeddings.append(embeddings dict[byte word])
        if keyword embeddings:
            average embedding = np.mean(keyword_embeddings, axis=0)
        else:
            average embedding = np.zeros(300)
        average embeddings.append(average embedding)
    return average_embeddings
train set['glove data'] = 0
train set['glove data'] = 0
train set['glove data'] = key to vec(train set['keywords'],embeddings dict)
test set['glove data'] = key to vec(test set['keywords'],embeddings dict)
X = np.vstack(train set['glove data'].values)
y = train set['binary root label'].values
X_test = np.vstack(test_set['glove_data'].values)
y_test = test_set['binary_root_label'].values
```

```
In [ ]: train_and_evaluate_logistic_regression('l1', 100, X, y, X_test, y_test, categories)
```



Accuracy: 0.9626436781609196 Recall: 0.9510869565217391 Precision: 0.9776536312849162 F-1 Score: 0.9641873278236915



Question 11 text answer

A feature engineering process will be needed, and it will need to be firstly tockenized to make each documents into indivudual tockens. Then, words embedding loopup will be needed to look up the corresponding GLOVE ords vector for each document After that, normalization will be needed to normalize each word vector to ensure consistent scale and mitigate the impact of different document length. After that aggregation is needed to aggregate the normalized words vectors to form a single vector representation for the whole document. The aggregation can be done using averaging the vectors. To aviod the curse of dimensionalty, dimensionality reduction can also be needed. Finally the resulting vector will be the final representation of the document using GLOVE.

Q12

```
import numpy as np

def load_glove_model(file_path):
    embeddings_dict = {}
    with open(file_path, 'rb') as f: # if 'r' fails with unicode error, please use 'rb'
    for line in f:
        values = line.split()
        word = values[0]
        vector = np.asarray(values[1:], "float32")
        embeddings_dict[word] = vector
    return embeddings_dict

file_prefix = "glove.6B.7"
files = ['glove.6B.50d.txt', 'glove.6B.100d.txt', 'glove.6B.200d.txt', 'glove.6B.300d.txt']

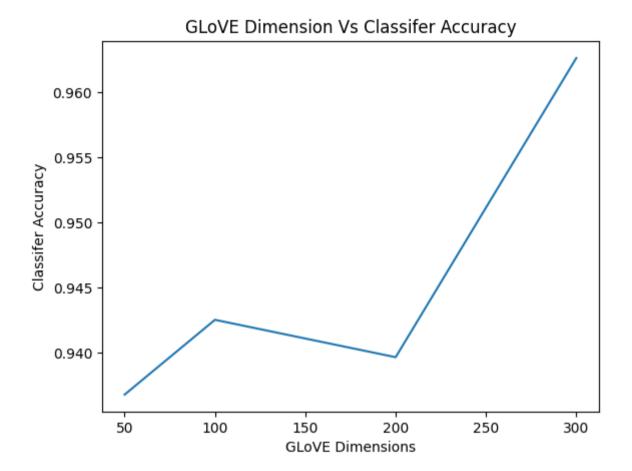
embeddings_dicts = [load_glove_model(file_prefix + file_name) for file_name in files]
```

```
In [ ]:
    dimension_list = [50, 100, 200, 300]
    X_train = {}
    X_test = {}
    y = train_set['binary_root_label'].values
    y_test = test_set['binary_root_label'].values

    for i, dim in enumerate(dimension_list):
        glove_data_key = f'glove_data_{dim}'
        train_set[glove_data_key] = key_to_vec(train_set['keywords'], embeddings_dicts[i])
```

```
test set[glove data key] = key to vec(test set['keywords'], embeddings dicts[i])
            X_train[dim] = np.vstack(train_set[glove_data_key].values)
            X_test[dim] = np.vstack(test_set[glove_data_key].values)
         accuracies = []
         Log = LogisticRegression(penalty='l1', C=1000, solver='saga', max iter=100000)
         for dim in dimension_list:
             Log.fit(X train[dim], y)
             predictions = Log.predict(X test[dim])
             accuracy = metrics.accuracy score(y test, predictions)
             accuracies.append(accuracy)
             print(f"Accuracy for {dim} dimensions: {accuracy}")
        Accuracy for 50 dimensions: 0.9367816091954023
        Accuracy for 100 dimensions: 0.9425287356321839
        Accuracy for 200 dimensions: 0.9396551724137931
        Accuracy for 300 dimensions: 0.9626436781609196
In [ ]:
         plt.plot(dimension_list,accuracies)
         plt.xlabel('GLoVE Dimensions')
         plt.ylabel('Classifer Accuracy')
         plt.title('GLoVE Dimension Vs Classifer Accuracy')
```

Out[]: Text(0.5, 1.0, 'GLoVE Dimension Vs Classifer Accuracy')



Q12 text answer:

The trend is expected.

The plot shows when dimension increases, the accuracy overall increases, although this is not always the case, but the overall trend still shows this. This tells us that higher dimension representation allows better preservation of the semantic relational information and contact, producing better classification accuracy.

Q13

```
In [ ]:
         !pip install umap-learn
         !pip install umap-learn[plot]
        Collecting umap-learn
          Downloading umap-learn-0.5.5.tar.gz (90 kB)
        Requirement already satisfied: numpy>=1.17 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn) (1.19.5)
        Requirement already satisfied: scipy>=1.3.1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn) (1.5.4)
        Requirement already satisfied: scikit-learn>=0.22 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn) (0.24.
        Collecting numba>=0.51.2
          Downloading numba-0.53.1-cp36-cp36m-win amd64.whl (2.3 MB)
        Collecting pynndescent>=0.5
          Downloading pynndescent-0.5.11-py3-none-any.whl (55 kB)
        Requirement already satisfied: tddm in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn) (4.64.1)
        Requirement already satisfied: setuptools in c:\users\michael mi\.conda\envs\219\lib\site-packages (from numba>=0.51.2->umap-learn)
        (58.0.4)
        Collecting llvmlite<0.37.>=0.36.0rc1
          Downloading llvmlite-0.36.0-cp36-cp36m-win amd64.whl (16.0 MB)
        Requirement already satisfied: importlib-metadata>=4.8.1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from pynndescent
        >=0.5- umap-learn) (4.8.3)
        Requirement already satisfied: joblib>=0.11 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from pynndescent>=0.5->umap-1
        earn) (1.1.1)
        Requirement already satisfied: typing-extensions>=3.6.4 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from importlib-me
        tadata>=4.8.1->pynndescent>=0.5->umap-learn) (4.1.1)
        Requirement already satisfied: zipp>=0.5 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from importlib-metadata>=4.8.1->
        pvnndescent>=0.5->umap-learn) (3.6.0)
        Requirement already satisfied: threadpoolctl>=2.0.0 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from scikit-learn>=0.
        22->umap-learn) (3.1.0)
        Requirement already satisfied: colorama in c:\users\michael mi\.conda\envs\219\lib\site-packages (from tqdm->umap-learn) (0.4.5)
        Requirement already satisfied: importlib-resources in c:\users\michael mi\.conda\envs\219\lib\site-packages (from tqdm->umap-learn)
        (5.4.0)
        Building wheels for collected packages: umap-learn
          Building wheel for umap-learn (setup.py): started
          Building wheel for umap-learn (setup.py): finished with status 'done'
          Created wheel for umap-learn: filename=umap learn-0.5.5-pv3-none-anv.whl size=86853 sha256=489f770a3f0aff6dd69017de6bf56da23d4e4c
        4bb6f9e0f0392f2279e3de610f
          Stored in directory: c:\users\michael mi\appdata\local\pip\cache\wheels\fa\bf\ee\50cca124d5a1e2f411b47ff63c6dd5c0a152d9622e4c9944
        Successfully built umap-learn
        Installing collected packages: llvmlite, numba, pynndescent, umap-learn
        Successfully installed llvmlite-0.36.0 numba-0.53.1 pynndescent-0.5.11 umap-learn-0.5.5
        Requirement already satisfied: umap-learn[plot] in c:\users\michael mi\.conda\envs\219\lib\site-packages (0.5.5)
        Requirement already satisfied: tqdm in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot]) (4.64.1)
        Requirement already satisfied: pynndescent>=0.5 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot])
        (0.5.11)
        Requirement already satisfied: numpy>=1.17 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot]) (1.19.
```

```
Requirement already satisfied: scikit-learn>=0.22 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot])
(0.24.2)
Requirement already satisfied: scipy>=1.3.1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot]) (1.5.
Requirement already satisfied: numba>=0.51.2 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot]) (0.5
3.1)
Collecting colorcet
  Downloading colorcet-3.0.1-pv2.pv3-none-anv.whl (1.7 MB)
Collecting holoviews
  Downloading holoviews-1.14.9-py2.py3-none-any.whl (4.3 MB)
Requirement already satisfied: matplotlib in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot]) (3.3.4)
Requirement already satisfied: pandas in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot]) (1.1.5)
Requirement already satisfied: seaborn in c:\users\michael mi\.conda\envs\219\lib\site-packages (from umap-learn[plot]) (0.11.2)
Collecting datashader
  Downloading datashader-0.13.0-pv2.pv3-none-anv.whl (15.8 MB)
Collecting bokeh
  Downloading bokeh-2.3.3.tar.gz (10.7 MB)
Collecting scikit-image
  Downloading scikit image-0.17.2-cp36-cp36m-win amd64.whl (11.5 MB)
Requirement already satisfied: llvmlite<0.37,>=0.36.0rc1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from numba>=0.5
1.2->umap-learn[plot]) (0.36.0)
Requirement already satisfied: setuptools in c:\users\michael mi\.conda\envs\219\lib\site-packages (from numba>=0.51.2->umap-learn
[plot]) (58.0.4)
Requirement already satisfied: joblib>=0.11 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from pynndescent>=0.5->umap-l
earn[plot]) (1.1.1)
Requirement already satisfied: importlib-metadata>=4.8.1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from pynndescent
\geq =0.5- \text{vmap-learn[plot]}) (4.8.3)
Requirement already satisfied: typing-extensions>=3.6.4 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from importlib-me
tadata>=4.8.1->pynndescent>=0.5->umap-learn[plot]) (4.1.1)
Requirement already satisfied: zipp>=0.5 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from importlib-metadata>=4.8.1->
pynndescent>=0.5->umap-learn[plot]) (3.6.0)
Requirement already satisfied: threadpoolctl>=2.0.0 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from scikit-learn>=0.
22->umap-learn[plot]) (3.1.0)
Collecting PyYAML>=3.10
  Downloading PyYAML-6.0.1-cp36-cp36m-win amd64.whl (153 kB)
Requirement already satisfied: python-dateutil>=2.1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from bokeh->umap-lear
n[plot]) (2.8.2)
Requirement already satisfied: Jinja2>=2.9 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from bokeh->umap-learn[plot])
(3.0.3)
Requirement already satisfied: pillow>=7.1.0 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from bokeh->umap-learn[plo
t]) (8.4.0)
Requirement already satisfied: packaging>=16.8 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from bokeh->umap-learn[plo
Requirement already satisfied: tornado>=5.1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from bokeh->umap-learn[plot])
(6.1)
Requirement already satisfied: MarkupSafe>=2.0 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from Jinja2>=2.9->bokeh->u
```

```
map-learn[plot]) (2.0.1)
Requirement already satisfied: pyparsing!=3.0.5,>=2.0.2 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from packaging>=1
6.8->bokeh->umap-learn[plot]) (3.1.1)
Requirement already satisfied: six>=1.5 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from python-dateutil>=2.1->bokeh-
>umap-learn[plot]) (1.16.0)
Collecting pyct>=0.4.4
  Downloading pvct-0.4.8-pv2.pv3-none-anv.whl (15 kB)
Collecting param>=1.7.0
  Downloading param-1.13.0-pv2.pv3-none-anv.whl (87 kB)
Collecting dask[complete]>=0.18.0
  Downloading dask-2021.3.0-py3-none-any.whl (925 kB)
Collecting datashape>=0.5.1
  Downloading datashape-0.5.2.tar.gz (76 kB)
Collecting xarray>=0.9.6
  Downloading xarray-0.16.2-py3-none-any.whl (736 kB)
Collecting fsspec>=0.6.0
  Downloading fsspec-2022.1.0-py3-none-any.whl (133 kB)
Collecting distributed>=2021.03.0
  Downloading distributed-2021.3.0-pv3-none-anv.whl (675 kB)
Collecting cloudpickle>=0.2.2
  Downloading cloudpickle-2.2.1-py3-none-any.whl (25 kB)
Collecting toolz>=0.8.2
  Downloading toolz-0.12.0-py3-none-any.whl (55 kB)
Collecting partd>=0.3.10
  Downloading partd-1.2.0-py3-none-any.whl (19 kB)
Collecting multipledispatch>=0.4.7
  Downloading multipledispatch-1.0.0-py3-none-any.whl (12 kB)
Collecting msgpack>=0.6.0
  Downloading msgpack-1.0.5-cp36-cp36m-win amd64.whl (69 kB)
Collecting zict>=0.1.3
  Downloading zict-2.1.0-py3-none-any.whl (11 kB)
Collecting psutil>=5.0
  Downloading psutil-5.9.8-cp36-cp36m-win amd64.whl (258 kB)
Collecting sortedcontainers!=2.0.0,!=2.0.1
  Downloading sortedcontainers-2.4.0-py2.py3-none-any.whl (29 kB)
Collecting tblib>=1.6.0
  Downloading tblib-1.7.0-py2.py3-none-any.whl (12 kB)
Collecting contextvars
  Downloading contextvars-2.4.tar.gz (9.6 kB)
Requirement already satisfied: click>=6.6 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from distributed>=2021.03.0->da
sk[complete]>=0.18.0->datashader->umap-learn[plot]) (8.0.4)
Requirement already satisfied: colorama in c:\users\michael mi\.conda\envs\219\lib\site-packages (from click>=6.6->distributed>=202
1.03.0->dask[complete]>=0.18.0->datashader->umap-learn[plot]) (0.4.5)
Requirement already satisfied: pytz>=2017.2 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from pandas->umap-learn[plo
t]) (2023.3.post1)
Collecting locket
  Downloading locket-1.0.0-py2.py3-none-any.whl (4.4 kB)
```

```
Collecting heapdict
  Downloading HeapDict-1.0.1-py3-none-any.whl (3.9 kB)
Collecting immutables>=0.9
  Downloading immutables-0.19-cp36-cp36m-win amd64.whl (58 kB)
Collecting panel>=0.8.0
  Downloading panel-0.13.1-py2.py3-none-any.whl (15.7 MB)
Collecting pvviz-comms>=0.7.4
  Downloading pyviz comms-2.2.1-py2.py3-none-any.whl (42 kB)
Collecting requests
  Downloading requests-2.27.1-py2.py3-none-any.whl (63 kB)
Collecting panel>=0.8.0
  Downloading panel-0.13.0-py2.py3-none-any.whl (15.6 MB)
  Downloading panel-0.12.7-py2.py3-none-any.whl (12.9 MB)
  Downloading panel-0.12.6-py2.py3-none-any.whl (12.9 MB)
  Downloading panel-0.12.5-py2.py3-none-any.whl (12.9 MB)
 Downloading panel-0.12.4-py2.py3-none-any.whl (12.9 MB)
 Downloading panel-0.12.3-py2.py3-none-any.whl (12.8 MB)
  Downloading panel-0.12.2-py2.py3-none-any.whl (12.8 MB)
  Downloading panel-0.12.1-py2.py3-none-any.whl (12.8 MB)
Requirement already satisfied: bleach in c:\users\michael mi\.conda\envs\219\lib\site-packages (from panel>=0.8.0->holoviews->umap-
learn[plot]) (4.1.0)
Collecting markdown
  Downloading Markdown-3.3.7-py3-none-any.whl (97 kB)
Requirement already satisfied: importlib-resources in c:\users\michael mi\.conda\envs\219\lib\site-packages (from tqdm->umap-learn
[plot]) (5.4.0)
Requirement already satisfied: webencodings in c:\users\michael mi\.conda\envs\219\lib\site-packages (from bleach->panel>=0.8.0->ho
loviews->umap-learn[plot]) (0.5.1)
Requirement already satisfied: kiwisolver>=1.0.1 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from matplotlib->umap-le
arn[plot]) (1.3.1)
Requirement already satisfied: cycler>=0.10 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from matplotlib->umap-learn[p
lot]) (0.11.0)
Collecting urllib3<1.27,>=1.21.1
  Downloading urllib3-1.26.18-py2.py3-none-any.whl (143 kB)
Requirement already satisfied: certifi>=2017.4.17 in c:\users\michael mi\.conda\envs\219\lib\site-packages (from requests->panel>=
0.8.0->holoviews->umap-learn[plot]) (2021.5.30)
Collecting idna<4,>=2.5
  Downloading idna-3.6-py3-none-any.whl (61 kB)
Collecting charset-normalizer~=2.0.0
  Downloading charset normalizer-2.0.12-py3-none-any.whl (39 kB)
Collecting networkx>=2.0
  Downloading networkx-2.5.1-py3-none-any.whl (1.6 MB)
Collecting tifffile>=2019.7.26
  Downloading tifffile-2020.9.3-py3-none-any.whl (148 kB)
Collecting PyWavelets>=1.1.1
  Downloading PyWavelets-1.1.1-cp36-cp36m-win amd64.whl (4.2 MB)
Collecting imageio>=2.3.0
  Downloading imageio-2.15.0-py3-none-any.whl (3.3 MB)
```

```
Collecting decorator<5.>=4.3
          Downloading decorator-4.4.2-py2.py3-none-any.whl (9.2 kB)
        Building wheels for collected packages: bokeh, datashape, contextvars
          Building wheel for bokeh (setup.pv): started
          Building wheel for bokeh (setup.py): finished with status 'done'
          Created wheel for bokeh: filename=bokeh-2.3.3-pv3-none-anv.whl size=11342794 sha256=84afc0c4232244aa18b268d62dde7521f0b17d4cf8a2a
        a006b23e7043b78a1fd
          Stored in directory: c:\users\michael mi\appdata\local\pip\cache\wheels\8b\59\97\257265b741bab184e0cc8f5676309cb1fe6fbda22011bbb3
          Building wheel for datashape (setup.pv): started
          Building wheel for datashape (setup.py): finished with status 'done'
          Created wheel for datashape: filename=datashape-0.5.2-py3-none-any.whl size=59453 sha256=143467c07b323a95e4ef548b8f008268c45042a7
        a89ae0aec7aba512d2524ada
          Stored in directory: c:\users\michael mi\appdata\local\pip\cache\wheels\81\61\fc\7d268954f6907b2a547c7895012769cde53af58f1aaf95a5
        4c
          Building wheel for contextvars (setup.py): started
          Building wheel for contextvars (setup.py): finished with status 'done'
          Created wheel for contextvars: filename=contextvars-2.4-py3-none-any.whl size=7681 sha256=fed357c9403f3bb6aef6e8673bb151bf01356ec
        b89b5228e3621aef5df5c2708
          Stored in directory: c:\users\michael_mi\appdata\local\pip\cache\wheels\41\11\53\911724983aa48deb94792432e14e518447212dd6c5477d49
        d3
        Successfully built bokeh datashape contextvars
        Installing collected packages: PyYAML, immutables, heapdict, zict, urllib3, toolz, tblib, sortedcontainers, psutil, param, msgpack,
        locket, idna, dask, contextvars, cloudpickle, charset-normalizer, requests, pyviz-comms, pyct, partd, multipledispatch, markdown, f
        sspec, distributed, decorator, bokeh, xarray, tifffile, PyWavelets, panel, networkx, imageio, datashape, colorcet, scikit-image, ho
        loviews, datashader
          Attempting uninstall: decorator
            Found existing installation: decorator 5.1.1
            Uninstalling decorator-5.1.1:
              Successfully uninstalled decorator-5.1.1
        Successfully installed PyWavelets-1.1.1 PyYAML-6.0.1 bokeh-2.3.3 charset-normalizer-2.0.12 cloudpickle-2.2.1 colorcet-3.0.1 context
        vars-2.4 dask-2021.3.0 datashader-0.13.0 datashape-0.5.2 decorator-4.4.2 distributed-2021.3.0 fsspec-2022.1.0 heapdict-1.0.1 holovi
        ews-1.14.9 idna-3.6 imageio-2.15.0 immutables-0.19 locket-1.0.0 markdown-3.3.7 msgpack-1.0.5 multipledispatch-1.0.0 networkx-2.5.1
        panel-0.12.1 param-1.13.0 partd-1.2.0 psutil-5.9.8 pyct-0.4.8 pyviz-comms-2.2.1 requests-2.27.1 scikit-image-0.17.2 sortedcontainer
        s-2.4.0 tblib-1.7.0 tifffile-2020.9.3 toolz-0.12.0 urllib3-1.26.18 xarray-0.16.2 zict-2.1.0
In [ ]:
         import umap.umap_ as umap
         import umap.plot
         import seaborn as sns
         umap model = umap.UMAP(random state=42)
         labels = train['binary root label']
         def apply umap projection(data, labels):
```

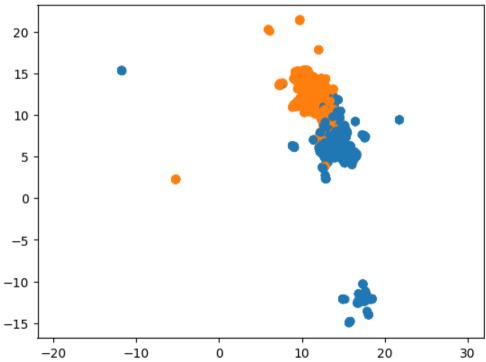
umap_model = umap.UMAP(random_state=42)
umap result = umap model.fit transform(data)

```
plt.scatter(
    umap_result[:, 0],
    umap_result[:, 1],
    c=[sns.color_palette()[x] for x in labels])
plt.gca().set_aspect('equal', 'datalim')
plt.title('UMAP Projection of the Normalized GLoVE-based Embeddings', fontsize=24)
plt.show()

# Apply UMAP on the GLoVE training data (300 dimensions)
apply_umap_projection(X_train[300], train['binary_root_label'])
```

c:\Users\Michael_Mi\.conda\envs\219-ass01\lib\site-packages\umap\umap_.py:1943: UserWarning: n_jobs value -1 overridden to 1 by set ting random_state. Use no seed for parallelism. warn(f"n jobs value {self.n jobs} overridden to 1 by setting random state. Use no seed for parallelism.")

UMAP Projection of the Normalized GLoVE-based Embeddings

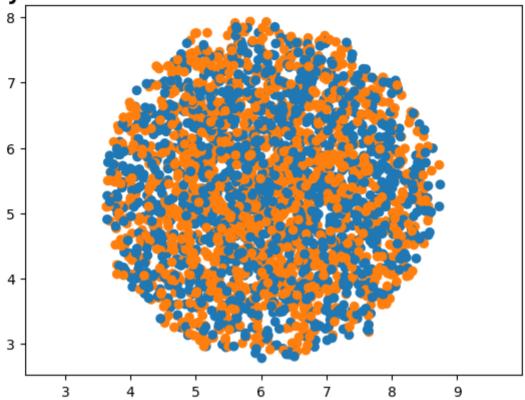


```
# Generate random points and normalize them
def generate_and_normalize_points(num_samples, vector_size):
    random_samples = np.random.normal(loc=0.0, scale=1.0, size=(num_samples, vector_size))
```

```
normalized samples = [sample / np.linalg.norm(sample) for sample in random samples]
   return normalized samples
# Apply UMAP on the random normalized vectors
def apply umap on normalized vectors(normalized vectors, labels):
    embedding result = umap model.fit transform(normalized vectors)
   # PLot
   plt.scatter(
       embedding result[:, 0],
       embedding result[:, 1],
       c=[sns.color palette()[x] for x in labels])
   plt.gca().set aspect('equal', 'datalim')
   plt.title('UMAP Projection of the Random Normalized Vector', fontsize=24)
   plt.show()
# Generate and normalize random points
normalized random vectors = generate and normalize points(2780, 300)
# Apply UMAP on the random normalized vectors
apply umap on normalized vectors(normalized random vectors, labels)
```

c:\Users\Michael_Mi\.conda\envs\219-ass01\lib\site-packages\umap\umap_.py:1943: UserWarning: n_jobs value -1 overridden to 1 by set ting random_state. Use no seed for parallelism. warn(f"n jobs value {self.n jobs} overridden to 1 by setting random state. Use no seed for parallelism.")

UMAP Projection of the Random Normalized Vector



Q13 Text Answer:

When comparing the two virtualizations, the plot of normalized GLOVE shows clusters, however, the plot with random vectors lacks clusters, and does not show any clear meaningful distribution. This means that GLOVE is effective when capture semantic similar contents, and this is the cause of the distinct clusters in the graph. On the other hand This emphasized the importance of meaningful embedding, and this also underscored the GLOVE's ability to representing content in document classification.