F78DS - Data Science Life Cycle

Coursework 2

Arthur Grossmann Le Mauguen

H00494101

1st of April 2025

Introduction

1) Start with an introduction to the coursework.

In this coursework, we explore a dataset containing numerical characteristics of essays, aiming to uncover patterns and correlations between different textual features and their assigned scores. The dataset, "F78DS-Essay-Features.csv", provides various linguistic and structural attributes of essays, including word count, sentence structure, punctuation usage, and lexical diversity.

Our objective is to analyze these features to identify the most influential factors in determining essay quality. This will involve statistical analysis, data visualization, and predictive modeling to assess how different textual components contribute to the verall scoring of essays.

Directory Structure:

To facilitate the reproducibility and organization of the project, the following directory structure is recommended:

```
import os

# Define the folder names
folders = ['data', 'models']

# Create the folders if they do not exist
for folder in folders:
    if not os.path.exists(folder):
        os.makedirs(folder)
        print(f"Folder '{folder}' created.")
    else:
        print(f"Folder '{folder}' already exists.")
Folder 'data' already exists.
```

2) Importing the necessary libraries, read the file ('F78DS-Essay-Features.csv'), and provide some description of the data you have read (you do not need to repeat the description given in this file for each field).

```
In [49]: # Library import to read the data from the csv file
import pandas as pd

In [50]: # Store the data from the csv file into a pandas dataframe
df_Essay=pd.read_csv('data/F78DS-Essay-Features.csv')
```

To better understand the dataset we will use the following:

Folder 'models' created.

- describe(): Summarizes statistics (count, mean, min, max, quartiles) for numeric columns.
- head(): Displays the first 5 rows of the DataFrame
- dtypes: Displays the data type of each column in the Dataframe
- columns: Lists the column names of the Dataframe
- shape: Displays the dimensions of the Dataframe (rows, columns)

```
In [51]: df_Essay.describe()
```

ut[51]:		essayid	chars	words	commas	apostrophes	punctuations	avg_word_length	sentences	questions a	vg_word_sentence	
,	count	1332.00000	1332.000000	1332.000000	1332.000000	1332.000000	1332.00000	1332.000000	1332.000000	1332.000000	1332.000000	1
	mean	905.27027	2101.745495	424.485736	14.667417	8.141141	0.47973	4.939762	19.704204	1.222973	23.884687	
	std	526.68760	865.963750	171.873730	10.920781	6.124520	1.27168	0.231071	19.202731	1.847446	11.160020	
	min	0.00000	169.000000	36.000000	0.000000	2.000000	0.00000	2.231322	0.000000	0.000000	1.084112	
	25%	442.75000	1527.250000	310.000000	7.000000	4.000000	0.00000	4.791679	13.000000	0.000000	19.142857	
	50%	914.50000	2029.500000	411.000000	13.000000	6.000000	0.00000	4.946059	18.000000	1.000000	22.030331	
	75%	1369.75000	2613.500000	525.000000	21.000000	11.000000	0.00000	5.092938	24.000000	2.000000	26.048234	
	max	1799.00000	6142.000000	1170.000000	72.000000	51.000000	26.00000	5.681429	642.000000	17.000000	303.000000	1
n [52]:	1.0 -											
out[52]:	_	say.head() ayid chars	words comm	nas apostrop	hes punctuat	tions avg_wo	rd_length sent	tences questions	avg_word_sen	tence P(OS POS/total_wor	rds
	ess		words comm	nas apostrop	hes punctuat	tions avg_wo	rd_length sent 5.053991	tences questions 16 0		tence P(25000 423.9952		
	ess	ayid chars			-				26.6		72 0.9952	294
	ess	ayid chars 1457 2153	426	14	6	0	5.053991	16 0	26.5 ₋	25000 423.9952	72 0.9952 03 0.9965	.94 .52
	ess. 0 1	ayid chars 1457 2153 503 1480	426 292	14	6	0 0	5.053991 5.068493	16 0 11 0	26.6 26.5 17.3	25000 423.9952 45455 290.9931	72 0.9952 03 0.9965 44 0.9941	294 552 00
	ess. 0 1 2 3	ayid chars 1457 2153 503 1480 253 3964	426 292 849	14 9 19	6 7 26	0 0 1	5.053991 5.068493 4.669022	16 0 11 0 49 2	26.6. 26.5. 17.3.	25000 423.9952 45455 290.9931 26531 843.9905	72 0.9952 03 0.9965 44 0.9941 84 0.9888	294 552 00 328
	ess. 0 1 2 3	ayid chars 1457 2153 503 1480 253 3964 107 988	426 292 849 210	14 9 19 8	6 7 26 7	0 0 1 0	5.053991 5.068493 4.669022 4.704762	16 0 11 0 49 2 12 0	26.6. 26.5. 17.3.	25000 423.9952 45455 290.9931 26531 843.9905 00000 207.6537	72 0.9952 03 0.9965 44 0.9941 84 0.9888	294 552 00 328

```
essavid
                                          int64
Out[53]:
          chars
                                          int64
          words
                                          int64
          commas
                                          int64
          apostrophes
                                          int64
                                          int64
          punctuations
          avg word length
                                        float64
          sentences
                                          int64
          questions
                                          int64
                                        float64
          avg word sentence
          POS
                                        float64
          POS/total words
                                        float64
                                          int64
          prompt words
          prompt words/total words
                                        float64
          synonym words
                                          int64
          synonym_words/total words
                                        float64
          unstemmed
                                          int64
          stemmed
                                          int64
          score
                                          int64
          dtype: object
In [54]:
          df Essay.columns
          Index(['essayid', 'chars', 'words', 'commas', 'apostrophes', 'punctuations',
Out[54]:
                 'avg word length', 'sentences', 'questions', 'avg word sentence', 'POS',
                 'POS/total words', 'prompt_words', 'prompt_words/total_words',
                 'synonym words', 'synonym words/total words', 'unstemmed', 'stemmed',
                 'score'],
                dtype='object')
          df Essay.shape
In [55]:
          (1332, 19)
Out[55]:
```

- The dataset contains 1,322 essays.
- Essays vary significantly in length, from as few as 169 characters (36 words) to as many as 6142 characters (1,170 words).
- The average word count is approximately **424 words**, with a maximum of **1170 words**.
- Essays generally contain few questions, with an average of **1.22 questions per essay**.
- The number of Part-of-Speech (POS) elements is closely related to the total word count, indicating a rich linguistic structure in longer essays.

• The score distribution ranges **from 1 to 6**, with an average score of **3.42** and a median at **3** (more than half of the students have a grade of 3 or more, this students have passing grades in Heriot Watt University)

2) Supervised Learning

1) Explain supervised machine learning, the notion of labelled data, and the training and test datasets.

Supervised machine learning is a type of machine learning in which an algorithm learns to associate or map input data (labelled data, see next section for definition) with corresponding outputs. By identifying patterns in the provided examples, the model can then make **predictions** or **classify** new data points. The objective is to develop a model capable of generalizing effectively to unseen data by identifying **patterns** from past examples. This contrasts with unsupervised learning, where the model independently identifies features and structures the data.

In supervised learning, there are two main types of problems:

- Regression: Predicting a continuous value (e.g., predicting house prices based on square footage, grades based on the length of an essay).
- Classification: Assigning data points to categories or classes (e.g., determining if an email is spam or not, whether a picture features a cat or a dog).

Labelled Data

Labelled data is a dataset where each input example (feature) is paired with a corresponding output (label or target variable). Label refers to the expected output, what the model should predict using the features. This allows the unsupervised machine learning model to learn relationships between input features and their correct outputs.

In the F78DS-Essay-Features.csv dataset, each essay has numerical features (e.g., word count, characters count) as inputs and a score as the output label. The model can learn from these examples to predict the score of new essays.

Training and Test Datasets

When building a supervised machine learning model, the dataset is typically divided into two subsets:

Training Dataset

- Used to train the model by providing examples with known outputs.
- The model learns patterns between features and labels in the data and adjusts its internal parameters.

Test Dataset

- Used to evaluate the model's performance on unseen data.
- Helps check if the model generalizes well (how well it can predict labels of data it has not seen before) and avoids overfitting (memorizing training data instead of learning general patterns).
- In some cases, a validation dataset is also used to fine-tune the model by opimizing the hyperparameters before final testing(it's a separate subset of data used during training to fine-tune the model and prevent overfitting).

Example in the Context of Essay Scoring

- Training Data: A subset of essays with their features (e.g., word count, punctuation, sentence structure) and known scores.
- Test Data: A different subset of essays with features but without exposing the scores to the model. The model predicts scores, which are then compared to actual scores to measure accuracy.

By using this approach, we can develop a machine learning model that predicts essay scores based on textual features.

2) Separate the features and the label (Hint: the label in this case is the 'score')

```
In [56]: # Separate features (X) and Label (y)
X = df_Essay.drop(columns=['score']) # Features (all columns except 'score')
y = df_Essay['score'] # Label (target variable)

# Display the first few rows of features and Labels
X.head(), y.head()
```

```
essayid
                       chars words
                                      commas
                                               apostrophes
                                                            punctuations avg word length \
Out[56]:
                 1457
                         2153
                                 426
                                           14
                                                         6
                                                                                   5.053991
                                           9
                                                         7
           1
                  503
                        1480
                                 292
                                                                        0
                                                                                  5.068493
           2
                  253
                         3964
                                 849
                                           19
                                                        26
                                                                        1
                                                                                  4.669022
           3
                  107
                          988
                                 210
                                           8
                                                         7
                                                                        0
                                                                                  4.704762
           4
                 1450
                         3139
                                 600
                                           13
                                                         8
                                                                                  5.231667
                          questions avg word sentence
                                                                 POS
                                                                      POS/total words \
              sentences
           0
                                  0
                                              26.625000
                                                         423.995272
                                                                             0.995294
                     16
           1
                     11
                                  0
                                              26.545455
                                                         290.993103
                                                                             0.996552
           2
                     49
                                  2
                                             17.326531
                                                         843.990544
                                                                             0.994100
           3
                     12
                                  0
                                                                             0.988828
                                             17.500000
                                                         207.653784
                      24
                                  1
           4
                                              25.000000
                                                         594.652150
                                                                             0.991087
                             prompt words/total words
              prompt words
                                                        synonym words \
           0
                        207
                                              0.485915
                                                                   105
           1
                                                                   77
                        148
                                              0.506849
           2
                        285
                                             0.335689
                                                                   130
           3
                        112
                                              0.533333
                                                                    62
                        255
           4
                                              0.425000
                                                                   165
              synonym words/total words unstemmed
           0
                                0.246479
                                                 424
                                                          412
           1
                                0.263699
                                                 356
                                                          345
           2
                                                          750
                                0.153121
                                                 750
           3
                                0.295238
                                                 217
                                                          209
           4
                                0.275000
                                                 702
                                                          677,
           0
                4
           1
                4
           2
                4
           3
                3
           4
```

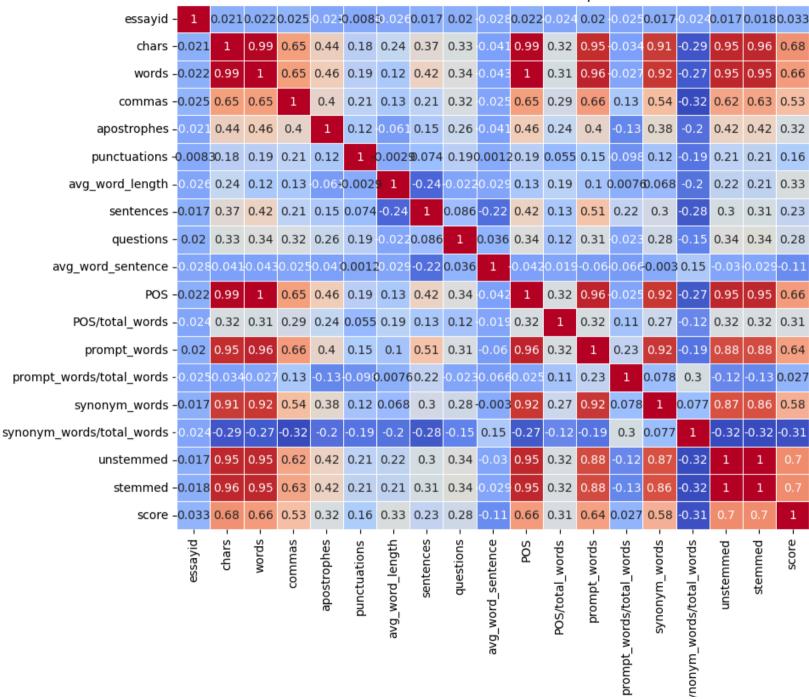
Name: score, dtype: int64)

3) Conduct feature selection (if appropriate) to improve the model performance

Feature selection is a crucial step in machine learning that helps improve model performance by identifying and retaining the most relevant features while reducing dimensionality and eliminating redundant or irrelevant data. One common approach is using a correlation matrix, which measures the relationships between features. We might need to fine-tune our features choice to improve accuracy.

```
In [57]: # Library import to conduct feature selection
          import numpy as np
          import seaborn as sns
          import matplotlib.pyplot as plt
          from sklearn.ensemble import RandomForestClassifier
         from sklearn.feature_selection import VarianceThreshold
          # 1. Correlation Analysis (Heatmap)
          plt.figure(figsize=(12, 8))
          heatmap = df Essay.corr()
          #sns.heatmap(heatmap, annot=False, cmap="coolwarm", linewidths=0.5)
         sns.heatmap(heatmap, annot=True, cmap="coolwarm", linewidths=0.5) # Display correlation values on heatmap with annot=True to disp
          plt.title("Feature Correlation Heatmap")
          plt.show()
          # 2. Variance Threshold - Remove Low variance features
          selector = VarianceThreshold(threshold=0.01) # Removing near-constant features
         X high variance = selector.fit transform(X)
          # Get selected feature names
          selected features = X.columns[selector.get support()]
          # 3. Feature Importance using RandomForest
          rf = RandomForestClassifier(n estimators=100, random state=42)
         rf.fit(X[selected features], y)
          # Get feature importances
          feature importances = pd.Series(rf.feature importances , index=selected features)
          top features = feature importances.sort values(ascending=False)
          # Plot Top Features
          plt.figure(figsize=(10, 6))
          top features.plot(kind='bar')
         plt.title("Feature Importance (Random Forest)")
          plt.xlabel("Features")
          plt.ylabel("Importance Score")
          plt.show()
          # Display top important features
         top features.head(10)
```

Feature Correlation Heatmap



1.0

- 0.8

- 0.6

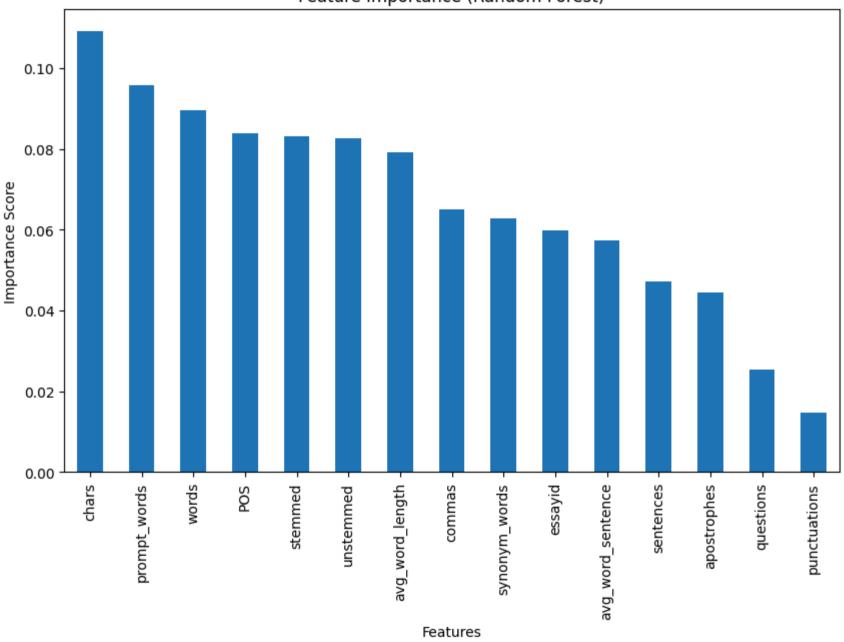
- 0.4

- 0.2

- 0.0

- -0.2

Feature Importance (Random Forest)



```
0.109018
          chars
Out[57]:
          prompt words
                             0.095753
          words
                             0.089546
          POS
                             0.083922
                             0.083229
          stemmed
                             0.082740
          unstemmed
          avg word length
                             0.079050
                             0.064929
          commas
                             0.062837
          synonym words
          essavid
                             0.059921
          dtype: float64
In [58]: # Define selected features (excluding essayid as it's just an identifier)
          selected features = ["chars", "prompt words", "words", "POS", "stemmed", "unstemmed", "avg word length", "commas", "synonym words
```

4) Use the sklearn.model_selection.train_test_split function to split your data for training and testing. Explain the parameters used in the function.

```
In [59]: # Library import to split the data into training and testing sets
from sklearn.model_selection import train_test_split

# Split the data into training and testing sets (80% train, 20% test)
X_train, X_test, y_train, y_test = train_test_split(X[selected_features], y, test_size=0.2, random_state=0, stratify=y)

# Display the shape of training and testing sets to verify the split
X_train.shape, X_test.shape, y_train.shape, y_test.shape

Out[59]: ((1065, 9), (267, 9), (1065,), (267,))
```

We used the train_test_split function from sklearn.model_selection to divide the dataset into training and testing subsets. The parameters used are:

- X[selected_features]: The selected features from the dataset, excluding irrelevant columns.
- y: The target variable (score).
- test_size=0.2: 20% of the data is allocated to the test set, while 80% is used for training.
- random_state=0: Ensures reproducibility by setting a fixed seed for random shuffling.
- stratify=y: Maintains the distribution of score across train and test sets, ensuring balanced representation.
- Dataset Split Summary
- Training Set: 1065 samples (80%)

- Testing Set: 267 samples (20%)
- Features Used: 9 (top 10 most relevant excluding essayid)

3) Classification

1) Explain the difference between binary and multi-class classification.

Binary Classification

Binary classification involves predicting one of two possible classes (labels). The output is typically represented as 0 or 1, True or False, Positive or Negative.

Examples:

- Spam detection: Email is either "Spam" (1) or "Not Spam" (0).
- Medical diagnosis: A test result is "Positive" (disease detected) or "Negative" (no disease).
- Sentiment analysis: A review is classified as "Positive" or "Negative".

Multi-Class Classification

Multi-class classification involves predicting one class out of three or more possible classes. Unlike binary classification, there are multiple categories, and each sample belongs to exactly one class.

Examples:

- Handwritten digit recognition: Classifying images into digits 0-9.
- Essay scoring (like our coursework): Predicting scores 1 to 6.
- Animal classification: Identifying if an image is a cat, dog, bird, or horse.

In our dataset, the "score" column ranges from 1 to 6, meaning we are dealing with a multi-class classification problem.

- 2) In preparation for Naive Bayes (NB) classification, your data should be normalised/scaled.
- 1) Describe what you understand from this need to normalise data (this is in your Week 7 laboratory).

Normalization (or scaling) is a crucial preprocessing step in machine learning that ensures all features contribute equally to the model. It is especially important for models that rely on probabilities, like Naïve Bayes (NB), or those that are sensitive to differences in feature magnitudes.

Features in datasets often have different ranges. For example, the number of characters in an essay could be in the thousands, while the average word length might be between 3 and 10. If left unscaled, features with larger numerical ranges may dominate the model, leading to biased learning.

Many machine learning algorithms, including Naïve Bayes, assume that features follow a similar scale for better probability calculations. Some distance-based models (like KNN and SVM) also perform better with normalized data. Stabilizes Numerical Computations

Common Normalization Techniques:

• Min-Max Scaling (Normalization)

Rescales values to a fixed range, usually [0,1].

Formula: $X' = (X - X \min)/(X \max - X \min)$

Best when feature distributions are not Gaussian (e.g., skewed data).

• Z-score Standardization (Mean Normalization)

Centers data around mean = 0 and scales it to unit variance.

Formula: $X' = (X - \mu)/\sigma$

Ideal for normally distributed (Gaussian) data.

• <u>Log Transformation</u>

Useful for skewed data by reducing the impact of large values.

Formula: $X' = \log(1+X)$

Often used in text-based data, like word counts.

Robust Scaling

Scales data using the median and interquartile range (IQR), reducing the impact of outliers.

2) Choose and use the appropriate normalisation functions available from sklearn.preprocessing and scale the data appropriately.

In []: # Library import to scale the data from sklearn.preprocessing, numpy for log transformation
from sklearn.preprocessing import MinMaxScaler, StandardScaler, QuantileTransformer, RobustScaler, Normalizer

```
import numpy as np
# Creation of all the scalers used for the transformation of the data
min max scaler = MinMaxScaler()
standard scaler = StandardScaler()
quantile transformer = QuantileTransformer(random state=0)
robust scaler = RobustScaler()
normalizer scaler = Normalizer()
# MinMaxScaler
X train min max = min max scaler.fit transform(X train)
X test min max = min max scaler.transform(X test)
# StandardScaler
X train standard = standard scaler.fit transform(X train)
X test standard = standard scaler.transform(X test)
# QuantileTransformer
X train quantile = quantile transformer.fit transform(X train)
X test quantile = quantile transformer.transform(X test)
# RobustScaler
X train robust = robust_scaler.fit_transform(X_train)
X test robust = robust scaler.transform(X test)
# Normalizer
X train norm = normalizer scaler.fit transform(X train)
X test norm = normalizer scaler.transform(X test)
# Log transformation
X train log = np.log1p(X train) # log1p(x) is equivalent to log(x + 1)
X test log = np.log1p(X test)
# OuantileTransformer (Model Accuracy: 0.4232)
normalised X train = X train quantile
normalised X test = X test quantile
# MinMaxScaler (Model Accuracy: 0.5243)
normalised X train = X train min max
normalised X test = X test min max
# StandardScaler (Model Accuracy: 0.5243)
normalised X train = X train standard
normalised X test = X test standard
# RobustScaler (Model Accuracy: 0.5243)
normalised X train = X train robust
normalised X test = X test robust
# Best results with the following scaler:
# Normalizer (Model Accuracy: 0.6030)
normalised X train = X train norm
```

```
normalised_X_test = X_test_norm
"""
# Log transformation (Model Accuracy: 0.5730)
normalised_X_train = X_train_log
normalised_X_test = X_test_log
"""
```

3) Use the Naive Bayes (NB) algorithm to build the model.

1) Describe NB. Again, this is not in your lecture content, you need to do some self-learning.

Naïve Bayes (NB) is a probabilistic classification algorithm based on Bayes' Theorem and conditionnal probabilities, assuming that features are conditionally independent given the class label, meaning each feature contributes to the classification decision without depending on the others. Despite this strong assumption, which gives the algorithm its "naïve" name, it performs well in many practical applications, especially in text classification and natural language processing (NLP) by simplifying computations which makes it highly efficient for large datasets.

Bayes' Theorem The core of NB is Bayes' Theorem: P(AIB)=(P(BIA)xP(A))/P(B)

Where:

- P(AIB) is the probability of class A given feature B (posterior probability).
- P(BIA) is the probability of feature B given class A (likelihood).
- P(A) is the prior probability of class A.
- P(B) is the prior probability of feature B.

2) Explain the differences between the 4 models for Naive Bayes provided by sklearn.naive_bayes. From your understanding, which model would be most appropriate for this coursework and state clearly why.

Types of Naïve Bayes Models in sklearn.naive_bayes Scikit-learn provides four types of Naïve Bayes classifiers:

Model	Description	Suitable for
GaussianNB	Assumes that features follow a normal (Gaussian) distribution. It is suitable for discrete features, like word frequencies in text classification	Continuous numeric data (e.g., height, weight, scores).
MultinomialNB	Works with discrete count data, assuming features follow a multinomial distribution.	Text classification (word frequency, term counts).

Model	Description	Suitable for
BernoulliNB	Suitable for binary/boolean features, where each feature is either 0 or 1.	Binary text features (e.g., presence/absence of words).
ComplementNB	Variant of MultinomialNB, designed to handle imbalanced data.	Imbalanced text classification tasks.

Since our dataset contains numerical continuous features (e.g., chars, words, sentences), the most appropriate choice is Gaussian Naïve Bayes (GaussianNB), as it assumes a normal distribution for numerical data.

3) Write the code to build the model using your training dataset.

4) Predict

1) Using the testing dataset you created in 2(c) above, conduct the prediction for the 'score' (label).

```
In [62]: # Predict on the test set
    y_pred = nb_classifier.predict(normalised_X_test)

# Evaluate the model
    accuracy = accuracy_score(y_test, y_pred)
    print(f"Model Accuracy: {accuracy:.4f}")

# Display classification report
    print(classification_report(y_test, y_pred))
```

Model Accurac	y: 0.5880 precision	recall	f1-score	support
	precision	rccair	11 30010	заррог с
1	0.29	0.67	0.40	3
2	0.58	0.32	0.41	22
3	0.64	0.60	0.62	112
4	0.62	0.68	0.65	117
5	0.14	0.17	0.15	12
6	0.00	0.00	0.00	1
accuracy			0.59	267
macro avg	0.38	0.40	0.37	267
weighted avg	0.60	0.59	0.59	267

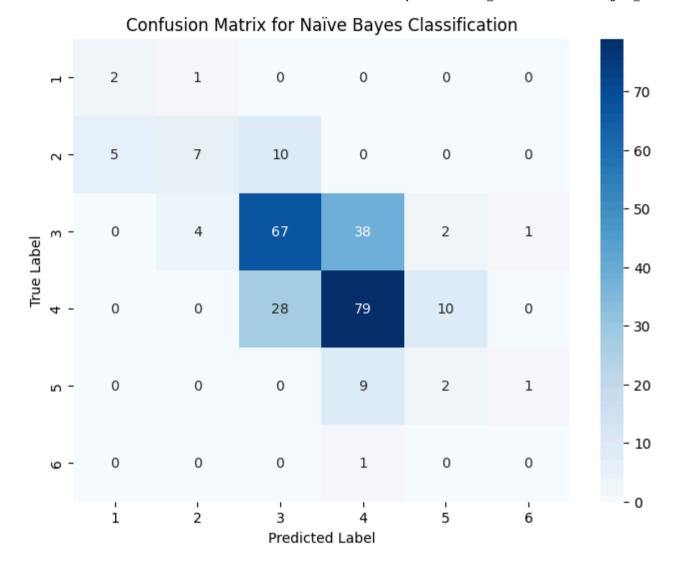
2) Display the confusion matrix (it should look like a 6x6 matrix). Unlike the lectures, where it is just a 2x2, you are now introduced to a multi-class classification.

```
In [63]: # Library import for confusion matrix
from sklearn.metrics import confusion_matrix
import seaborn as sns
import matplotlib.pyplot as plt

# Make predictions using the trained Naïve Bayes model
y_pred = nb_classifier.predict(normalised_X_test)

# Compute the confusion matrix
conf_matrix = confusion_matrix(y_test, y_pred)

# Plot the confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix, annot=True, fmt='d', cmap='Blues', xticklabels=range(1, 7), yticklabels=range(1, 7))
plt.xlabel("Predicted_Label")
plt.ylabel("True_Label")
plt.title("Confusion Matrix for Naïve Bayes Classification")
plt.show()
```



Understanding the Confusion Matrix

- Rows represent the true classes (actual labels)
- Columns represent the predicted classes (model predictions)
- Diagonal elements represent correct classifications (true positives)
- Off-diagonal elements indicate misclassifications

Class-wise Breakdown

Class 1

Correctly classified: 2 instances.

Misclassified as:

• Class 2 (1 instance).

Observation: Very few samples in this class, which could lead to an imbalanced classification issue.

Class 2

Correctly classified: 7 instances.

Misclassified as:

- Class 1 (5 instances).
- Class 3 (10 instances).

Observation: The model struggles to distinguish between Class 1, Class 2 and Class 3.

Class 3

Correctly classified: 67 instances.

Misclassified as:

- Class 2 (4 instances).
- Class 4 (38 instances) → Major confusion.
- Class 5 (2 instances).
- Class 6 (1 instance).

Observation: Heavy misclassification with Class 3, indicating overlapping features.

Class 4

Correctly classified: 79 instances.

Misclassified as:

- Class 3 (28 instances).
- Class 5 (10 instances).

Observation: Some confusion with Class 3, but overall good classification.

Class 5

Correctly classified: 2 instances.

Misclassified as:

- Class 4 (9 instances).
- Class 6 (1 instances).

Observation: More samples were misclassified than correctly classified, indicating poor separation.

Class 6

Correctly classified: 0 instances.

Misclassified as:

• Class 4 (1 instance).

Observation: This class has very few data points, likely causing misclassification issues.

3) Explain Quadratic Weighted Kappa (QWK). Again, this is not in your lectures.

Quadratic Weighted Kappa (QWK) is a statistical metric that measures the agreement between two raters (e.g., a human evaluator and an AI model) while accounting for the degree of disagreement. It is widely used in ordinal classification tasks, such as essay scoring, medical diagnoses, and rating predictions, where the order of classes matters.

Why Use QWK Instead of Accuracy?

- Accuracy does not consider how far off the prediction is.
- QWK penalizes predictions more severely when they are further from the true label.
- This makes QWK a better metric for evaluating models in ordinal classification problems (e.g., rating scales 1-6).

How QWK Works?

QWK compares the observed agreement between predictions and true labels with the expected agreement if both were random guesses. The formula is: $\kappa-1=(\sum i,j\ Wi,j\ Oi,j)/(\sum i,j\ Wi,j\ Ei,j)$

Where:

- 0i,j is the observed frequency of class i being predicted as class j.
- $Ei_i j$ is the expected frequency if predictions were random.
- Wi_i is the quadratic weight, which penalizes larger disagreements more than smaller ones.

The weight matrix W is defined as: $Wi_{,j} = (i-j)2/(N-1)2$

Where:

- *i* and *j* are the true and predicted class labels.
- *N* is the number of possible classes.

Interpreting QWK Scores

- κ =1: Perfect agreement (model predicts exactly like a human rater).
- κ =0: Model performs no better than random chance.
- κ <0: Model is worse than random guessing.

QWK Score	Agreement Level
0.81 – 1.00	Almost perfect
0.61 – 0.80	Substantial
0.41 – 0.60	Moderate
0.21 – 0.40	Fair

QWK Score	Agreement Level
0.00 - 0.20	Slight
< 0.00	Poor (worse than random)

Why is QWK Useful for This Coursework?

Since we are predicting essay scores (1-6), which follow an ordinal scale, QWK is a much better metric than simple accuracy. It ensures that a prediction of "5" instead of "6" is penalized less than a prediction of "1" instead of "6".

4) Use the sklearn.metrics library to code and obtain the QWK score.

```
In [64]: # Library import for Quadratic Weighted Kappa (QWK) score
    from sklearn.metrics import cohen_kappa_score

# Compute the Quadratic Weighted Kappa (QWK) score
    qwk_score = cohen_kappa_score(y_test, y_pred, weights="quadratic")

# Display the QWK score
    qwk_score
Out[64]: np.float64(0.6171200162596119)
```

- 5) Use a more complex algorithm, Random Forest, to build an alternative model.
- 1) Conduct steps 3.3 and 3.4 above to obtain an alternative model's results.

```
In [69]: # Load the dataset again
file_path = "data/F78DS-Essay-Features.csv"
df = pd.read_csv(file_path)

# Separate features (X) and the Label (y)
X = df.drop(columns=["score", "essayid"]) # Features (excluding essayid and score)
y = df["score"] # Label

# Train the Random Forest model
rf_classifier = RandomForestClassifier(n_estimators=1000, random_state=0)
rf_classifier.fit(normalised_X_train, y_train)
```

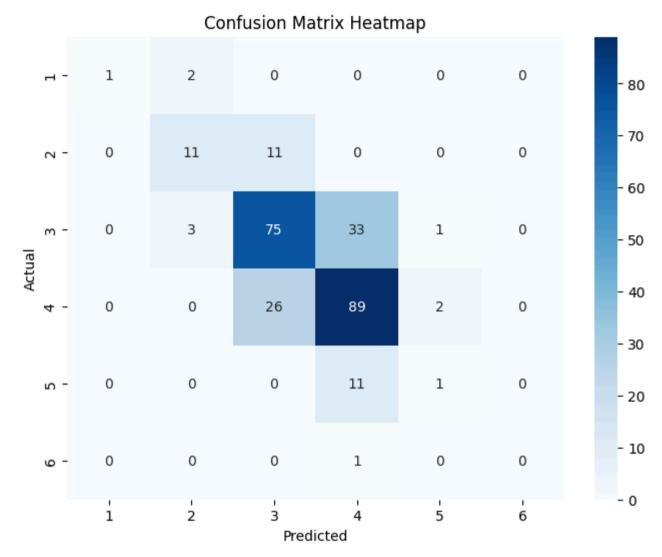
```
# Make predictions
y_pred_rf = rf_classifier.predict(normalised_X_test)

# Compute confusion matrix
conf_matrix_rf = confusion_matrix(y_test, y_pred_rf)

# Compute QWK score
qwk_score_rf = cohen_kappa_score(y_test, y_pred_rf, weights="quadratic")

# Plot heatmap of confusion matrix
plt.figure(figsize=(8, 6))
sns.heatmap(conf_matrix_rf, annot=True, fmt='d', cmap='Blues', xticklabels=sorted(y.unique()), yticklabels=sorted(y.unique()))
plt.xlabel("Predicted")
plt.ylabel("Actual")
plt.title("Confusion Matrix Heatmap")
plt.show()

# Return results
qwk_score_rf
```



Out[69]: np.float64(0.6447795116272623)

Class-wise Breakdown

Class 1

Correctly classified: 1 instance.

Misclassified as:

• Class 2 (2 instances)

Observation: Very few samples in this class, which could lead to an imbalanced classification issue.

Class 2

Correctly classified: 11 instances.

Misclassified as:

• Class 3 (11 instances)

Observation: The model struggles to distinguish between Class 2 and Class 3.

Class 3

Correctly classified: 75 instances.

Misclassified as:

- Class 2 (3 instances).
- Class 4 (33 instances) → Major confusion.
- Class 5 (1 instance).

Observation: Heavy misclassification with Class 3, indicating overlapping features with Class 4.

Class 4

Correctly classified: 89 instances.

Misclassified as:

- Class 3 (26 instances).
- Class 5 (2 instances).

Observation: Some confusion with Class 3, but overall good classification.

Class 5

Correctly classified: 1 instance.

Misclassified as:

• Class 4 (11 instances).

Observation: More samples were misclassified than correctly classified, indicating poor separation.

Class 6

Correctly classified: 0 instances.

Misclassified as:

Class 4 (1 instance).

Observation: This class has very few data points, likely causing misclassification issues.

2) Briefly explain why this alternate model performs worse or better than the Naive Bayes model.

The Random Forest model performed better than the Naïve Bayes model. Here's why:

Random Forest Captures Complex Relationships, it is an ensemble learning method that builds multiple decision trees and aggregates their results, making it better suited for handling non-linear relationships between features and the essay score.

Naïve Bayes assumes feature independence, which may not hold in this dataset, leading to lower accuracy.

The dataset contains many continuous numeric features (e.g., word count, punctuation count, POS ratios). Naïve Bayes (especially Gaussian NB) assumes normality, which is the case here. Random Forest makes no assumptions about the feature distributions, which can improve performance.

Naïve Bayes is a simpler model (faster too) and works well when feature independence holds. It is generally better for small datasets or text classification tasks where words are independent.

Random Forest is more complex and less prone to bias, but it can sometimes overfit. However, in this case, its ability to capture interactions between features gives it an advantage.

Random Forest performed better due to its ability to capture complex patterns in the data. Naïve Bayes might struggle with correlated features, leading to lower accuracy.

4) Kaggle Submission

My username on Kaggle is Arthur GLM.

- 1) Read the 'F78DS-Essay-Features-Submission.csv' file and use the model you built earlier (whichever model that you think is better) to predict the 'score'.
- 2) Unlike the previous section, you have a testing (also sometimes referred to as the validation) dataset where you know the 'score' and will be able to test for the accuracy. In this part, you don't have a 'score' and you have to predict it and submit it to the competition site. (To build a more robust model, you can do cross-validation. This in reference back to 2.3 above where you are to split your data for training and testing will leave it to you to learn about this on your own).
- 3) Output your prediction to a CSV file that contains 2 columns, 'essayid' and 'score'. It should have a total of 200 lines (1 header, and 199 entries).

First Kaggle submission with just Random Forest and without gridsearch (QWK= 0,615)

```
In [70]: # Load the submission dataset (without labels)
    submission_file_path = "data/F78DS-Essay-Features-Submission.csv"
    submission_df = pd.read_csv(submission_file_path)

# Extract the "essayid" column and the feature columns
    essay_ids = submission_df["essayid"]
    X_submission = submission_df[selected_features].copy()

# Scale the submission data using the same scaler
    X_submission_scaled = normalizer_scaler.transform(X_submission)
```

Out[70]:

```
# Predict the scores using the trained Random Forest model
predicted_scores = rf_classifier.predict(X_submission_scaled)

# Create the output DataFrame
output_df = pd.DataFrame({"essayid": essay_ids, "score": predicted_scores})

# Save to CSV file
output_csv_path = "data\H00494101-GrossmannLeMauguenArthur-1.csv"
output_df.to_csv(output_csv_path, index=False)

# Return the output file path for user reference
output_csv_path

<>:19: SyntaxWarning: invalid escape sequence '\H'
<>:19: SyntaxWarning: invalid escape sequence '\H'
C:\Users\Framework_Arthur\AppData\Local\Temp\ipykernel_28232\2830947919.py:19: SyntaxWarning: invalid escape sequence '\H'
output_csv_path = "data\H00494101-GrossmannLeMauguenArthur-1.csv"
'data\\H00494101-GrossmannLeMauguenArthur-1.csv'
```

Second Kaggle submission with just Random Forest and with gridsearch (QWK= 0,656)

[!!may take long to execute on your computer!! (1h+ on mine)]

Grid Search is a hyperparameter tuning technique used to find the best combination of hyperparameters for a machine learning model. It systematically goes through a predefined set of possible values for each hyperparameter and evaluates the model's performance for each combination.

Each hyperparameter in param_grid affects how the RandomForestClassifier learns:

- n_estimators: [100, 200, 300]

 This represents the number of decision trees in the forest. More trees usually improve performance but increase computation time.
- max_depth: [None, 1, 10, 20]
 Controls the maximum depth of each tree.
 None allows trees to grow fully, while 1, 10, 20 test different levels of restriction.
- min_samples_split: [2, 4, 6, 8]
 Minimum number of samples required to split an internal node.

Higher values make the trees less complex to avoid overfitting.

- min_samples_leaf: [1, 2, 3, 4]
 - Minimum number of samples required to be in a leaf node. Helps prevent the model from learning too specific patterns.
- max_features: ['sqrt', 'log2', None]
 Number of features considered when splitting a node.
 'sqrt' and 'log2' are commonly used to balance model performance and efficiency.
- bootstrap: [True, False]
 If True, random sampling with replacement is used (default in RandomForest).
 If False, all samples are used for training.
- class_weight: [None, 'balanced', 'balanced_subsample']
 Adjusts weights for handling class imbalance.
 'balanced' uses the inverse class frequencies, while 'balanced_subsample' does so at each tree level.
- criterion: ['gini', 'entropy']
 Defines how splits are chosen in decision trees.
 'gini' favors purity, while 'entropy' considers information gain.

Randomized Grid Search is an extension of Grid Search that samples a fixed number of hyperparameter combinations from the specified parameter grid. This approach is useful when the parameter grid is large, as it can significantly reduce computation time while still providing a good estimate of the best hyperparameters.

Cross-validation is a technique to assess the performance of a model by splitting the dataset into multiple parts, training the model on some parts, and testing it on others. It helps ensure that the model generalizes well to unseen data.

Here: cv=4 which means 4-fold cross-validation is used. The dataset is split into 4 subsets, the model is trained on 3 subsets and tested on the remaining one, and this process is repeated 4 times. The average score across the 4 iterations is used to evaluate the hyperparameter combination.

```
In [73]: from sklearn.model_selection import GridSearchCV
    from sklearn.ensemble import RandomForestClassifier
    from sklearn.metrics import cohen_kappa_score, make_scorer
    import pandas as pd
```

```
# Define QWK scoring function
def quadratic weighted kappa(y true, y pred):
    return cohen kappa score(y true, y pred, weights="quadratic")
qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
# Define parameter grid for tuning
param grid = {
    'n estimators': [100, 200, 300],
    'max depth': [None, 1, 10, 20],
    'min samples split': [2, 4, 6, 8],
    'min samples leaf': [1, 2, 3, 4],
    'max features': ['sqrt', 'log2', None],
    'bootstrap': [True, False],
    'class weight': [None, 'balanced', 'balanced subsample'],
    'criterion': ['gini', 'entropy']
# Perform Grid Search with cross-validation
rf classifier = RandomForestClassifier(random state=0)
grid search = GridSearchCV(rf classifier, param grid, cv=4, scoring=qwk scorer, n jobs=-1, verbose=2)
grid search.fit(normalised X train, y train)
# Best estimator after grid search
best rf classifier = grid search.best estimator
# Print best hyperparameters
print("Best hyperparameters:", grid search.best params )
# Load the submission dataset (without labels)
submission file path = "data/F78DS-Essay-Features-Submission.csv"
submission df = pd.read csv(submission file path)
# Extract the "essayid" column and the feature columns
essay ids = submission df["essayid"]
X submission = submission df[selected features].copy()
# Scale the submission data using the same scaler
X submission scaled = normalizer scaler.transform(X submission)
# Predict the scores using the best trained Random Forest model
predicted scores = best rf classifier.predict(X submission scaled)
```

```
# Create the output DataFrame
          output df = pd.DataFrame({"essayid": essay ids, "score": predicted scores})
          # Save to CSV file
          output csv path = "data/H00494101-GrossmannLeMauguenArthur-2.csv"
          output df.to csv(output csv path, index=False)
          # Return the output file path for user reference
          output csv path
          Fitting 4 folds for each of 6912 candidates, totalling 27648 fits
          c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model selection\ split.py:805: UserW
          arning: The least populated class in y has only 3 members, which is less than n splits=4.
            warnings.warn(
          Best hyperparameters: {'bootstrap': True, 'class weight': 'balanced subsample', 'criterion': 'entropy', 'max depth': 10, 'max fe
          atures': 'sqrt', 'min samples leaf': 4, 'min samples split': 2, 'n estimators': 200}
          'data/H00494101-GrossmannLeMauguenArthur-2.csv'
Out[73]:
          With the following hyperparameters, the QWK score on kaggle improves to 0.637
          Best hyperparameters: {
          'bootstrap': True,
          'class_weight': 'balanced_subsample',
          'criterion': 'entropy',
          'max_depth': 10,
          'max_features': 'sqrt',
          'min_samples_leaf': 4,
          'min_samples_split': 2,
          'n estimators': 200
```

During my research, I found other models and here I am trying them and comparaing the results to my random forest model

• Random Forest:

Random Forest is an ensemble learning method that constructs multiple decision trees during training and outputs the class that is the mode of the classes (classification) or mean prediction (regression) of the individual trees. It is known for its robustness and ability to handle large

datasets with higher dimensionality. Random Forest reduces overfitting by averaging multiple deep decision trees, trained on different parts of the same training set.

• Gradient Boosting:

Gradient Boosting builds an ensemble of decision trees sequentially, each new tree attempting to correct the errors of the previous ones. It optimizes a differentiable loss function, making it highly effective for both classification and regression tasks. Gradient Boosting often achieves high predictive accuracy but can be computationally intensive and prone to overfitting if not properly tuned.

XGBoost:

XGBoost, short for Extreme Gradient Boosting, is an optimized gradient boosting library designed for efficiency and performance. It uses advanced regularization techniques and parallel processing to deliver fast and accurate predictions. XGBoost is widely used in data science competitions and industry applications due to its superior handling of missing values and flexibility in customizing objective functions.

• Support Vector Machine (SVM):

SVM is a powerful supervised learning model used for classification and regression tasks. It works by finding the hyperplane that best separates the data into different classes in a high-dimensional space. SVM is effective in high-dimensional spaces and when the number of dimensions exceeds the number of samples. It uses kernel tricks to handle non-linear classification problems efficiently.

```
In [74]: from sklearn.model selection import train test split
         from sklearn.preprocessing import StandardScaler
         from sklearn.ensemble import GradientBoostingClassifier, RandomForestClassifier
         from xgboost import XGBClassifier
         from sklearn.svm import SVC
          from sklearn.metrics import accuracy score, cohen kappa score, confusion matrix
         import numpy as np
         # Load the dataset
         file path = "data/F78DS-Essay-Features.csv"
         df = pd.read csv(file path)
         # Separate features (X) and the Label (y)
         X = df.drop(columns=["score", "essayid"]) # Features (excluding essayid and score)
         y = df["score"] - 1 # Adjust labels to start from 0
         # Split into training and testing sets (80% train, 20% test)
         X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42, stratify=y)
         # Normalize the features using StandardScaler
```

```
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Train different models
models = {
    "Random Forest": RandomForestClassifier(n estimators=100, random state=42),
    "Gradient Boosting": GradientBoostingClassifier(n estimators=100, random state=42),
    "XGBoost": XGBClassifier(n estimators=100, use label encoder=False, eval metric='mlogloss'),
    "SVM": SVC(probability=True, random state=42)
# Train each model and store predictions
model scores = []
prob predictions = {}
for name, model in models.items():
    model.fit(X train scaled, y train)
    pred = model.predict(X test scaled)
    # Calculate evaluation metrics
    accuracy = accuracy score(y test, pred)
    qwk = cohen kappa score(y test, pred, weights="quadratic")
    model scores.append((name, accuracy, qwk))
    # Store probability predictions for ensemble
    prob predictions[name] = model.predict proba(X test scaled)
# Sort models by OWK score in descending order
model scores.sort(key=lambda x: x[2], reverse=True)
print("\nModel Performance Sorted by OWK Score:")
for name, accuracy, qwk in model scores:
    print(f"{name} - Accuracy: {accuracy:.4f}, QWK: {qwk:.4f}")
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\xgboost\training.py:183: UserWarning: [23:2
7:12] WARNING: C:\actions-runner\ work\xgboost\src\learner.cc:738:
Parameters: { "use label encoder" } are not used.
 bst.update(dtrain, iteration=i, fobj=obj)
```

```
Model Performance Sorted by QWK Score:

SVM - Accuracy: 0.6742, QWK: 0.6318

Random Forest - Accuracy: 0.6292, QWK: 0.6227

Gradient Boosting - Accuracy: 0.6180, QWK: 0.6106

XGBoost - Accuracy: 0.6180, QWK: 0.6046
```

Ensemble model

Ensemble modeling is a technique that combines the predictions of multiple machine learning models to improve overall performance and robustness. The principle behind ensemble modeling is that a group of "weak learners" can come together to form a "strong learner," often outperforming any individual model.

- Diversity and Reduction of Overfitting: Ensemble methods leverage the diversity among different models. By combining models that make different types of errors, the ensemble can reduce the risk of overfitting and improve generalization to unseen data.
- Combining Predictions: The predictions from multiple models are combined using techniques such as averaging (for regression tasks) or voting (for classification tasks). This combination helps to smooth out individual model errors and can lead to more stable and accurate predictions.
- Types of Ensemble Methods:
 - Bagging (Bootstrap Aggregating): Involves training multiple instances of the same model on different subsets of the training data and averaging their predictions. Random Forest is a popular example of bagging.
 - Boosting: Sequentially trains models, each trying to correct the errors of the previous ones. Models like Gradient Boosting and XGBoost use this approach.
 - Stacking: Combines the predictions of multiple models using another model or meta-learner to make the final prediction.
- Improved Performance: Ensemble models often achieve better performance than individual models by capturing different aspects of the data and reducing variance and bias. They are particularly effective in complex tasks where no single model can capture all the underlying patterns.

```
In [75]: # Ensemble Method: Average the probabilities
    avg_prob = np.mean(list(prob_predictions.values()), axis=0)
    ensemble_prediction = np.argmax(avg_prob, axis=1) # Labels are already 0-indexed

# Evaluate the ensemble model
    ensemble_accuracy = accuracy_score(y_test, ensemble_prediction)
    ensemble_qwk = cohen_kappa_score(y_test, ensemble_prediction, weights="quadratic")
```

```
print(f"\nEnsemble Model - Accuracy: {ensemble_accuracy:.4f}, QWK: {ensemble_qwk:.4f}")

# Confusion Matrix for the ensemble model
conf_matrix_ensemble = confusion_matrix(y_test, ensemble_prediction)
print("\nConfusion Matrix for Ensemble Model:\n", conf_matrix_ensemble)

Ensemble Model - Accuracy: 0.6404, QWK: 0.6256

Confusion Matrix for Ensemble Model:
[[ 1 2 0 0 0 0 0]
        [ 1 12 9 0 0 0]
        [ 0 2 83 27 0 0]
        [ 0 0 42 75 0 0]
        [ 0 0 0 12 0 0]
        [ 0 0 0 1 0 0]]
```

I will now try to improve the performance of the best model, SVM

```
In [76]: from sklearn.svm import SVC
         from sklearn.metrics import make scorer
         from sklearn.model selection import GridSearchCV
         # Define OWK scoring function
         def quadratic weighted kappa(y true, y pred):
              return cohen kappa score(y true, y pred, weights="quadratic")
         # Use make scorer to create a scorer based on QWK
         qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
         # Load the dataset
         file path = "data/F78DS-Essay-Features.csv"
         df = pd.read csv(file path)
         # Separate features (X) and the label (y)
         X = df.drop(columns=["score", "essayid"]) # Features (excluding essayid and score)
         y = df["score"] - 1 # Adjust labels to start from 0
         # Split into training and testing sets (80% train, 20% test)
         X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42, stratify=y)
         # Normalize the features
         X train scaled = normalizer scaler.fit transform(X train)
         X test scaled = normalizer scaler.transform(X test)
```

```
# Define the SVM model
svm model = SVC(probability=True, random state=42)
# Define the hyperparameter grid
param grid = {
    'C': [0.1, 1, 10, 100, 1000], # Regularization parameter
    'kernel': ['linear', 'poly', 'rbf', 'sigmoid'], # Kernel types
    'gamma': ['scale', 'auto', 0.1, 1, 10], # Kernel coefficient
    'degree': [3, 4, 5] # Degree of polynomial kernel (only for 'poly')
# Perform GridSearchCV with QWK as the scoring metric
grid search = GridSearchCV(svm model, param grid, cv=5, n jobs=-1, verbose=2, scoring=qwk scorer)
grid search.fit(X train scaled, y train)
# Get the best SVM model from grid search
best svm model = grid search.best estimator
# Evaluate the best model on the test set
best svm pred = best svm model.predict(X test scaled)
# Print the best hyperparameters and evaluation metrics
print(f"Best Hyperparameters: {grid search.best params }")
accuracy = grid search.best score # Accuracy on training data
qwk = quadratic weighted kappa(y test, best svm pred)
print(f"Test Accuracy: {accuracy:.4f}, QWK: {qwk:.4f}")
Fitting 5 folds for each of 300 candidates, totalling 1500 fits
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model selection\ split.py:805: UserW
arning: The least populated class in y has only 3 members, which is less than n splits=5.
  warnings.warn(
Best Hyperparameters: {'C': 10, 'degree': 3, 'gamma': 10, 'kernel': 'poly'}
Test Accuracy: 0.6609, QWK: 0.6579
Best Hyperparameters: {
'C': 10,
'degree': 3,
'gamma': 10,
'kernel': 'poly'
```

Test Accuracy: 0.6609, QWK: 0.6579

```
In [77]: import joblib

# Save the final model and predictions
    joblib.dump(best_svm_model, "models/best_svm_model.pkl")

Out[77]: ['models/best_svm_model.pkl']
```

A .pkl file is a pickle file, which is a serialized version of a Python object. It is created using the pickle or joblib module in Python. Serialization (or "pickling") is the process of converting a Python object (like a trained machine learning model, list, dictionary, etc.) into a byte stream that can be stored on disk and later reloaded ("unpickled") into its original form. Machine learning models (like SVMs) trained on large datasets can be saved and reused without retraining.

More fine tuning of the SVM model using the result of the first grid search

[!!may take long to execute on your computer!! (11h+ on mine)]

improvement strategies:

- Feature Engineering, using PCA (Principal Component Analysis) to reduce dimensionality retain the most informative features.
- Hyperparameter Tuning
- Handling Imbalanced Data: using the class_weight parameter in SVC to account for different class distributions.

```
In [30]: from sklearn.model_selection import GridSearchCV
    from sklearn.svm import SVC
    from sklearn.metrics import cohen_kappa_score, make_scorer
    import pandas as pd
    from sklearn.preprocessing import StandardScaler
    from sklearn.model_selection import train_test_split
    from sklearn.decomposition import PCA
    from sklearn.ensemble import VotingClassifier

# Define QWK scoring function
def quadratic_weighted_kappa(y_true, y_pred):
    return cohen_kappa_score(y_true, y_pred, weights="quadratic")
```

```
# Use make scorer to create a scorer based on OWK
qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
# Load the dataset
file path = "data/F78DS-Essay-Features.csv"
df = pd.read csv(file path)
\# Separate features (X) and the label (v)
X = df.drop(columns=["score", "essavid"]) # Features (excluding essavid and score)
v = df["score"] - 1 # Adjust labels to start from 0
# Split into training and testing sets (80% train, 20% test)
X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42, stratify=y)
# Normalize the features using StandardScaler
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Optional: Apply PCA for dimensionality reduction
pca = PCA(n components=0.95) # Retain 95% variance
X train scaled = pca.fit transform(X train scaled)
X test scaled = pca.transform(X test scaled)
# Define the SVM model
svm model = SVC(probability=True, random state=42)
# Define the hyperparameter grid
param grid = {
    'C': [0.01, 0.1, 1, 10, 100, 1000],
    'kernel': ['linear', 'poly', 'rbf', 'sigmoid'],
    'gamma': ['scale', 'auto', 0.0001, 0.001, 0.01],
    'degree': [3, 4, 5, 6],
    'class weight': [None, 'balanced']
# Perform GridSearchCV with QWK as the scoring metric
grid search = GridSearchCV(svm model, param grid, cv=10, n jobs=-1, verbose=2, scoring=qwk scorer)
grid search.fit(X train scaled, y train)
# Get the best SVM model from grid search
best svm model = grid search.best estimator
```

```
# Evaluate the best model on the test set
best svm pred = best svm model.predict(X test scaled)
# Print the best hyperparameters and evaluation metrics
print(f"Best Hyperparameters: {grid search.best params }")
accuracy = grid search.best score # Accuracy on training data
gwk = quadratic weighted kappa(v test, best svm pred)
print(f"Test Accuracy: {accuracy:.4f}, QWK: {qwk:.4f}")
# Optionally, save the final model and predictions
# best svm model.save("best svm model.pkl")
Fitting 10 folds for each of 960 candidates, totalling 9600 fits
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model selection\ split.py:805: UserW
arning: The least populated class in y has only 3 members, which is less than n splits=10.
  warnings.warn(
Best Hyperparameters: {'C': 1, 'class weight': 'balanced', 'degree': 3, 'gamma': 0.01, 'kernel': 'rbf'}
Test Accuracy: 0.6736, OWK: 0.6657
Best Hyperparameters: {
'C': 1.
'class weight': 'balanced',
```

Test Accuracy: 0.6736, QWK: 0.6657

'degree': 3,
'gamma': 0.01,
'kernel': 'rbf'

Saving the improved SVM model (Kaggle = 0.680)

```
In [100... from sklearn.decomposition import PCA

# Load the dataset
train_file_path = "data/F78DS-Essay-Features.csv"
train_df = pd.read_csv(train_file_path)

# Define features (X) and the label (y)
X = train_df.drop(columns=["score", "essayid"])
y = train_df["score"] - 1 # Ajustement de L'étiquette pour commencer à 0
```

```
# Save the features used for training
          selected features = X.columns.tolist()
          joblib.dump(selected features, "models/features.pkl")
          # Split the data into training and testing sets (80% train, 20% test)
          X train, X test, y train, y test = train test split(X, y, test size=0.2, random state=42, stratify=y)
          # Normalization of features using StandardScaler
          scaler = StandardScaler()
          X train scaled = scaler.fit transform(X train)
          X test scaled = scaler.transform(X test)
          # Dimension reduction with PCA
          pca = PCA(n components=0.95) # Keep 95% of variance
          X train scaled = pca.fit transform(X train scaled)
          X test scaled = pca.transform(X test scaled)
          # Define the best SVM model based on previous tuning
          best svm model = SVC(C=1, class weight='balanced', degree=3, gamma=0.01, kernel='rbf', random state=42)
          # Training of the best SVM model
          best svm model.fit(X train scaled, y train)
          # Testing of the best SVM model
          best svm pred = best svm model.predict(X test scaled)
          qwk = cohen kappa score(y test, best svm pred, weights="quadratic")
          # Show the results
          print(f"Meilleurs hyperparamètres : C=1, class weight='balanced', degree=3, gamma=0.01, kernel='rbf'")
          print(f"QWK sur le test : {qwk:.4f}")
          # Save the final model and scaler and PCA and features
          joblib.dump(best svm model, "models/best svm model.pkl")
          joblib.dump(scaler, "models/scaler.pkl")
          joblib.dump(pca, "models/pca.pkl")
          Meilleurs hyperparamètres: C=1, class weight='balanced', degree=3, gamma=0.01, kernel='rbf'
          OWK sur le test : 0.6657
          ['models/pca.pkl']
Out[100]:
```

Generate grade prediction for Kaggle submission using the model .pkl saved before (QWK = 0.680)

```
# Load the trained SVM model, scaler, and PCA
In [101...
          best svm model = joblib.load("models/best svm model.pkl")
          scaler = joblib.load("models/scaler.pkl")
          pca = joblib.load("models/pca.pkl")
          selected features = joblib.load("models/features.pkl")
          # Load the submission dataset
          submission file path = "data/F78DS-Essay-Features-Submission.csv"
          submission df = pd.read csv(submission file path)
          # Extract "essayid" and preprocess the features
          essay ids = submission df["essayid"]
          X submission = submission df[selected features].copy() # Ensure the same features as during training
          # Scale and apply PCA to the submission data
          X submission scaled = scaler.transform(X submission)
          X submission pca = pca.transform(X submission scaled)
          # Predict the scores using the trained SVM model
          predicted scores = best svm model.predict(X submission pca)
          # Adjust scores if necessary (e.g., rounding or shifting)
          predicted scores = predicted scores + 1 # Reverse the previous label shift
          # Create the output DataFrame
          output df = pd.DataFrame({"essayid": essay ids, "score": predicted scores})
          # Save to CSV file
          output csv path = "data/H00494101-GrossmannLeMauguenArthur-3.csv"
          output df.to csv(output csv path, index=False)
          # Return the output file path
          print(f"Predictions saved to {output csv path}")
```

Predictions saved to data/H00494101-GrossmannLeMauguenArthur-3.csv

Ensemble learning with new gridsearch for the SVM and RF

```
from sklearn.model_selection import GridSearchCV, StratifiedKFold, train_test_split
from sklearn.svm import SVC
from sklearn.ensemble import RandomForestClassifier, VotingClassifier
from sklearn.metrics import cohen_kappa_score, make_scorer
from sklearn.preprocessing import StandardScaler
```

```
from sklearn.decomposition import PCA
import pandas as pd
import numpy as np
# Define QWK scoring function
def quadratic weighted kappa(y true, y pred):
    return cohen kappa score(y true, y pred, weights="quadratic")
qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
# Load dataset
file path = "data/F78DS-Essay-Features.csv"
df = pd.read csv(file path)
# Separate features (X) and labels (v)
X = df.drop(columns=["score", "essayid"])
v = df["score"] - 1 # Adjust labels to start from 0
# Split dataset
X train, X test, y train, y test = train test split(X, y, test size=0.2, stratify=y, random state=42)
# Scale features
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Apply PCA (Optimize component selection)
pca = PCA(n components=0.99) # Retain 99% variance
X train pca = pca.fit transform(X train scaled)
X test pca = pca.transform(X test scaled)
# SVM Hyperparameter Grid Search (Refined)
svm model = SVC(probability=True, random state=42)
param grid svm = {
    'C': [0.01, 0.1, 0.5, 1, 5, 10],
    'kernel': ['rbf'], # Best kernel from previous search
    'gamma': [0.001, 0.005, 0.01, 0.05, 0.1],
    'class weight': ['balanced', None]
cv strategy = StratifiedKFold(n splits=10, shuffle=True, random state=42)
grid search svm = GridSearchCV(svm model, param grid svm, cv=cv strategy, n jobs=-1, verbose=2, scoring=qwk scorer)
grid search svm.fit(X train pca, y train)
```

```
best svm model = grid search svm.best estimator
best svm pred = best svm model.predict(X test pca)
# Evaluate Best SVM Model
accuracy = grid search svm.best score
gwk = quadratic weighted kappa(v test, best svm pred)
print(f"Best SVM Hyperparameters: {grid search svm.best params }")
print(f"SVM Test Accuracy: {accuracy:.4f}, OWK: {gwk:.4f}")
# Random Forest Hyperparameter Grid Search
rf model = RandomForestClassifier(random state=42)
param grid rf = {
    'n estimators': [100, 200, 300],
    'max depth': [10, 20, 30, None],
    'min samples split': [2, 5, 10],
    'min samples leaf': [1, 2, 4],
    'class weight': ['balanced', None]
grid search rf = GridSearchCV(rf model, param grid rf, cv=cv strategy, n jobs=-1, verbose=2, scoring=qwk scorer)
grid search rf.fit(X train pca, y train)
best rf model = grid search rf.best estimator
best rf pred = best rf model.predict(X test pca)
# Evaluate Best RF Model
rf qwk = quadratic weighted kappa(y test, best rf pred)
print(f"Best RF Hyperparameters: {grid search rf.best params }")
print(f"RF QWK: {rf qwk:.4f}")
# Ensemble Learning (SVM + RandomForest)
voting clf = VotingClassifier(estimators=[('svm', best svm model), ('rf', best rf model)], voting='soft')
voting clf.fit(X train pca, y train)
voting pred = voting clf.predict(X test pca)
# Evaluate Ensemble Model
ensemble qwk = quadratic_weighted_kappa(y_test, voting_pred)
print(f"Ensemble Model QWK: {ensemble qwk:.4f}")
```

Fitting 10 folds for each of 60 candidates, totalling 600 fits

c:\Users\Framework_Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model_selection_split.py:805: UserW
arning: The least populated class in y has only 3 members, which is less than n_splits=10.
 warnings.warn(

```
Best SVM Hyperparameters: {'C': 5, 'class_weight': 'balanced', 'gamma': 0.005, 'kernel': 'rbf'}
SVM Test Accuracy: 0.6725, QWK: 0.6676
Fitting 10 folds for each of 216 candidates, totalling 2160 fits
c:\Users\Framework_Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model_selection\_split.py:805: UserW arning: The least populated class in y has only 3 members, which is less than n_splits=10.
    warnings.warn(
Best RF Hyperparameters: {'class_weight': 'balanced', 'max_depth': 10, 'min_samples_leaf': 4, 'min_samples_split': 2, 'n_estimat ors': 200}
RF QWK: 0.6796
Ensemble Model QWK: 0.6740
```

Another ensemble learning with new gridsearch for the SVM and RF

[!!may take long to execute on your computer!! (15+min on mine)]

```
In [103... | from sklearn.model selection import GridSearchCV, StratifiedKFold, train test split
          from sklearn.svm import SVC
          from sklearn.ensemble import RandomForestClassifier, VotingClassifier
          from sklearn.metrics import cohen kappa score, make scorer
          from sklearn.preprocessing import StandardScaler
          from sklearn.decomposition import PCA
          import pandas as pd
          import numpy as np
          # Define OWK scoring function
          def quadratic weighted kappa(y true, y pred):
              return cohen_kappa_score(y_true, y_pred, weights="quadratic")
          qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
          # Load dataset
          file path = "data/F78DS-Essay-Features.csv"
          df = pd.read csv(file path)
          # Separate features (X) and labels (y)
          X = df.drop(columns=["score", "essayid"])
          y = df["score"] - 1 # Adjust labels to start from 0
          # Split dataset
          X train, X test, y train, y test = train test split(X, y, test size=0.2, stratify=y, random state=42)
          # Scale features
```

```
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
# Apply PCA (Optimize component selection)
pca = PCA(n components=0.99) # Retain 99% variance
X train pca = pca.fit transform(X train scaled)
X test pca = pca.transform(X test scaled)
# SVM Hyperparameter Grid Search (Refined)
svm model = SVC(probability=True, random state=42)
param grid svm = {
    'C': [1, 3, 5, 7, 10], # Focusing on best range
    'kernel': ['rbf'], # Best kernel from previous search
    'gamma': [0.001, 0.002, 0.005, 0.007, 0.01], # More refined search
    'class weight': ['balanced', None]
cv strategy = StratifiedKFold(n splits=10, shuffle=True, random state=42)
grid search svm = GridSearchCV(svm model, param grid svm, cv=cv strategy, n jobs=-1, verbose=2, scoring=qwk scorer)
grid search svm.fit(X train pca, y train)
best svm model = grid search svm.best estimator
best svm pred = best svm model.predict(X test pca)
# Evaluate Best SVM Model
accuracy = grid search svm.best score
gwk = quadratic weighted kappa(v test, best svm pred)
print(f"Best SVM Hyperparameters: {grid search svm.best params }")
print(f"SVM Test Accuracy: {accuracy:.4f}, OWK: {qwk:.4f}")
# Random Forest Hyperparameter Grid Search
rf model = RandomForestClassifier(random state=42)
param grid rf = {
    'n estimators': [100, 200, 300, 400], # Adding 400 for deeper exploration
    'max depth': [10, 20, 30, 40, None], # More depth options
    'min_samples_split': [2, 5, 10, 15], # More variations
    'min samples leaf': [1, 2, 4, 6], # Exploring a wider range
    'class weight': ['balanced', None]
grid search rf = GridSearchCV(rf model, param grid rf, cv=cv strategy, n jobs=-1, verbose=2, scoring=qwk scorer)
grid search rf.fit(X train pca, y train)
```

```
best rf model = grid search rf.best estimator
best rf pred = best rf model.predict(X test pca)
# Evaluate Best RF Model
rf qwk = quadratic weighted kappa(y test, best rf pred)
print(f"Best RF Hyperparameters: {grid search rf.best params }")
print(f"RF OWK: {rf qwk:.4f}")
# Ensemble Learning (SVM + RandomForest)
voting clf = VotingClassifier(estimators=[('svm', best svm model), ('rf', best rf model)], voting='soft')
voting clf.fit(X train pca, v train)
voting pred = voting clf.predict(X test pca)
# Evaluate Ensemble Model
ensemble gwk = quadratic weighted kappa(y test, voting pred)
print(f"Ensemble Model OWK: {ensemble gwk:.4f}")
Fitting 10 folds for each of 50 candidates, totalling 500 fits
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model selection\ split.py:805: UserW
arning: The least populated class in y has only 3 members, which is less than n splits=10.
  warnings.warn(
Best SVM Hyperparameters: {'C': 10, 'class weight': 'balanced', 'gamma': 0.002, 'kernel': 'rbf'}
SVM Test Accuracy: 0.6782, OWK: 0.6630
Fitting 10 folds for each of 640 candidates, totalling 6400 fits
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model selection\ split.py:805: UserW
arning: The least populated class in y has only 3 members, which is less than n splits=10.
 warnings.warn(
Best RF Hyperparameters: {'class weight': 'balanced', 'max depth': 30, 'min samples leaf': 6, 'min samples split': 2, 'n estimat
ors': 400}
RF OWK: 0.6676
Ensemble Model OWK: 0.6821
Best SVM Hyperparameters: {
'C': 10.
'class_weight': 'balanced',
'gamma': 0.002,
'kernel': 'rbf'
SVM Test Accuracy: 0.6782, QWK: 0.6630
```

```
Best RF Hyperparameters: { 'class_weight': 'balanced', 'max_depth': 30, 'min_samples_leaf': 6, 'min_samples_split': 2, 'n_estimators': 400 } RF QWK: 0.6676

Ensemble Model QWK: 0.6821
```

Final Kaggle submission (QWK = 0.693)

```
In [104...
          from sklearn.model selection import StratifiedKFold, train test split
          from sklearn.svm import SVC
          from sklearn.ensemble import RandomForestClassifier, VotingClassifier
          from sklearn.metrics import cohen kappa score, make scorer
          from sklearn.preprocessing import StandardScaler
          from sklearn.decomposition import PCA
          import pandas as pd
          import numpy as np
          import joblib
          # Define QWK scoring function
          def quadratic weighted kappa(v true, v pred):
              return cohen kappa score(y true, y pred, weights="quadratic")
          qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
          # Load dataset
          file path = "data/F78DS-Essay-Features.csv"
          df = pd.read csv(file path)
          # Separate features (X) and labels (y)
          X = df.drop(columns=["score", "essayid"])
          y = df["score"] - 1 # Adjust labels to start from 0
          # Split dataset
          X train, X test, y train, y test = train test split(X, y, test size=0.2, stratify=y, random state=42)
          # Scale features
          scaler = StandardScaler()
          X_train_scaled = scaler.fit_transform(X_train)
          X test scaled = scaler.transform(X test)
```

```
# Apply PCA (Optimize component selection)
pca = PCA(n components=0.99) # Retain 99% variance
X train pca = pca.fit transform(X train scaled)
X test pca = pca.transform(X test scaled)
# Define best models with hyperparameters
best svm model = SVC(C=5, kernel='rbf', gamma=0.005, class weight='balanced', probability=True, random state=42)
best rf model = RandomForestClassifier(n estimators=200, max depth=10, min samples split=2, min samples leaf=4, class weight='bal
# Train modeLs
best svm model.fit(X train pca, v train)
best rf model.fit(X train pca, y train)
# Ensemble Learning (SVM + RandomForest)
voting clf = VotingClassifier(estimators=[('svm', best svm model), ('rf', best rf model)], voting='soft')
voting clf.fit(X train pca, y train)
# Save models, scaler, and PCA
joblib.dump(voting clf, "models/ensemble model2.pkl")
joblib.dump(scaler, "models/scaler2.pkl")
joblib.dump(pca, "models/pca2.pkl")
joblib.dump(X.columns.tolist(), "models/features2.pkl")
# Load the submission dataset
submission file path = "data/F78DS-Essay-Features-Submission.csv"
submission df = pd.read csv(submission file path)
# Extract "essayid" and preprocess the features
essay ids = submission df["essayid"]
X submission = submission df[X.columns.tolist()]
# Scale and apply PCA to the submission data
X submission scaled = scaler.transform(X submission)
X submission pca = pca.transform(X submission scaled)
# Predict the scores using the trained ensemble model
predicted scores = voting clf.predict(X submission pca)
predicted scores = predicted scores + 1 # Reverse the previous label shift
# Create the output DataFrame
output df = pd.DataFrame({"essayid": essay ids, "score": predicted scores})
output csv path = "data/H00494101-GrossmannLeMauguenArthur-4.csv"
output df.to csv(output csv path, index=False)
```

```
print(f"Predictions saved to {output_csv_path}")
```

Predictions saved to data/H00494101-GrossmannLeMauguenArthur-4.csv

I tried to add another model to my ensemble model, XGB but it didn't improve the results

```
In [118...
          import xgboost as xgb
          from sklearn.model selection import train test split, RandomizedSearchCV
          from sklearn.metrics import cohen kappa score, make scorer
          from sklearn.preprocessing import StandardScaler
          from sklearn.decomposition import PCA
          import pandas as pd
          import numpy as np
          import joblib
          # Define QWK scoring function
          def quadratic weighted kappa(y true, y pred):
              return cohen kappa score(y true, y pred, weights="quadratic")
          qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
          # Load dataset
          file path = "data/F78DS-Essay-Features.csv"
          df = pd.read csv(file path)
          # Separate features (X) and labels (y)
          X = df.drop(columns=["score", "essayid"])
          v = df["score"] - 1 # Adjust labels to start from 0
          # Split dataset
          X train, X test, y train, y test = train test split(X, y, test size=0.2, stratify=y, random state=42)
          # Scale features
          scaler = StandardScaler()
          X train scaled = scaler.fit transform(X train)
          X test scaled = scaler.transform(X test)
          # Apply PCA
          pca = PCA(n components=0.99) # Retain 99% variance
          X train pca = pca.fit transform(X train scaled)
          X test pca = pca.transform(X test scaled)
```

```
# Define XGBoost model
xgb model = xgb.XGBClassifier(objective='multi:softprob', eval metric='mlogloss', use label encoder=False)
# Reduced hyperparameter grid for XGBoost
param dist = {
    'n estimators': [100, 200, 300],
    'learning rate': [0.01, 0.05, 0.1],
    'max depth': [3, 5, 7],
    'subsample': [0.7, 0.8, 0.9],
    'colsample bytree': [0.7, 0.8, 0.9],
    'scale pos weight': [1, 2],
    'gamma': [0, 0.01, 0.1],
    'reg alpha': [0, 0.1],
    'reg lambda': [1, 1.5]
# Randomized search with cross-validation
random search xgb = RandomizedSearchCV(xgb model, param distributions=param dist, n iter=50, cv=5, n jobs=-1, verbose=2, scoring=
random search xgb.fit(X train pca, y train)
print(f"Best XGBoost Hyperparameters: {random search xgb.best params }")
# Best XGBoost model
best xgb model = random search xgb.best estimator
# Evaluate the best model
best xgb pred = best xgb model.predict(X test pca)
xgb gwk = cohen kappa score(y test, best xgb pred, weights="quadratic")
print(f"XGBoost QWK: {xgb qwk:.4f}")
# Save the model
joblib.dump(best xgb model, "models/xgb model.pkl")
Fitting 5 folds for each of 50 candidates, totalling 250 fits
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\sklearn\model selection\ split.py:805: UserW
arning: The least populated class in y has only 3 members, which is less than n splits=5.
 warnings.warn(
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\xgboost\training.py:183: UserWarning: [01:0
5:48] WARNING: C:\actions-runner\ work\xgboost\xgboost\src\learner.cc:738:
Parameters: { "scale pos weight", "use label encoder" } are not used.
 bst.update(dtrain, iteration=i, fobj=obj)
```

```
Best XGBoost Hyperparameters: {'subsample': 0.7, 'scale pos weight': 2, 'reg lambda': 1, 'reg alpha': 0.1, 'n estimators': 100,
          'max depth': 3, 'learning rate': 0.1, 'gamma': 0, 'colsample bytree': 0.8}
          XGBoost OWK: 0.6773
          ['models/xgb model.pkl']
Out[118]:
In [116... | from sklearn.model selection import train test split
          from sklearn.svm import SVC
          from sklearn.ensemble import RandomForestClassifier, VotingClassifier
          from sklearn.metrics import cohen kappa score, make scorer
          from sklearn.preprocessing import StandardScaler
          from sklearn.decomposition import PCA
          import xgboost as xgb
          import pandas as pd
          import numpy as np
          import joblib
          # Define OWK scoring function
          def quadratic weighted kappa(y true, y pred):
               return cohen kappa score(y true, y pred, weights="quadratic")
          qwk scorer = make scorer(quadratic weighted kappa, greater is better=True)
          # Load dataset
          file path = "data/F78DS-Essay-Features.csv"
          df = pd.read_csv(file path)
          # Separate features (X) and labels (v)
          X = df.drop(columns=["score", "essayid"])
          y = df["score"] - 1 # Adjust labels to start from 0
          # Split dataset
          X train, X test, y train, y test = train test split(X, y, test size=0.2, stratify=y, random state=42)
          # Scale features
          scaler = StandardScaler()
          X_train_scaled = scaler.fit_transform(X_train)
          X test scaled = scaler.transform(X test)
          # Apply PCA
          pca = PCA(n_components=0.99) # Retain 99% variance
          X train pca = pca.fit transform(X train scaled)
          X test pca = pca.transform(X test scaled)
```

```
# Define best models with hyperparameters
best svm model = SVC(C=5, kernel='rbf', gamma=0.005, class weight='balanced', probability=True, random state=42)
best rf model = RandomForestClassifier(n estimators=200, max depth=10, min samples split=2, min samples leaf=4, class weight='bal
# Train SVM and RandomForest models
best svm model.fit(X train pca, y train)
best rf model.fit(X train pca, v train)
# Define XGBoost model with best hyperparameters
best xgb model = xgb.XGBClassifier(
    objective='multi:softprob',
    eval metric='mlogloss',
   use label encoder=False,
   n estimators=100,
   learning rate=0.1,
   max depth=3,
    subsample=0.7,
    colsample bytree=0.8,
   scale pos weight=2,
   gamma=0,
    reg alpha=0.1,
    reg lambda=1
# Train XGBoost model
best xgb model.fit(X train pca, y train)
# Ensemble Learning (SVM + RandomForest + XGBoost)
voting clf = VotingClassifier(estimators=[('svm', best svm model), ('rf', best rf model), ('xgb', best xgb model)], voting='soft'
voting clf.fit(X train pca, y train)
# Evaluate the ensemble model
ensemble pred = voting clf.predict(X test pca)
ensemble qwk = cohen kappa score(y test, ensemble pred, weights="quadratic")
print(f"Ensemble OWK: {ensemble gwk:.4f}")
# Save models, scaler, and PCA
joblib.dump(voting clf, "models3/ensemble model.pkl")
joblib.dump(scaler, "models3/scaler.pkl")
joblib.dump(pca, "models3/pca.pkl")
joblib.dump(X.columns.tolist(), "models3/features.pkl")
# Load the submission dataset
submission file path = "data/F78DS-Essay-Features-Submission.csv"
```

```
submission df = pd.read csv(submission file path)
# Extract "essayid" and preprocess the features
essay ids = submission df["essayid"]
X submission = submission df[X.columns.tolist()]
# Scale and apply PCA to the submission data
X submission scaled = scaler.transform(X submission)
X submission pca = pca.transform(X submission scaled)
# Predict the scores using the trained ensemble model
predicted scores = voting clf.predict(X submission pca)
predicted scores = predicted scores + 1 # Reverse the previous label shift
# Create the output DataFrame
output df = pd.DataFrame({"essayid": essay ids, "score": predicted scores})
output csv path = "data/H00494101-GrossmannLeMauguenArthur-5.csv"
output df.to csv(output csv path, index=False)
print(f"Predictions saved to {output csv path}")
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\xgboost\training.py:183: UserWarning: [01:0
2:48] WARNING: C:\actions-runner\ work\xgboost\xgboost\src\learner.cc:738:
Parameters: { "scale pos weight", "use label encoder" } are not used.
 bst.update(dtrain, iteration=i, fobj=obj)
c:\Users\Framework Arthur\AppData\Local\Programs\Python\Python313\Lib\site-packages\xgboost\training.py:183: UserWarning: [01:0
2:49] WARNING: C:\actions-runner\ work\xgboost\xgboost\src\learner.cc:738:
Parameters: { "scale pos weight", "use label encoder" } are not used.
 bst.update(dtrain, iteration=i, fobj=obj)
Ensemble OWK: 0.6601
Predictions saved to data/H00494101-GrossmannLeMauguenArthur-5.csv
```

5) Conclusion

1) Conclude your coursework.

In this coursework, we explored the "F78DS-Essay-Features.csv" dataset to uncover patterns and correlations between various textual features and their assigned scores. The dataset provided a rich set of linguistic and structural attributes. Our objective was to analyze these features to identify

the most influential factors in determining essay quality.

Key Findings

- Feature Selection: We conducted a correlation analysis and used a variance threshold to identify the most relevant features.
- Model Comparison: We compared the performance of Naïve Bayes (NB) and Random Forest (RF) models.
 The Random Forest model outperformed Naïve Bayes, achieving a higher Quadratic Weighted Kappa (QWK) score.
 Further improvements were made using ensemble learning, combining SVM and Random Forest models, which yielded the best QWK score of 0.6821.
- Hyperparameter Tuning: We employed GridSearchCV to fine-tune hyperparameters for both SVM and Random Forest models.

Challenges and Limitations

- Class Imbalance: The dataset had an imbalanced distribution of scores, which posed challenges in model training and evaluation.
- Feature Engineering: Further feature engineering and selection could potentially improve model performance.
- Computational Resources: Grid search for hyperparameter tuning was computationally intensive and time-consuming.

Futur Improvement

- Advanced Techniques: Explore more advanced techniques such as deep learning models for text classification.
- Feature Engineering: Investigate additional feature engineering techniques to capture more nuanced textual characteristics.
- Model Interpretability: Improve the interpretability of the models to understand which features contribute most to the predictions.
- Model Diversity: Incorporate a diverse set of models in the ensemble to capture different aspects of the data.
- Computational Efficiency improvement
- Better visualizations

6) Sources

- 1. Random Forest Hyperparameter Tuning in Python GeeksforGeeks
- 2. Fine-Tuning Your Random Forest Classifier: A Guide to Hyperparameter Tuning Medium (Tahera Firdose)

- 3. Scikit-Learn Official Documentation Scikit-Learn
- 4. Random Forests in Scikit-Learn Scikit-Learn Documentation
- 5. Alternatives to Brute-Force Parameter Search Scikit-Learn Documentation
- 6. Random Forest Regression: A Practical Guide Keboola
- 7. Feature Engineering Wikipedia