Subreddit prediction

1. Description of the project

Project overview

This project aims to develop machine learning models for **analyzing Reddit text** to determine the origin subreddit of a given post or comment. Reddit, a popular social media platform, is organized into a variety of thematic communities known as *subreddits*, where users share content and engage in discussions.

Objective

The primary objective is to build a model that can **predict the subreddit** of a Reddit post or comment. Given a text entry from Reddit, the model will identify which of the following subreddits it originally came from:

- Toronto
- Brussels
- London
- Montreal

This defines a multiclass classification problem

Approach

This project consists of two main parts:

1. Implement a Bernoulli Naïve Bayes Classifier from Scratch

First, a Bernoulli Naïve Bayes classifier will be developed from the ground up, without relying on external libraries for the core algorithm. This implementation will provide a deeper understanding of how the Bernoulli Naïve Bayes method works and how it can be applied to text classification.

2. Utilize a Classifier from Scikit-Learn

In the second part, a pre-built classifier from the scikit-learn library will be used to perform the same task. This comparison will allow us to evaluate the effectiveness of our custom implementation against a widely used, optimized machine learning library.

2. Modules importation

Module importation

import numpy as np

```
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns
from sklearn.feature extraction.text import CountVectorizer
from sklearn.feature extraction.text import TfidfVectorizer
from sklearn.decomposition import PCA
from sklearn.feature_selection import mutual info classif
from sklearn.feature selection import SelectKBest
import time
import nltk
from nltk.tokenize import word tokenize
from nltk.stem import WordNetLemmatizer
from nltk.corpus import stopwords, words
import langid
# Ensure required NLTK resources are downloaded
try:
    nltk.download('punkt')
    nltk.download('stopwords')
    nltk.download('words')
except Exception as e:
    print(f"Error downloading NLTK resources: {e}")
# Define stopwords list
specific stopwords = ["https", "subreddit", "www", "com"] ## some
specific words for the given dataset
stopwords list = stopwords.words('english') +specific stopwords +
stopwords.words('french') # dataset is both in english and in french
[nltk data] Downloading package punkt to /home/clatimie/nltk data...
[nltk data]
              Package punkt is already up-to-date!
[nltk data] Downloading package stopwords to
                /home/clatimie/nltk data...
[nltk data]
[nltk_data]
              Package stopwords is already up-to-date!
[nltk data] Downloading package words to /home/clatimie/nltk data...
[nltk data]
              Package words is already up-to-date!
```

3. Bernoulli Naïve Bayes Classifier

```
# Bernoulli Naïve Bayes
class NaiveBayesClassifier:
    def __init__(self, laplace_alpha, unique_labels):
        self.alpha = laplace_alpha # true for performing Laplace
smoothing
    self.classes = unique_labels
```

```
self.thetak = None
        self.theta j k = None
    def fit(self, X, y):
        # Laplace smoothing parameters
        n k = self.classes.shape[0] # number of classes
        n j = X.shape[1] # number of features
        n samples = X.shape[0] # number of samples
        self.theta k = np.zeros(n k) # probability of class k
        self.theta j k = np.zeros((n k, n j)) # probability of
feature j given class k
        # compute parameters
        for k in range(n k):
            count k = (y==self.classes[k]).sum()
            self.theta k[k] = count_k / n_samples
            for j in range(n j):
                self.theta j k[k][j] = (X[y==self.classes[k], j].sum()
+self.alpha) / (count k+2*self.alpha)
    def predict(self, X):
        theta k = self.theta k \# Prior probabilities P(y)
        theta j k = self.theta j k # Conditional probabilities P(X|y)
for each feature and class
        # Calculate log probabilities for P(y) and P(X|y)
        log theta k = np.log(theta k) # Shape (num classes,)
        log theta j k = np.log(theta j k) # Shape (num classes,
num features)
        log one minus theta j k = np.log(1 - theta j k) # Shape
(num classes, num features)
        # Calculate the log probabilities of each sample in X for each
class
        probs = (X @ log theta j k.T) + ((1 - X) @
log one minus theta j k.T) + log theta k
        # Choose the class with the highest probability
        y pred = np.argmax(probs, axis=1)
        # Transform back to text-based values (class labels)
        return self.classes[y pred]
    def accu eval(self, X, y):
        # Predict the classes for the input data
        predicted classes = self.predict(X)
        # Ensure the predicted classes are in the correct shape
```

```
# If predicted classes is already 1D, reshaping is not
necessary
        if predicted classes.ndim == 1:
            predicted classes = predicted classes.reshape((-1, 1))
        # Convert y to a NumPy array if it's a Pandas Series
        if isinstance(y, pd.Series):
            y = y.to numpy()
        # Calculate accuracy: compare predicted classes with true
labels
        accuracy = np.mean(predicted classes.flatten() == y.flatten())
        accuracy per class = np.zeros((len(self.classes)))
        # Calculate accuracy per class
        for i, cls in enumerate(self.classes):
            # Find indices where the true label is the current class
            class indices = (y == cls)
            # Calculate the accuracy for the current class
            if np.sum(class indices) > 0: # Avoid division by zero
                accuracy_per_class[i] =
np.mean(predicted classes[class indices] == y[class_indices])
        return accuracy, accuracy per class
   def k fold cross validation(self, k, X, y, print info=True):
        # Performs k-fold cross-validation to evaluate the model's
performance
        num samples = X.shape[0] # Get number of samples in dataset
        indices = np.arange(num samples)
        np.random.seed(10)
        np.random.shuffle(indices) # Shuffle the indices
        X = X[indices] # Apply shuffled indices to X
        y = y[indices] # Apply shuffled indices to y to maintain
correspondence
        fold size = num samples // k # Calculate size of each fold
        accuracies = [] # Initialize list to store accuracies for
each fold
        accuracies training = [] # Initialize list for training
accuracies
        accuracies per class = []
        for fold in range(k):
            if print info:
                print(f"\nFold : {fold + 1}") # Print current fold
number
```

```
test start = fold * fold size # Start index for test set
            test end = (fold + 1) * fold size if fold < k - 1 else
num samples # End index for test set
            X test = X[test start:test end, :] # Create test set
           y test = y[test start:test end] # Corresponding target
values for test set
            X train = np.vstack((X[:test start, :], X[test end:, :]))
# Create training set
            y train = np.concatenate((y[:test start], y[test end:]))
# Corresponding target values for training set
            if print info:
                print(f"Class distribution within training dataset :")
# Print class distribution
                for k in range(0, len(self.classes)):
                    print(f'Proportion of class {self.classes[k]} :
{np.sum(y train==self.classes[k])/len(y train)*100} %')
            self.fit(X train, y train) # Fit model on training set
            accu valid, accu valid per class = self.accu eval(X test,
y test) # Evaluate accuracy on test set
            accuracies.append(accu valid)
            accuracies per class.append(accu valid per class)
            accu_training,_ = self.accu_eval(X_train, y_train)
            accuracies training.append(accu training) # Evaluate
accuracy on training set
            if print info:
                print(f"\n Accuracy = {accuracies[-1]}") # Print
accuracy for current fold
                print(f"\n Accuracies per class
{accuracies_per_class[-1]}")
        accuracies = np.array(accuracies) # Convert accuracies list
to NumPy array
        mean accuracies = np.mean(accuracies) # Calculate mean
accuracy across folds
        mean accuracies training = np.mean(accuracies training) #
Calculate mean training accuracy across folds
        std accuracies = np.std(accuracies) # Calculate standard
deviation of accuracies
        mean accu per class = np.mean(np.array(accuracies per class),
axis=0)
        return mean accuracies, std accuracies,
mean accuracies training, mean accu per class
   def predict and save(self, x, path):
        # Example of how to predict classes
```

4. Lemma and STEM Tokenizer

```
class LemmaTokenizer:
   def init (self, stopwords=None):
        self.wnl = WordNetLemmatizer()
        self.stop words = stopwords
   def call (self, doc):
        # Tokenize the document and apply lemmatization and filtering
        return [
            self.wnl.lemmatize(t, pos="v") for t in word tokenize(doc)
            if t.isalpha() and t.lower() not in self.stop words]
class StemTokenizer:
   def __init__(self, stop words=None):
        # Initialize the Porter Stemmer
        self.wnl = nltk.stem.PorterStemmer()
        self.stop words = stop words
   def call (self, doc):
        # Tokenize the document
        tokens = word tokenize(doc)
        # Process tokens
        return [self.wnl.stem(t) for t in tokens if t.isalpha() and
t.lower() not in self.stop words]
```

5. Dataset analysis

Load training dataset

```
np.random.seed(10) # set a random seed to make results reproductible
# Define the path to the training data file
path_training = "../datasets/Train.csv"
```

```
# Read the CSV file into a pandas DataFrame
training data = pd.read csv(path training, delimiter=',')
# Set column names explicitly for better readability
training data.columns = ['text', 'subreddit']
# Shuffle dataset
training data = training data.sample(frac=1,
random state=42).reset index(drop=True)
# Separate the training data into two series: texts and subreddit
labels
x train = training data['text'] # Contains the Reddit posts
or comments
y train = training data['subreddit'] # Contains the subreddit each
post originates from
# Get unique subreddit labels
unique labels = np.unique(y train) # List of unique subreddits in
the dataset
n samples training = x train.shape[0]
n classes = unique labels.shape[0]
print(f"Training dataset has {n samples training} examples and there
are {n classes} classes")
Training dataset has 1399 examples and there are 4 classes
```

Load test dataset

```
# Define the path to the training data file
path_test = "../datasets/Test.csv"

# Read the CSV file into a pandas DataFrame
x_test = pd.read_csv(path_test, delimiter=',')["body"]

n_samples_test = x_test.shape[0]
print(f"Test dataset has {n_samples_test} examples")

Test dataset has 600 examples
```

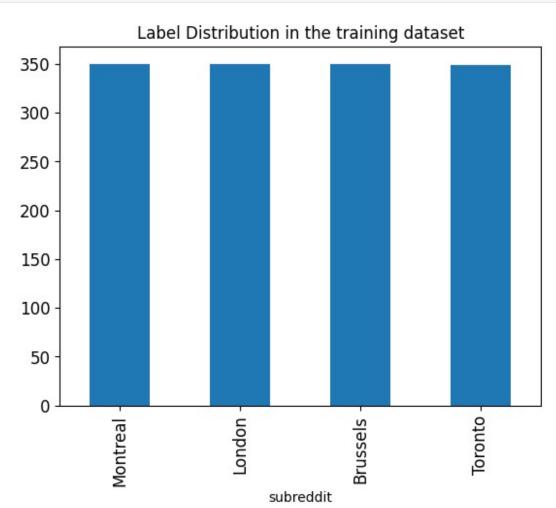
Inspect training dataset

Labels distribution

```
# Show distribution of examples per class
df = pd.DataFrame(training_data)
# Count the number of samples for each label
label_counts = df['subreddit'].value_counts()
```

```
# Plot the distribution
label_counts.plot(kind='bar', title='Label Distribution in the
training dataset', fontsize=12)

<Axes: title={'center': 'Label Distribution in the training dataset'},
xlabel='subreddit'>
```



Text lenght distribution

```
# Calculate the length of each text (in words) for both training and
test datasets
text_lengths_train = x_train.apply(lambda x: len(x.split()))
text_lengths_test = x_test.apply(lambda x: len(x.split()))

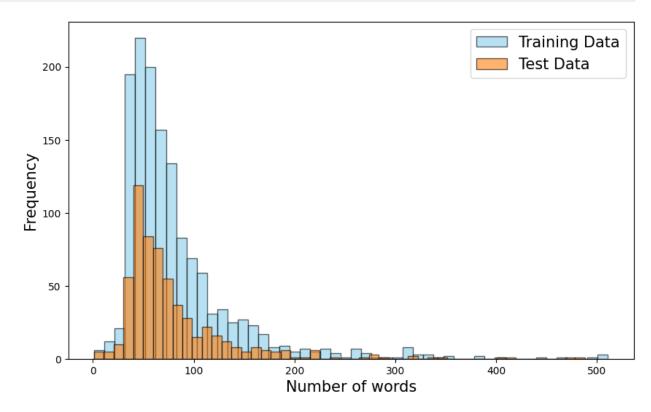
# Plot both histograms on the same figure
plt.figure(figsize=(10, 6))

# Plot the training dataset histogram
plt.hist(text_lengths_train, bins=50, color='skyblue',
edgecolor='black', alpha=0.6, label='Training Data')
```

```
# Plot the test dataset histogram
plt.hist(text_lengths_test, bins=50, color='tab:orange',
edgecolor='black', alpha=0.6, label='Test Data')

# Add labels and title
plt.xlabel('Number of words', fontsize=15)
plt.ylabel('Frequency', fontsize=15)
# Add legend
plt.legend(fontsize=15)

# Show the plot
plt.show()
```



Most distinctive words analysis

```
def classify_language(comment):
    language, _ = langid.classify(comment)
    return 'Montreal (english)' if language == 'en' else 'Montreal
(french)' if language == 'fr' else 'Montreal (english)'

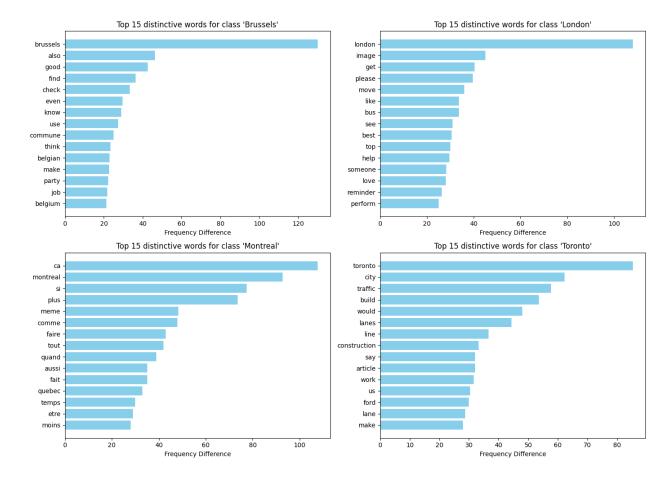
# Modify the labels for comments in the Montreal class
y_train_mtl_distinct = [] # To hold modified labels

for comment, label in zip(x_train, y_train):
    if label == 'Montreal':
        language = classify_language(comment)
```

```
v train mtl distinct.append(language)
    else:
        y train mtl distinct.append(label)
def plot most distinctive words frequency(top n plot, texts train,
y train, top n selected, plots=True):
    unique_labels = sorted(set(y_train)) # Get unique classes
    label texts = {label: [] for label in unique labels} # Dictionary
to hold texts per class
    # Separate texts by label
    for text, label in zip(texts train, y train):
        label_texts[label].append(text)
    # Fit CountVectorizer with the custom tokenizer
    vectorizer = CountVectorizer(
        token_pattern=r'\b[a-zA-Z]{2,}\b',
        stop words=stopwords list,
        tokenizer=LemmaTokenizer(stopwords=stopwords list),
        strip accents="unicode"
    )
    vectorizer.fit(texts train)
    feature names = vectorizer.get feature names out()
    # Initialize a dictionary to store word frequencies per class
    word frequencies = {label: np.zeros(len(feature names)) for label
in unique labels}
    # Calculate word frequencies for each word in each class
    for label in unique labels:
        count matrix = vectorizer.transform(label texts[label])
        word frequencies[label] =
np.array(count matrix.sum(axis=0)).flatten()
    # List to hold the top distinctive words across all classes
    all distinctive words = []
    if plots:
        # Set up the figure with subplots
        n labels = len(unique labels)
        n cols = 2 # Number of columns for subplots
        n rows = (n labels + n cols - 1) // n cols # Calculate number
of rows required
        fig, axes = plt.subplots(n rows, n cols, figsize=(14, 10)) #
Adjust grid size
        axes = axes.flatten() # Flatten axes array for easy indexing
    for i, label in enumerate(unique labels):
        # Calculate distinctiveness by comparing word frequency of
```

```
this class to the average in other classes
        other classes = [lbl for lbl in unique labels if lbl != label]
        if label == "montreal english":
            avg freg other classes =
np.mean([word frequencies[other_label] for other_label in
other_classes if other_label != "montreal_french"], axis=0)
        elif label == "montreal french":
            avg_freq_other_classes =
np.mean([word frequencies[other label] for other label in
other_classes if other_label != "montreal_english"], axis=0)
        else:
            avg freq other classes =
np.mean([word frequencies[other label] for other label in
other classes], axis=0)
        # Calculate distinctiveness score (frequency in this class
minus average frequency in other classes)
        distinctiveness scores = word frequencies[label] -
avg freq other classes
        # Get the indices of the top N distinctive words
        if label == "montreal english" or label == "montreal french":
            top n selected mt\overline{l} = int(top n selected*0.6)
            top indices = np.argsort(distinctiveness scores)[-
top n selected mtl:][::-1] # Indices of top N scores in descending
order
        else:
            top indices = np.argsort(distinctiveness scores)[-
top n selected:][::-1] # Indices of top N scores in descending order
        # Select the top N distinctive words and their scores
        distinctive words = [feature names[idx] for idx in
top_indices]
        distinctive scores = [distinctiveness scores[idx] for idx in
top indices]
        # Extend the all distinctive words list with the current
class's words
        all distinctive words.extend(distinctive words)
        if plots:
            ax = axes[i]
            ax.barh(distinctive words[0:top n plot],
distinctive scores[0:top n plot], color='skyblue')
            ax.set xlabel("Frequency Difference")
            ax.set title(f"Top {top n plot} distinctive words for
class '{label}'")
            ax.invert yaxis() # Invert y-axis to have the most
distinctive words on top
```

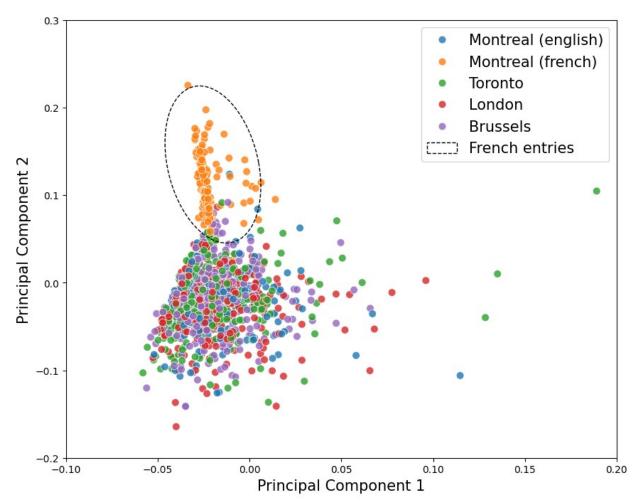
```
# Adjust layout and show the figure
    if plots:
        for j in range(i + 1, len(axes)):
            axes[i].axis('off')
        plt.tight layout()
        plt.show()
    # Return the merged list of top distinctive words across all
    return list(set(all distinctive words)) # Convert to set to
remove duplicates and back to list
token = plot most distinctive words frequency(15, x train, y train,
top n selected=500, plots=True)
/home/clatimie/myenv/lib/python3.12/site-packages/sklearn/
feature extraction/text.py:521: UserWarning: The parameter
'token pattern' will not be used since 'tokenizer' is not None'
  warnings.warn(
/home/clatimie/myenv/lib/python3.12/site-packages/sklearn/feature_extr
action/text.py:406: UserWarning: Your stop_words may be inconsistent
with your preprocessing. Tokenizing the stop words generated tokens
['could', 'etaient', 'etais', 'etait', 'etant', 'etante', 'etantes',
'etants', 'ete', 'etee', 'etees', 'etiez', 'etions', 'eumes', 'eutes', 'fume', 'futes', 'meme', 'might', 'must', 'need', 'sha',
'wo', 'would'] not in stop_words.
  warnings.warn(
```



PCA Analysis

```
from matplotlib.patches import Ellipse
# PCA Analysis with TF-IDF vectorization
vectorizer = TfidfVectorizer(
    lowercase=True,
    tokenizer=LemmaTokenizer(stopwords=stopwords_list)
X_tfidf = vectorizer.fit_transform(x_train)
# Use PCA to reduce dimensionality to 2D
pca = PCA(n components=2)
X_pca = pca.fit_transform(X_tfidf)
# Plot the PCA result with labels
plt.figure(figsize=(10, 8))
sns.scatterplot(x=X_pca[:, 0], y=X_pca[:, 1],
hue=y_train_mtl_distinct, palette='tab10', s=60, alpha=0.8)
# Define the ellipse properties
ellipse = Ellipse(
    xy=(-0.02, 0.135), # Center of the ellipse (mean of the points)
    width=0.18, # Width of the ellipse
```

```
height=0.05, # Height of the ellipse
    angle=95, # Rotation angle of the ellipse
    edgecolor='black', # Color of the ellipse edge
    facecolor='none', # No fill inside the ellipse
    lw=1.
    linestyle='--',
    label="French entries"
)
# Add the ellipse to the plot
plt.gca().add patch(ellipse)
# Add titles and labels
plt.xlabel("Principal Component 1", fontsize=15)
plt.xlim(-0.1, 0.2)
plt.ylim(-0.2, 0.3)
plt.ylabel("Principal Component 2", fontsize=15)
plt.legend(loc='best', fontsize=15)
plt.show()
```



6. Vectorization of the Training Texts (BNB)

To utilize the texts in machine learning models, it is essential to convert them into a vectorized format. Below are several methods available for encoding texts as vectors.

Codes

Hyperparameter for BNB

```
def grid search naive bayes distinctiveness(x train, y train,
max_features_list, y_train_mtl, k_cv=10):
    best accuracy = 0
    best params = {}
    results = []
    # Iterate over all max features
    for max features in (max features list):
        print(f"Testing max features={max features}")
        vocab =
np.unique(np.array(plot most distinctive words frequency(20, x train,
y_train_mtl, top_n_selected=max_features, plots=False)))
        vectorizer = CountVectorizer(
            binary=True, # vectorized vector must be binary for Naive
Bayes
            lowercase=True, # words must be in lowercases
            vocabulary=vocab
        x train distinctiveness = vectorizer.fit transform(x train)
        classifier = NaiveBayesClassifier(laplace alpha=1,
unique labels=unique labels)
        time start = time.time()
        mean accuracy, mean std, mean training accuracy,
mean accu per class = classifier.k fold cross validation(k cv,
x_train_distinctiveness.todense(), y_train, print_info=False)
        mean computation time = 1/k cv * (time.time() - time start)
        # Calculate mean accuracy across folds
        results.append((max features, mean accuracy, mean std,
mean training accuracy, mean computation time, mean accu per class))
        # Update best params if current mean accuracy is the highest
        if mean accuracy > best accuracy:
            best accuracy = mean accuracy
            best_params = {'max_features': max features}
    # Output the results of the grid search
```

```
print("\nGrid search results:")
    for max features, accuracy, std, mean training accuracy,
mean computation time, mean accu per class in results:
        print(f"max features: {max features} -> Mean Accuracy:
{accuracy:.4f}")
    max features values = [result[0] for result in results]
    mean accuracies = [result[1] for result in results]
    mean stds = [result[2] for result in results]
    mean training accuracies = [result[3] for result in results]
    mean accu per class = np.array([result[5] for result in results])
    # Create a new figure for plotting
    plt.figure(figsize=(10, 6))
    plt.plot(max features values, mean training accuracies,
label='Training Accuracy', color='g', marker='o', linewidth=2)
    # Add labels and title
    plt.xlabel("Max features per class labels", fontsize=15)
    plt.ylabel("Mean accuracy", color='k', fontsize=15)
plt.title("Feature selection using distinctiveness scoring")
    plt.legend(loc='upper left')
    # Create a secondary y-axis for validation accuracy
    ax2 = plt.qca().twinx()
    ax2.plot(max_features_values, mean_accuracies, label='Validation
Accuracy', color='b', marker='o', linewidth=2)
    ax2.plot(max features values, mean accu per class[:,0],
label='Validation Accuracy - Brussels', color='tab:orange',
marker='+', linestyle='--')
    ax2.plot(max_features_values, mean_accu_per_class[:,1],
label='Validation Accuracy - London', color='tab:red', marker='+',
linestyle='--')
    ax2.plot(max features values, mean accu per class[:,2],
label='Validation Accuracy - Montreal', color='tab:purple',
marker='+', linestyle='--')
    ax2.plot(max features values, mean accu per class[:,3],
label='Validation Accuracy - Toronto', color='tab:grey', marker='+',
linestyle='--')
    ax2.set_ylabel("Validation Accuracy", fontsize=15)
    ax2.tick params(axis='y')
    # Show both leaends
    ax2.legend(loc='lower right')
    # Show the plot
    plt.show()
```

```
print(f"\nBest parameter:
max features={best params['max features']} with
accuracy={best accuracy:.4f}")
    return best params, best accuracy
def grid search naive bayes mutual information(x train, y train,
max features list, k cv=10):
    best accuracy = 0
    best params = {}
    results = []
    # Iterate over all max features
    for max features in (max features list):
        print(f"Testing max features={max features}")
        vectorizer = CountVectorizer(
            binary=True, # vectorized vector must be binary for Naive
Bayes
            lowercase=True, # words must be in lowercases
            tokenizer=LemmaTokenizer(stopwords=stopwords list)
        )
        x train = vectorizer.fit transform(x train)
        x train new = SelectKBest(mutual info classif,
k=max features).fit transform(x train, y train)
        classifier = NaiveBayesClassifier(laplace alpha=1,
unique labels=unique labels)
        time start = time.time()
        mean accuracy, mean_std, mean_training_accuracy,
mean_accu_per_class = classifier.k_fold_cross_validation(k cv,
x train_new.todense(), y_train, print_info=False)
        mean computation time = 1/k cv * (time.time() - time start)
        # Calculate mean accuracy across folds
        results.append((max features, mean accuracy, mean std,
mean training accuracy, mean computation time, mean accu per class))
        # Update best params if current mean accuracy is the highest
        if mean accuracy > best accuracy:
            best accuracy = mean accuracy
            best params = {'max features': max features}
    # Output the results of the grid search
    print("\nGrid search results:")
    for max_features, accuracy, std, mean_training_accuracy,
mean computation time, mean accu per_class in results:
        print(f"max features: {max features} -> Mean Accuracy:
```

```
{accuracy:.4f}")
    max features values = [result[0] for result in results]
    mean accuracies = [result[1] for result in results]
    mean stds = [result[2] for result in results]
    mean training accuracies = [result[3] for result in results]
    mean accu per class = np.array([result[5] for result in results])
    # Create a new figure for plotting
    plt.figure(figsize=(10, 6))
    plt.plot(max features values, mean training accuracies,
label='Training Accuracy', color='g', marker='o', linewidth=2)
    # Add labels and title
    plt.xlabel("Max features per class labels", fontsize=15)
    plt.ylabel("Mean accuracy", color='k', fontsize=15)
    plt.title("Feature selection using mutual information scoring")
    plt.legend(loc='upper left')
    # Create a secondary y-axis for validation accuracy
    ax2 = plt.qca().twinx()
    ax2.plot(max features values, mean accuracies, label='Validation
Accuracy', color='b', marker='o', linewidth=2)
    ax2.plot(max features values, mean accu per class[:,0],
label='Validation Accuracy - Brussels', color='tab:orange',
marker='+', linestyle='--')
    ax2.plot(max features values, mean accu per class[:,1],
label='Validation Accuracy - London', color='tab:red', marker='+',
linestyle='--')
    ax2.plot(max features values, mean accu per class[:,2],
label='Validation Accuracy - Montreal', color='tab:purple',
marker='+', linestyle='--')
    ax2.plot(max features values, mean accu per class[:,3],
label='Validation Accuracy - Toronto', color='tab:grey', marker='+',
linestyle='--')
    ax2.set ylabel("Validation Accuracy", fontsize=15)
    ax2.tick params(axis='y')
    # Show both legends
    ax2.legend(loc='lower right')
    # Show the plot
    plt.show()
    print(f"\nBest parameter:
max_features={best_params['max_features']} with
accuracy={best accuracy:.4f}")
```

```
return best_params, best_accuracy

#grid_search_naive_bayes_distinctiveness(x_train, y_train,
np.arange(50, 2000, 200), y_train_mtl_distinct, k_cv=10)

#grid_search_naive_bayes_mutual_information(x_train, y_train,
np.arange(50, 4000, 200), k_cv=10)
```

7. K-fold cross validation (BNB + Distinctiveness)

```
k cv = 10
vocab = np.unique(np.array(plot most distinctive words frequency(20,
x train, y train mtl distinct, top n selected=650, plots=False)))
vectorizer = CountVectorizer(
    binary=True, # vectorized vector must be binary for Naive Bayes
    lowercase=True, # words must be in lowercases
    vocabulary=vocab
)
x train distinctiveness = vectorizer.fit transform(x train)
print(f"Feature selection based on distinctiveness ranking: vectorized
training dataset has {x train distinctiveness.shape[1]}
tokens/features")
classifier = NaiveBayesClassifier(laplace alpha=1,
unique labels=unique labels)
time start = time.time()
mean_accuracy, mean_std, mean_training_accuracy, mean_accu_per_class =
classifier.k fold cross validation(k cv,
x_train_distinctiveness.todense(), y_train, print_info=False)
mean computation time = (time.time() - time start)
print(f'Mean accuracy (training) accross {k cv}-fold cross
validation : {mean training accuracy}')
print(f'Mean variance of validation accuracy accross {k cv}-fold cross
validation : {mean_std}')
print(f'Mean validation accuracy accross {k cv}-fold cross
validation : {mean accuracy}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class Brussels : {mean accu per class[0]}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class London : {mean accu per class[1]}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class Montreal : {mean accu per class[2]}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class Toronto : {mean accu per class[3]}')
print(f'Computation time accross {k cv}-fold cross validation:
{mean computation time}')
```

```
/home/clatimie/myenv/lib/python3.12/site-packages/sklearn/
feature extraction/text.py:521: UserWarning: The parameter
'token_pattern' will not be used since 'tokenizer' is not None'
  warnings.warn(
/home/clatimie/myenv/lib/python3.12/site-packages/sklearn/feature extr
action/text.py:406: UserWarning: Your stop_words may be inconsistent
with your preprocessing. Tokenizing the stop words generated tokens
['could', 'etaient', 'etais', 'etait', 'etant', 'etante', 'etantes',
'etants', 'ete', 'etee', 'etees', 'etes', 'etiez', 'etions', 'eumes', 'eutes', 'fume', 'futes', 'meme', 'might', 'must', 'need', 'sha',
'wo', 'would'] not in stop words.
 warnings.warn(
Feature selection based on distinctiveness ranking: vectorized
training dataset has 2732 tokens/features
Mean accuracy (training) accross 10-fold cross validation :
0.8695141030033117
Mean variance of validation accuracy accross 10-fold cross
validation: 0.04333134284106084
Mean validation accuracy accross 10-fold cross validation :
0.7106455376239549
Mean validation accuracy accross 10-fold cross validation for class
Brussels: 0.8056171622402777
Mean validation accuracy accross 10-fold cross validation for class
London: 0.7796315645274643
Mean validation accuracy accross 10-fold cross validation for class
Montreal: 0.5035571753937008
Mean validation accuracy accross 10-fold cross validation for class
Toronto: 0.7484169322511749
Computation time accross 10-fold cross validation: 4.5911760330200195
```

8. K-fold cross validation (BNB + Mutual Information)

```
classifier = NaiveBayesClassifier(laplace alpha=1,
unique labels=unique labels)
time start = time.time()
mean accuracy, mean std, mean training accuracy, mean accu per class =
classifier.k fold cross validation(k cv, x train mi.todense(),
y train, print info=False)
mean_computation_time = (time.time() - time start)
print(f'Mean accuracy (training) accross {k_cv}-fold cross
validation : {mean training accuracy}')
print(f'Mean variance of validation accuracy accross {k cv}-fold cross
validation : {mean std}')
print(f'Mean validation accuracy accross {k cv}-fold cross
validation : {mean accuracy}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class Brussels : {mean accu per class[0]}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class London : {mean accu per class[1]}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class Montreal : {mean accu per class[2]}')
print(f'Mean validation accuracy accross {k cv}-fold cross validation
for class Toronto : {mean_accu_per_class[3]}')
print(f'Computation time accross {k cv}-fold cross validation:
{mean computation time}')
classifier.fit(x train mi.todense(), y train)
x test = vectorizer.transform(x test)
x test mi = selector.transform(x test)
y pred = classifier.predict(x test mi.todense())
y pred = y pred.flatten() if len(y pred.shape) > 1 else y pred
# Construct the DataFrame and save to CSV
results df = pd.DataFrame({
    'id': range(len(y pred)),
    'subreddit': y pred
})
# Save predictions to CSV
results df.to csv("../output/submissions mutual information bnb.csv",
index=False)
print("Predictions saved to
../output/submissions mutual information bnb.csv")
```

/home/clatimie/myenv/lib/python3.12/site-packages/sklearn/
feature_extraction/text.py:521: UserWarning: The parameter
'token_pattern' will not be used since 'tokenizer' is not None'
warnings.warn(

Feature selection based on mutual information ranking: vectorized training dataset has 2850 tokens/features

Mean accuracy (training) accross 10-fold cross validation: 0.8590310608655933

Mean variance of validation accuracy accross 10-fold cross validation: 0.04346068097414368

Mean validation accuracy accross 10-fold cross validation: 0.749863892669648

Mean validation accuracy accross 10-fold cross validation for class Brussels: 0.8170787233493352

Mean validation accuracy accross 10-fold cross validation for class London: 0.8835763419696704

Mean validation accuracy accross 10-fold cross validation for class Montreal: 0.5223190772951375

Mean validation accuracy accross 10-fold cross validation for class Toronto : 0.7713861288476179

Computation time accross 10-fold cross validation: 4.608723878860474 Predictions saved to ../output/submissions_mutual_information_bnb.csv

File overview

This notebook implements **Support Vector Machines (SVM)** classification for the subreddit prediction dataset. Hyperparameter tuning is performed, and the model's accuracy is evaluated using **10-fold cross-validation**.

Load modules

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import warnings
warnings.filterwarnings("ignore", category=UserWarning) # This will
suppress UserWarnings
import time
from sklearn.feature extraction.text import TfidfVectorizer
from sklearn.preprocessing import StandardScaler
from sklearn.svm import SVC
from sklearn.model selection import KFold
from sklearn.metrics import accuracy score, classification report
from sklearn.feature_selection import mutual info classif
from sklearn.feature selection import SelectKBest
from sklearn.model selection import GridSearchCV
from sklearn.pipeline import Pipeline
import nltk
from nltk.tokenize import word tokenize
from nltk.stem import WordNetLemmatizer
from nltk.corpus import stopwords, words
# Ensure required NLTK resources are downloaded
try:
    nltk.download('punkt')
    nltk.download('stopwords')
    nltk.download('words')
except Exception as e:
    print(f"Error downloading NLTK resources: {e}")
# Define stopwords list
specific_stopwords = ["https", "subreddit", "www", "com"] ## some
specific words for the given dataset
stopwords list = stopwords.words('english') +specific stopwords +
stopwords.words('french') # dataset is both in english and in french
```

```
[nltk_data] Downloading package punkt to /home/clatimie/nltk_data...
[nltk_data] Package punkt is already up-to-date!
[nltk_data] Downloading package stopwords to
[nltk_data] /home/clatimie/nltk_data...
[nltk_data] Package stopwords is already up-to-date!
[nltk_data] Downloading package words to /home/clatimie/nltk_data...
[nltk_data] Package words is already up-to-date!
```

Load training dataset

```
# Define the path to the training data file
path_training = "../datasets/Train.csv"
# Read the CSV file into a pandas DataFrame
training data = pd.read csv(path training, delimiter=',')
# Set column names explicitly for better readability
training data.columns = ['text', 'subreddit']
# Shuffle dataset
training data = training data.sample(frac=1,
random state=42).reset index(drop=True)
# Separate the training data into two series: texts and subreddit
labels
x_train = training_data['text'] # Contains the Reddit posts
or comments
y_train = training_data['subreddit'] # Contains the subreddit each
post originates from
# Get unique subreddit labels
unique labels = np.unique(y train) # List of unique subreddits in
the dataset
n samples training = x train.shape[0]
n classes = unique labels.shape[0]
print(f"Training dataset has {n_samples_training} examples and there
are {n classes} classes")
Training dataset has 1399 examples and there are 4 classes
```

Load test dataset

```
# Define the path to the training data file
path_test = "../datasets/Test.csv"

# Read the CSV file into a pandas DataFrame
x_test = pd.read_csv(path_test, delimiter=',')["body"]
```

```
n_samples_test = x_test.shape[0]
print(f"Test dataset has {n_samples_test} examples")
Test dataset has 600 examples
```

Lemma Tokenizer from NLTK

```
class LemmaTokenizer:
    def __init__(self, stopwords=None):
        self.wnl = WordNetLemmatizer()
        self.stop_words = stopwords

def __call__(self, doc):
    # Tokenize the document and apply lemmatization and filtering
    return [
        self.wnl.lemmatize(t, pos="v") for t in word_tokenize(doc)
        if t.isalpha() and t.lower() not in self.stop_words]
```

Hyperparameters search

```
""" # Define the parameter grid for hyperparameter search
param grid = {
    'svc__kernel': ['linear', 'rbf', 'poly'], # Different kernel
    'select k': [1000, 2000, 3000, 4000], # Different values for top
k features
    'svc C': [0.1, 0.2], # Different values for C (controls slack in
SVM)
    'svc gamma': ['scale', 0.001, 0.01, 0.1] # Gamma values for RBF
and poly kernels
# Define the pipeline
pipeline = Pipeline([
    ('vectorizer', TfidfVectorizer(
        lowercase=True,
        tokenizer=LemmaTokenizer(stopwords=stopwords list)
    ('select', SelectKBest(mutual_info_classif)), # Placeholder for k
parameter
    ('scaler', StandardScaler(with mean=False)), # Use
with mean=False for sparse data
    ('svc', SVC()) # SVM classifier
1)
# Use GridSearchCV to find the best combination of hyperparameters
grid search = GridSearchCV(
   estimator=pipeline,
   param grid=param grid,
   cv=5, # 10-fold cross-validation
```

```
scoring='accuracy',
    verbose=3, # To display progress
    n jobs=-1 # Use all available cores
# Fit the model to the training data and search for best parameters
grid search.fit(x_train, y_train)
# Get the best parameters and corresponding score
best params = grid search.best params
best score = grid search.best score
print("Best Parameters:", best params)
print(f"Best Cross-Validated Accuracy: {best_score:.4f}") """
' # Define the parameter grid for hyperparameter search\nparam grid =
      \'svc_kernel\': [\'linear\', \'rbf\', \'poly\'], # Different
                   \'select__k\': [1000, 2000, 3000, 4000], #
kernel options\n
Different values for top k features\n
                                       \'svc C\': [0.1, 0.2],
Different values for C (controls slack in SVM)\n \'svc gamma\':
[\'scale\', 0.001, 0.01, 0.1] # Gamma values for RBF and poly
kernels\n\n# Define the pipeline\npipeline = Pipeline([\n
(\'vectorizer\', TfidfVectorizer(\n
                                          lowercase=True,\n
tokenizer=LemmaTokenizer(stopwords=stopwords list)\n
(\'select\', SelectKBest(mutual info classif)), # Placeholder for k
               (\'scaler\', StandardScaler(with mean=False)), # Use
parameter\n
with mean=False for sparse data\n (\'svc\', SVC()) # SVM
classifier\n])\n\n# Use GridSearchCV to find the best combination of
hyperparameters\ngrid search = GridSearchCV(\n
                                                 estimator=pipeline.\
                                cv=5, # 10-fold cross-validation\n
     param grid=param grid,\n
scoring=\'accuracy\',\n
                          verbose=3, # To display progress\n
n jobs=-1 # Use all available cores\n)\n\n# Fit the model to the
training data and search for best parameters\ngrid search.fit(x train,
y_train)\n\n# Get the best parameters and corresponding score\
nbest params = grid search.best params \nbest score =
grid search.best score \n\nprint("Best Parameters:", best params)\
nprint(f"Best Cross-Validated Accuracy: {best score:.4f}")
```

10-fold cross validation

```
vectorizer = TfidfVectorizer(
    lowercase=True,
    tokenizer=LemmaTokenizer(stopwords=stopwords_list)
)

x_train_tfidf = vectorizer.fit_transform(x_train)

selector = SelectKBest(mutual_info_classif, k=3000)
x_train_mi = selector.fit_transform(x_train_tfidf, y_train)
```

```
scaler = StandardScaler()
x train svc = scaler.fit transform(np.asarray(x train mi.todense()))
classifier = SVC(kernel="rbf",gamma='scale', C=1)
accuracies = []
class_accuracies = {class_name: [] for class_name in set(y_train)} #
To store accuracy for each class
kf = KFold(n splits=10, shuffle=True, random state=42)
fold = 0
# Start measuring time
start time = time.time()
accuracies = []
training accuracies = []
class accuracies = {class name: [] for class name in set(y train)} #
To store accuracy for each class
kf = KFold(n splits=10, shuffle=True, random state=42)
fold = 0
for train index, val index in kf.split(x train svc):
    fold += 1
    X train fold, X val fold = x train svc[train index],
x train svc[val index]
    y fold train, y fold val = y train[train index],
y train[val index]
    # Train the classifier
    classifier.fit(X train fold, y fold train)
    # Predict and evaluate on the validation set
    y pred = classifier.predict(X val fold)
    y_pred_training = classifier.predict(X train fold)
    # Display results for each fold
    print(f"\nFold n°{fold}:")
    # Get accuracy per class
    class accuracy = classification report(y fold val, y pred,
output dict=True)
    print("Classification Report:\n",
classification report(y fold val, y pred))
    accuracy = accuracy score(y fold val, y pred)
    accuracies.append(accuracy)
    accuracy training = accuracy score(y pred training, y fold train)
    training accuracies.append(accuracy training)
```

```
for label, metrics in class accuracy.items():
        if label != 'accuracy' and label!="macro avg" and label!=
"weighted avg":
            class accuracies[label].append(metrics['precision'])
# Compute total time
end time = time.time()
total time = end_time - start_time
print(f"\nTotal computing time for 10 folds: {total time:.2f}
seconds")
# Mean accuracy across 10 folds
mean_accuracy = np.mean(accuracies)
print(f"Mean Accuracy across 10 folds for SVM classifier:
{mean accuracy:.4f}")
# Average accuracy for each class
print("\nAverage Accuracy per Class:")
for label, accuracies in class accuracies.items():
    avg class accuracy = np.mean(accuracies)
    print(f"Class {label}: {avg class accuracy:.4f}")
# Mean training accuracy across 10 folds
mean training accuracy = np.mean(accuracy training)
print(f"Mean training accuracy across 10 folds for SVM classifier:
{mean training accuracy:.4f}")
Fold n°1:
Classification Report:
               precision recall f1-score support
    Brussels
                   0.69
                             0.87
                                       0.77
                                                   38
                   0.71
                             0.84
                                       0.77
                                                   32
      London
    Montreal
                   0.90
                             0.59
                                       0.72
                                                   32
    Toronto
                   0.85
                             0.74
                                       0.79
                                                   38
                                       0.76
                                                   140
    accuracy
                   0.79
                             0.76
                                       0.76
                                                   140
   macro avg
weighted avg
                   0.79
                             0.76
                                       0.76
                                                   140
```

File overview

This notebook implements a stacking model for subreddit prediction. The stacking classifier combines the predictions of multiple models, including Support Vector Machines (SVM) and Bernoulli Naive Bayes (BNB). Hyperparameter tuning is performed, and the model's performance is evaluated using 10-fold cross-validation.

Load modules

```
import numpy as np
import matplotlib.pyplot as plt
import pandas as pd
import seaborn as sns
import warnings
warnings.filterwarnings("ignore", category=UserWarning) # This will
suppress UserWarnings
import time
from sklearn.feature extraction.text import TfidfVectorizer,
CountVectorizer
from sklearn.preprocessing import StandardScaler
from sklearn.svm import SVC
from sklearn.naive bayes import BernoulliNB
from sklearn.linear model import LogisticRegression
from sklearn.model selection import KFold
from sklearn.feature selection import SelectKBest
from sklearn.metrics import accuracy score, classification report
from sklearn.feature selection import mutual info classif
from sklearn.metrics import confusion matrix
import nltk
from nltk.tokenize import word tokenize
from nltk.stem import WordNetLemmatizer
from nltk.corpus import stopwords, words
# Ensure required NLTK resources are downloaded
try:
    nltk.download('punkt')
    nltk.download('stopwords')
    nltk.download('words')
except Exception as e:
    print(f"Error downloading NLTK resources: {e}")
# Define stopwords list
specific stopwords = ["https", "subreddit", "www", "com"] ## some
specific words for the given dataset
```

```
stopwords_list = stopwords.words('english') +specific_stopwords +
stopwords.words('french') # dataset is both in english and in french

[nltk_data] Downloading package punkt to /home/clatimie/nltk_data...
[nltk_data] Package punkt is already up-to-date!
[nltk_data] Downloading package stopwords to
[nltk_data] /home/clatimie/nltk_data...
[nltk_data] Package stopwords is already up-to-date!
[nltk_data] Downloading package words to /home/clatimie/nltk_data...
[nltk_data] Package words is already up-to-date!
```

Load training dataset

```
# Define the path to the training data file
path training = "../datasets/Train.csv"
# Read the CSV file into a pandas DataFrame
training data = pd.read csv(path training, delimiter=',')
# Set column names explicitly for better readability
training_data.columns = ['text', 'subreddit']
# Shuffle dataset
training data = training data.sample(frac=1,
random state=42).reset index(drop=True)
# Separate the training data into two series: texts and subreddit
labels
x train = training data['text'] # Contains the Reddit posts
or comments
y train = training data['subreddit'] # Contains the subreddit each
post originates from
# Get unique subreddit labels
unique labels = np.unique(y_train) # List of unique subreddits in
the dataset
n samples trainings = x train.shape[0]
n classes = unique labels.shape[0]
print(f"Training dataset has {n samples training} examples and there
are {n classes} classes")
Training dataset has 1399 examples and there are 4 classes
```

I Oad test dataset

```
# Define the path to the training data file
path_test = "../datasets/Test.csv"
```

```
# Read the CSV file into a pandas DataFrame
x_test = pd.read_csv(path_test, delimiter=',')["body"]
n_samples_test = x_test.shape[0]
print(f"Test dataset has {n_samples_test} examples")
Test dataset has 600 examples
```

Lemma Tokenizer from NLTK

```
class LemmaTokenizer:
    def __init__(self, stopwords=None):
        self.wnl = WordNetLemmatizer()
        self.stop_words = stopwords

def __call__(self, doc):
    # Tokenize the document and apply lemmatization and filtering
    return [
        self.wnl.lemmatize(t, pos="v") for t in word_tokenize(doc)
        if t.isalpha() and t.lower() not in self.stop_words]
```

10 fold cross validation of the stacking model

```
y binary = [1 if label == "Montreal" else -1 for label in y train] #
for svm training
# Define vectorizers
vectorizer svm = TfidfVectorizer(lowercase=True,
tokenizer=LemmaTokenizer(stopwords=stopwords list),
strip_accents="unicode")
vectorizer bnb = CountVectorizer(lowercase=True,
tokenizer=LemmaTokenizer(stopwords=stopwords list),
strip accents="unicode")
# Define models
svm model = SVC(kernel='rbf', probability=True, gamma='scale', C=1)
bnb model = BernoulliNB()
# Define feature selectors
selector bnb = SelectKBest(mutual info classif, k=2850)
selector svm = SelectKBest(mutual info classif, k=3000)
# Define scaler
scaler svm = StandardScaler()
# Preprocess data before cross-validation
X train bnb = vectorizer bnb.fit transform(x train)
X train svm = vectorizer svm.fit transform(x train)
# Apply feature selection
```

```
X train bnb selected = selector bnb.fit transform(X train bnb,
y train)
X train svm selected = selector svm.fit transform(X train svm,
y binary)
# Scale the SVM features
X train svm scaled =
scaler svm.fit transform(np.asarray(X train svm selected.todense()))
# Prepare KFold cross-validation
kf = KFold(n splits=10, shuffle=True, random state=42)
accuracies = []
training accuracies = []
class accuracies = {class name: [] for class name in set(y train)} #
To store accuracy for each class
mean_conf_matrix = np.zeros((len(np.unique(y_train)),
len(np.unique(y train)))) # Initialize empty confusion matrix
fold = 0
# 10-Fold Cross-Validation
time start = time.time()
for train index, val index in kf.split(X train svm scaled):
    fold += 1
    # Split data into training and validation sets
    X_train_fold_svm, X_val_fold_svm =
X train svm scaled[train index], X train svm scaled[val index]
    X train fold bnb, X val fold bnb =
X train bnb selected[train index], X train bnb selected[val index]
    y train bnb fold, y val bnb fold = np.array(y train)[train index],
np.array(y train)[val index]
    y_train_svm_fold, y_val_svm_fold = np.array(y_binary)
[train_index], np.array(y_binary)[val_index]
    # Train the models
    svm_model.fit(X_train_fold_svm, y_train_svm_fold)
    bnb_model.fit(X_train fold bnb, y train bnb fold)
    # Get predictions from both models
    svm predictions = svm model.predict(X_val_fold_svm)
    bnb predictions = bnb model.predict(X val fold bnb)
    # Vectorized version of combining predictions
    final predictions = np.where(svm predictions == 1, "Montreal",
bnb predictions)
    # Get predictions from both models for training data
    svm predictions training = svm model.predict(X train fold svm)
```

```
bnb predictions training = bnb model.predict(X train fold bnb)
    # Vectorized version of combining predictions for training data
    final predictions training = np.where(svm predictions training ==
1, "Montreal", bnb predictions training)
    # Calculate accuracy for this fold
    accuracy = accuracy score(y val bnb fold, final predictions) #
Use y val bnb fold as the correct target variable
    accuracies.append(accuracy)
    training accuracy = accuracy score(y train bnb fold,
final predictions training)
    training accuracies.append(training accuracy)
    print("Classification Report:\n",
classification_report(y_val_bnb_fold, final_predictions))
    class accuracy = classification report(y val bnb fold,
final predictions, output dict=True)
    for label, metrics in class accuracy.items():
        if label != 'accuracy' and label != "macro avg" and label !=
"weighted avg":
            class accuracies[label].append(metrics['precision'])
    print(f"Validation accuracy for fold {fold}: {accuracy:.4f}")
    print(f"Training accuracy for fold {fold}:
{training accuracy:.4f}\n")
    # Confusion Matrix for this fold
    conf matrix = confusion matrix(y val bnb fold, final predictions)
    # Add this fold's confusion matrix to the cumulative confusion
matrix
    mean conf matrix += conf matrix
time end = time.time()
# Calculate the mean confusion matrix
mean_conf_matrix /= kf.get_n_splits() # Average the confusion matrix
# Plot the mean confusion matrix
plt.figure(figsize=(6, 6))
sns.heatmap(mean conf matrix, annot=True, fmt='.2f', cmap='Blues',
xticklabels=np.unique(y train), yticklabels=np.unique(y train))
plt.title("Mean Confusion Matrix - 10-Fold Cross-Validation")
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.show()
# Calculate the mean accuracy across all folds
```

```
mean accuracy = np.mean(accuracies)
print(f"Mean Accuracy across 10 folds: {mean accuracy:.4f}")
mean_training_accuracy = np.mean(training accuracies)
print(f"Mean Accuracy across 10 folds: {mean training accuracy:.4f}")
# Average accuracy for each class
print("\nAverage Accuracy per Class:")
for label, accuracies in class accuracies.items():
    avg class accuracy = np.mean(accuracies)
    print(f"Class {label}: {avg class accuracy:.4f}")
print(f"Computing time : {time end-time start} (s)")
# Fitting the models with the whole dataset
svm model.fit(X train svm scaled, y_binary)
bnb model.fit(X train bnb selected, y train)
# Preprocess x test
x test bnb = vectorizer bnb.transform(x test)
x test svm = vectorizer svm.transform(x test)
x test bnb selected = selector bnb.transform(x test bnb)
x test svm selected = selector svm.transform(x test svm)
x test svm scaled =
scaler_svm.transform(np.asarray(x test svm selected.todense()))
# Make predictions
svm predictions = svm model.predict(x test svm scaled)
bnb predictions = bnb model.predict(x test bnb selected)
final predictions = []
final predictions = np.where(svm predictions == 1, "Montreal",
bnb predictions)
results df = pd.DataFrame({
    'id': range(len(final predictions)),
    'subreddit': final predictions
})
results df.to csv("../output/stacking.csv", index=False)
Classification Report:
               precision recall f1-score support
    Brussels
                   0.79
                             0.79
                                       0.79
                                                   38
      London
                   0.62
                             0.88
                                       0.73
                                                   32
                   0.83
                                                   32
    Montreal
                             0.75
                                       0.79
```

Toronto	0.89	0.66	0.76	38
accuracy macro avg weighted avg	0.78 0.79	0.77 0.76	0.76 0.77 0.77	140 140 140

Validation accuracy for fold 1: 0.7643 Training accuracy for fold 1: 0.9333

Classification Report:

CCGSSTITCGCTOIL	report.			
	precision	recall	f1-score	support
Brussels	0.82	0.76	0.79	37
London	0.62	0.86	0.72	36
Montreal	1.00	0.82	0.90	28
Toronto	0.88	0.74	0.81	39
accuracy			0.79	140
macro avg	0.83	0.80	0.80	140
weighted avg	0.82	0.79	0.80	140

Validation accuracy for fold 2: 0.7929 Training accuracy for fold 2: 0.9388

Classification Report:

	precision	recall	f1-score	support
Brussels	0.86	0.71	0.78	45
London	0.59	0.87	0.70	30
Montreal	0.84	0.74	0.79	35
Toronto	0.82	0.77	0.79	30
accuracy			0.76	140
macro avg	0.78	0.77	0.77	140
weighted avg	0.79	0.76	0.77	140

Validation accuracy for fold 3: 0.7643 Training accuracy for fold 3: 0.9357

Classification Report:

Crassification	Report:			
	precision	recall	f1-score	support
Brussels	0.82	0.88	0.85	42
London	0.54	0.88	0.67	25
Montreal	0.96	0.58	0.72	43
Toronto	0.79	0.73	0.76	30
accuracy			0.76	140
macro avg	0.78	0.77	0.75	140
weighted avg	0.81	0.76	0.76	140

Validation accuracy for fold 4: 0.7571 Training accuracy for fold 4: 0.9285

Classification Report:

	precision	recall	f1-score	support
Brussels London Montreal Toronto	0.76 0.75 0.96 0.78	0.81 0.87 0.70 0.82	0.78 0.80 0.81 0.80	31 38 37 34
accuracy macro avg weighted avg	0.81 0.81	0.80 0.80	0.80 0.80 0.80	140 140 140

Validation accuracy for fold 5: 0.8000 Training accuracy for fold 5: 0.9444

Classification Report:

C CGCCTC	. оро. с.			
	precision	recall	f1-score	support
Brussels	0.59	0.79	0.68	29
London	0.76	0.84	0.80	38
Montreal	0.89	0.68	0.77	37
Toronto	0.90	0.78	0.84	36
accuracy			0.77	140
macro avg	0.79	0.77	0.77	140
weighted avg	0.80	0.77	0.78	140

Validation accuracy for fold 6: 0.7714 Training accuracy for fold 6: 0.9333

Classification Report:

CCGSSTITCGCTOIL	report :			
	precision	recall	f1-score	support
Brussels	0.82	0.87	0.84	31
London	0.81	0.83	0.82	36
Montreal	0.80	0.87	0.84	38
Toronto	0.86	0.71	0.78	35
accuracy			0.82	140
macro avg	0.82	0.82	0.82	140
weighted avg	0.82	0.82	0.82	140

Validation accuracy for fold 7: 0.8214 Training accuracy for fold 7: 0.9365

Classification Report:

precision	recall	f1-score	support
-----------	--------	----------	---------

Brussels	0.61	0.94	0.74	32
London	0.74	0.66	0.79	44
Montreal	0.94	0.81	0.87	36
Toronto	0.86	0.64	0.73	28
accuracy			0.76	140
macro avg	0.79	0.76	0.76	140
weighted avg	0.79	0.76	0.76	140

Validation accuracy for fold 8: 0.7571 Training accuracy for fold 8: 0.9420

Classification Report:

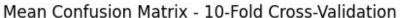
	precision	recall	f1-score	support
	p. 002020			
Brussels	0.73	0.77	0.75	35
London	0.59	0.83	0.69	29
Montreal	0.88	0.66	0.75	35
Toronto	0.89	0.78	0.83	41
accuracy			0.76	140
macro avg	0.77	0.76	0.76	140
weighted avg	0.79	0.76	0.76	140

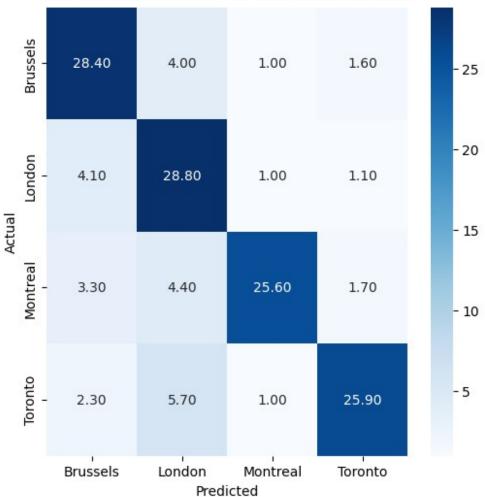
Validation accuracy for fold 9: 0.7571 Training accuracy for fold 9: 0.9380

Classification Report:

	precision	recall	f1-score	support
Brussels London Montreal Toronto	0.69 0.72 0.92 0.88	0.83 0.79 0.76 0.76	0.76 0.75 0.83 0.82	30 42 29 38
accuracy macro avg weighted avg	0.80 0.80	0.79 0.78	0.78 0.79 0.79	139 139 139

Validation accuracy for fold 10: 0.7842 Training accuracy for fold 10: 0.9341





Mean Accuracy across 10 folds: 0.7770 Mean Accuracy across 10 folds: 0.9365

Average Accuracy per Class:

Class Toronto: 0.8547 Class Montreal: 0.9025 Class London: 0.6739 Class Brussels: 0.7502

Computing time : 105.14465403556824 (s)

```
import pandas as pd
import numpy as np
from sklearn.feature extraction.text import TfidfVectorizer
from sklearn.naive bayes import MultinomialNB
from sklearn.linear model import LogisticRegression
from catboost import CatBoostClassifier
from sklearn.model selection import GridSearchCV
from sklearn.decomposition import TruncatedSVD
from sklearn.preprocessing import Normalizer, MinMaxScaler
from sentence transformers import SentenceTransformer
# Load training and test data
train file = 'Train.csv'
test file = 'Test.csv'
output file = 'submissions.csv'
# Training data does not have a header
train data = pd.read csv(train file, header=None, names=['text',
'subreddit'])
test data = pd.read csv(test file)
# Preprocessing metadata (TF-IDF and N-grams)
tfidf vectorizer = TfidfVectorizer(ngram range=(1, 2),
max features=5000)
tfidf features = tfidf vectorizer.fit transform(train data['text'])
# Perform dimensionality reduction on TF-IDF using TruncatedSVD
svd = TruncatedSVD(n components=300, random state=42)
reduced tfidf = svd.fit transform(tfidf features)
# Ensure non-negative features for MultinomialNB
minmax scaler = MinMaxScaler()
non negative tfidf = minmax scaler.fit transform(reduced tfidf)
# Normalize TF-IDF features for Logistic Regression and CatBoost
tfidf normalizer = Normalizer(norm='l2')
normalized train tfidf = tfidf normalizer.fit transform(reduced tfidf)
# Load Sentence Transformer model
sentence model = SentenceTransformer('paraphrase-multilingual-MiniLM-
L12-v2')
sentence embeddings =
sentence model.encode(train data['text'].tolist())
# Normalize Sentence Embeddings
sentence normalizer = Normalizer(norm='l2')
normalized train sentences =
sentence_normalizer.fit_transform(sentence_embeddings)
# Combine features (Normalized TF-IDF + Normalized Sentence
```

```
Embeddings)
X combined = np.hstack([
    normalized train tfidf,
                                         # Normalized TF-IDF features
(300 dims)
    normalized train sentences
                                        # Normalized Sentence
Embeddings (84 dims)
# Map labels
label_map = {label: idx for idx, label in
enumerate(train data['subreddit'].unique())}
y = train data['subreddit'].map(label map)
# Hyperparameter grids
param grid nb = {
    'alpha': [0.01, 0.1, 1.0]
param grid lr = {
    'C': [0.1, 1, 10],
    'solver': ['lbfgs']
}
param grid_cb = {
    'iterations': [100, 200],
    'learning rate': [0.05],
    'depth': [4, 6]
}
# Multinomial Naive Bayes
nb model = MultinomialNB()
grid search nb = GridSearchCV(
    nb model, param grid=param grid nb, cv=3, scoring='accuracy',
verbose=1, n jobs=-1
grid search nb.fit(non negative tfidf, y)
# Logistic Regression
lr model = LogisticRegression(max iter=1000)
grid search lr = GridSearchCV(
    lr model, param grid=param grid lr, cv=3, scoring='accuracy',
verbose=1, n jobs=-1
grid search lr.fit(X combined, y)
# CatBoost
cb model = CatBoostClassifier(verbose=0)
grid search cb = GridSearchCV(
    cb_model, param_grid=param_grid_cb, cv=3, scoring='accuracy',
verbose=1, n jobs=-1
```

```
grid search cb.fit(X combined, y)
# Save best estimators
nb best model = grid search nb.best estimator
lr best model = grid_search_lr.best_estimator_
cb_best_model = grid_search_cb.best_estimator_
# Metrics
metrics = {
    'Classifier': ['Multinomial Naive Bayes', 'Logistic Regression',
'CatBoost'],
    'Training Accuracy': [
        grid search nb.best score ,
        grid search lr.best_score_,
        grid search cb.best score
metrics df = pd.DataFrame(metrics)
print(metrics df)
# Process test set: TF-IDF
test tfidf features = tfidf vectorizer.transform(test data['body'])
test reduced tfidf = svd.transform(test tfidf features) # Reduce
dimensions to 300
test normalized tfidf = tfidf normalizer.transform(test reduced tfidf)
# Normalize TF-IDF
# Process test set: Sentence Embeddings
test sentence embeddings =
sentence model.encode(test data['body'].tolist())
test normalized sentence embeddings =
sentence_normalizer.transform(test sentence embeddings) # Normalize
Sentence Embeddings
# Combine test features (TF-IDF + Sentence Embeddings)
test combined = np.hstack([
   test normalized tfidf,
                               # Normalized TF-IDF features
(300 dims)
   test normalized sentence embeddings # Normalized Sentence
Embeddings (84 dims)
1)
# Predict on test set using best CatBoost model
test predictions = cb best model.predict(test combined)
# Map predictions back to labels
reverse label map = {idx: label for label, idx in label map.items()}
# Flatten predictions and map back to labels
```

```
test predictions = test predictions.flatten() if
len(test predictions.shape) > 1 else test predictions
test_data['subreddit'] = [reverse_label_map[int(pred)] for pred in
test predictions]
# Create submission file
submission = test_data[['id', 'subreddit']]
submission.to csv(output file, index=False)
print(f"Submission file saved as: {output file}")
Fitting 3 folds for each of 3 candidates, totalling 9 fits
Fitting 3 folds for each of 3 candidates, totalling 9 fits
Fitting 3 folds for each of 4 candidates, totalling 12 fits
                Classifier Training Accuracy
  Multinomial Naive Bayes
                                     0.589274
1
       Logistic Regression
                                     0.731398
2
                  CatBoost
                                     0.684257
Submission file saved as: submissions.csv
```