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Machine Learning Approaches to Ethical Analysis of Statistics

MATTHEW KENELY, KARL CHETCUTI, JAN LAWRENCE FORMOSA, and DAVID CACHIA ENRIQUEZ

I. INTRODUCTION

The dataset selected for this study is the Student Performance¹ dataset from Kaggle. This dataset provides comprehensive statistical information on a range of academic, behavioural, and demographic variables influencing students' performance.

The data was collected from two Portuguese schools and includes variables such as age, gender, family size, and grades in mathematics and Portuguese, among others. It is particularly well-suited for analysing relationships between educational and demographic factors to determine potential correlations that contribute to students' academic performance.

To analyse the dataset, four machine learning (ML) techniques are applied to classify and predict students' levels of academic performance. The techniques employed are:

- Linear Regression
- Ensemble Learning
- K-means Clustering
- Principal Component Analysis

The GitHub repository for this study is available at https://github.com/matthewkenely/ics5110

II. BACKGROUND

A. MACHINE LEARNING TECHNIQUES

1) Linear Regression

Linear Regression is a supervised machine learning algorithm used to predict a continuous numerical value. For each input, the algorithm estimates a corresponding output value that lies on a continuous scale [1].

Linear Regression combines the inputs and features in a linear way, where the predicted output is modeled as a linear combination of the input features and their corresponding weights, plus a bias term. There are two primary types of linear regression.

Simple Linear Regression (SLR) makes use of a single independent variable and a single dependent variable [1]. The equation for SLR is:

$$y = \beta_0 + \beta_1 X \tag{1}$$

where:

• y is the dependent variable,

¹https://www.kaggle.com/datasets/larsen0966/student-performance-dataset

- *X* is the independent variable,
- β_0 is the intercept,
- β_1 is the slope.

Multiple Linear Regression (MLR) makes use of multiple independent variables and one dependent variable [2]. The equation for MLR is:

$$y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_n X_n \tag{2}$$

where:

- y is the dependent variable,
- X_1, X_2, \dots, X_n are the independent variables,
- β_0 is the intercept,
- $\beta_1, \beta_2, \dots, \beta_n$ are the slopes.

The goal of linear regression is to determine the best-fit line for a model. This line is drawn to best represent the data points, serving as a predictor for the output based on a given input. To minimize the differences between the predicted and actual values, the best-fit line is positioned as close as possible to all the data points.

The cost function is used to assess how well the model's predictions match the actual data. Rather than simply indicating whether a prediction was correct or incorrect, the cost function calculates the magnitude of the error. If the prediction is close to the actual value, the cost is small; if the prediction deviates, the cost is larger. The objective of linear regression is to adjust the weights of each feature to minimise the total cost across all predictions, thereby improving the model's accuracy.

2) Ensemble Learning

Ensemble Learning is a machine learning technique that employs a multi-model approach to generate predictions. Ensemble learning algorithms train multiple smaller models, each making its own prediction. These predictions are then aggregated to produce the final output. The idea of this approach is that by having multiple learners and then combining their results, the model would overcome errors that are inherent to single individual learner models, which would lead to more accurate and reliable predictions [3].

There are various approaches to Ensemble Learning, all of which involve the use of multiple learners but differ in their training processes and decision-making mechanisms. These approaches include:

- Bagging (Bootstrap Aggregating): Trains multiple learners in parallel on different subsets of the input data. These learners operate independently, without sharing information. At the end of the training process, their predictions are consolidated. Depending on the task, this consolidation is typically done through majority voting (for classification) or averaging (for regression) [4].
- **Boosting**: Trains multiple learners sequentially, with each learner using the results of the previous one. The output of each learner is passed on to the next, which attempts to correct the errors made by its predecessor. This iterative process continues for a set number of iterations. The final prediction is a weighted average of all individual learner predictions [5].
- Stacking: Involves training multiple base models in parallel. These models can vary in type and are typically chosen for their complementary strengths. The final prediction is made by a meta-model, which is trained to predict based on the outputs of all the base models [3].
- Blending: Similar to stacking, blending also uses multiple base models and a final meta-model to make predictions. The key distinction is that the meta-model in blending is trained on a separate *holdout* dataset or aggregates predictions using a weighted average [6].
- **Voting**: Uses multiple learners to generate predictions but differs in how the results are combined. This aggregation can take two forms:
 - -- **Hard Voting**: The final prediction is made by taking the majority response from the learners.
 - Soft Voting: The prediction probabilities are averaged, and the class with the highest probability is selected.

For this study, we employ three Ensemble Learning techniques: Bagging (using Random Forest), Boosting (using Gradient Boosting), and Stacking (combining Random Forest and Gradient Boosting as base models, with Logistic Regression as the meta-model).

3) K-means Clustering

K-Means clustering is an unsupervised machine learning algorithm designed to partition data into a specified number, k, of clusters based on feature similarity. It is commonly used for segmentation tasks and works by minimising the distance between data points from a dataset using their centroids, which are essentially the centres of the clusters, to group them together into clusters.

The algorithm begins by randomly initializing centroids within the feature space of the dataset. The distance between each data point and the centroids is then calculated using the Euclidean distance, defined by the following equation:

$$d(x,y) = \sqrt{\sum_{i=1}^{n} (y_i - x_i)^2}$$

The data point will be assigned to the nearest centroid cluster based on this measurement.

Once all of the data points are assigned to a cluster, the centroids are re-initialised as the average position of all the data points of their respective cluster. The steps of assigning data points to their nearest cluster and then reinitialising the centroids will be repeated until the centroids are no longer changing, which means they are optimal, or until a predefined number of iterations is reached.

4) Principal Component Analysis

Principal Component Analysis (PCA) is a dimensionality reduction technique that converts data into a new coordinate system while maintaining important patterns and trends.

PCA can be broken down into five main steps:

- 1) Data Standardisation
- 2) Covariance Matrix Computation
- 3) Eigendecomposition
- 4) Component Selection
- 5) Data Transformation

First, the data is standardised so that the variables have a mean of 0 and a standard deviation of 1. This critical step ensures that there is no large difference in scale between the initial features, as such differences would bias the analysis. The standardisation process can be represented as:

$$z = \frac{x - \mu}{\sigma}$$

where z is the standardised value, x is the original value, μ is the mean, and σ is the standard deviation [7].

Secondly, the covariance matrix of the standardised data is computed. The covariance matrix highlights the relationships between pairs of variables in terms of their variance. It is calculated as:

$$Cov(X) = \frac{1}{n}X^T X$$

The third step is Eigendecomposition which is performed on the covariance matrix in order to extract the Eigenvectors and Eigenvalues. Eigenvectors define the directions of principle components while the Eigenvalues indicate the amount of variance captured by each principle component.

After Eigendecomposition, the eigenvectors are sorted based on their eigenvalues in descending order. This means that the first eigenvector corresponds to the direction of maximum variance in the data. k principal components are then chosen depending on how much of the total variance should be retained.

explained variance =
$$\frac{\sum_{i=1}^{k} eigenvalue_i}{\sum_{i=1}^{p} eigenvalue_i}$$

Finally, the selected principal components are used to create a projection matrix. The original data is then multiplied by this projection matrix to transform the data:

$$Z = X \cdot P$$

where Z is the transformed data, X is the original data, and P is the projection matrix.

B. RESCALING AND NORMALISATION

Before processing data, it is essential to standardise the distribution and range of different variables, particularly when they have different units or scales. Rescaling is the process used to achieve this standardization, which involves adjusting the data to a specific range, such as from 1 to 100. This helps ensure that the model interprets the variables on a comparable scale, improving its performance [8].

Normalisation, on the other hand, involves transforming the data to follow a standard distribution, such as a Gaussian distribution with a mean of 0 and a standard deviation of 1 [9]. This transformation is crucial for models like Support Vector Machines (SVM) and Logistic Regression, as it improves both the accuracy and convergence speed during training.

C. CROSS-VALIDATION

Cross Validation is a technique used in Machine Learning to test a model's performance on unseen data. This helps prevent overfitting, where a model learns the training data too well and has poor performance on new data [10]

The process of cross validation is done as follows:

- Available data is split into different folds
- The model is trained using all of the folds apart from one
- The model is tested on the remaining fold that was not used during training
- The process is repeated until every fold has acted as the test fold

After all the iterations are complete, the results are averaged to get an estimate of the model's performance. Through cross validation, the model can better generalise to unseen data.

D. DIMENSIONALITY REDUCTION AND FEATURE SELECTION

Dimensionality Reduction and Feature Selection are methods that are used to streamline the dataset by removing irrelevant and redundant features. By removing these and only focusing on the dataset's most relevant features, the performance of the model is improved.

Dimensionality Reduction techniques use a lower number of features (dimensions) of the data in a dataset while retaining as much of the important information as possible. By transforming data into a lower-dimensional space while still preserving the key aspects of said data, the complexity of the model can be reduced, which may also improve its generalisation performance [11]. The PCA technique used in this project follows this principle and is a common dimensionality reduction method.

Feature Selection works differently from Dimensionality Reduction, as instead of lowering the dimension of dataset features while retaining the essence of the data, it removes irrelevant and redundant features from the dataset. Instead of creating new features to transform the dataset into a lower-dimensional space, feature selection streamlines the dataset by selecting a subset of the most important features.

E. QUANTITATIVE MEASURES

1) Accuracy

Accuracy is one of the most simple and straightforward performance measures for evaluating machine learning models. It measures the percentage of correctly predicted instances out of the total instances. It can be expressed as:

$$Accuracy = \frac{Number of Correct Predictions}{Total Number of Predictions}$$
 (3)

2) PCA Visualisation

PCA visualisation refers to the process of projecting PCAreduced data onto a 2D or 3D space to facilitate easier interpretation and exploration of the underlying structure of the data.

By plotting the first two or three principal components, which capture the most variance in the data, PCA visualisation allows for the identification of clusters, outliers, and patterns that may be harder to interpret in higher dimensions.

III. DATA PREPARATION

For this study, we selected the Student Performance dataset for our experiments. This dataset contains information collected from two Portuguese schools, focusing on student performance in two subjects: Mathematics and Portuguese. As a result, the dataset is divided into two sections, each corresponding to one of the subjects.

For the purposes of this study, we chose to work with only the Portuguese section of the dataset, as it contained a larger volume of data, which better suited the requirements of our analysis.

A. DATA CLEANING AND TRANSFORMATION

Data Cleaning and Data Transformation are essential techniques employed by computer scientists to improve the quality, usability, and efficiency of data for analysis or application.

Data Cleaning focuses on identifying and handling errors, inconsistencies, missing values or even unneeded data from a datasets. This may include tasks such as removal of duplicate data, standardizing formats, or filling in gaps using statistical methods, such as by using interpolation

Data Transformation on the other hand, involves converting data into a more suitable or meaningful format. This might include operations such as scaling numerical values, encoding categorical variables, aggregating data, or restructuring datasets to meet the requirements of specific algorithms or workflows.

In this study, no single cleaning or transformation method was applied universally across the dataset. Instead, these processes were tailored individually to each experiment, as the handling of the data varied depending on the specific needs of the experiment. Examples of the techniques applied include:

• **Standardization**: A feature-scaling technique that transforms all values in the dataset to a common range, typically between 0 and 1.



- Feature Selection: The process of determining which features to include or exclude based on their relevance to the analysis. The approach and rationale for feature selection varied across experiments.
- Label Encoding: A method for translating categorical data into a numerical format. This is done by attaching a numerical label to each different instance of data in a feature.

These techniques will be discussed in greater detail in the following sections, where each experiment is explored comprehensively.

B. DISTRIBUTION ANALYSIS

Distribution analysis involves examining how the data points in a dataset are spread. This kind of analysis helps to identify patterns and trends in the data, as well as reveal outliers that could lead to unproductive experiments. Visualizations such as histograms or box plots are commonly used to facilitate a better understanding of the data distribution. By conducting distribution analysis, we can derive insights that guide the direction of an experiment and ensure that each experiment is sufficiently distinct from others.

C. FEATURE ANALYSIS

The initial examination of the dataset involved performing feature analysis on each feature, which was visualized using histograms. This analysis revealed the distribution of each feature and helped determine whether it would be worthwhile to train a machine learning model to predict that specific feature.

One of the ideas discussed early on was to train a model to predict the number of past failures without using any graderelated features. Although this idea seemed promising, the feature analysis showed a significant imbalance in the 'failures' feature. The experiment confirmed this, as the model learned to predict '0' for nearly all data points due to the fact that over 80

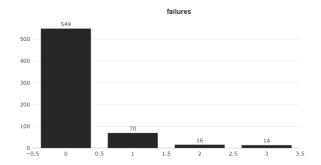


FIGURE 1. Feature distribution of the "failures" feature, demonstrating

This issue of imbalance was not limited to the 'failures' feature. Other features, such as "Pstatus", "schoolsup", "paid", and "Dalc", also exhibited imbalances. Despite these imbalances, most of the other features had distributions that were much more suitable for training machine learning models.

D. CORRELATION MATRIX

A second important analysis involved creating a correlation matrix to examine the relationships between the features in the dataset. The correlation matrix allows for a clear visualization of pairwise correlations, helping identify relationships and patterns between different features. This tool provided valuable insights that guided several experimental decisions.

One key finding from the correlation matrix was the high correlation between the three grade features: G1, G2, and G3. These features had an average correlation score of 87%, suggesting that they were strongly related. This observation aligns with logical reasoning: if a student performs well in one or two of the grades, it is likely that the third grade will also reflect similar performance. The same applies to poor grades, where all three features tend to show similar patterns. Consequently, these features can act as proxies for each other, meaning that when predicting one of them, the other two will have a significant impact on the model's decision.

Further instances of how correlations influenced the experimental setup will be discussed in later sections, providing additional context for these relationships.

IV. EXPERIMENTS

For this study, four distinct experiments were conducted on the dataset, each aimed at exploring different aspects of the data through various machine learning techniques.

An overview of the experiments is as follows:

- General Final Grade Prediction: investigates training a model to predict a student's final grade using an optimal number of features, constrained by specific correlation bounds.
- 2) **Tailored Final Grade Prediction**: examines the use of different feature subgroups to predict the final grade.
- Dimensionality Reduction: explores how reducing the number of features affects the accuracy of the model predicting the final grade.
- Student Segmentation: explores how students can be clustered into different segments based on their characteristics.

A. GENERAL FINAL GRADE PREDICTION

1) Overview

The objective of this experiment was to develop a model capable of predicting a student's final grade (denoted as G3) using the most relevant features, determined by specific correlation thresholds.

To understand the relationship between G3 and the other features, we referred to the previously generated correlation matrix, which highlighted two features—G1 and G2—as having extremely high correlations with G3. These features, which represent the grades for the first and second terms, showed a correlation score exceeding 80%, much higher than the correlations with other features.

As discussed earlier, this high correlation implies that any model trained to predict G3 using G1 and G2 would almost

entirely rely on these two features, making the model less informative. Therefore, for this experiment, G1 and G2 were excluded in order to train a more balanced model. To achieve this, an "upper-bound" threshold was set to exclude features with extremely high correlation values, either positive or negative. The optimal upper-bound was set to 0.8, meaning any feature with an absolute correlation above 80

Additionally, a "lower-bound" threshold was applied to remove features with extremely low correlation values (close to 0%) that were identified as noise and impeded the model's learning. The optimal lower-bound value was set to 0.065, meaning features with an absolute correlation below 6.5% were excluded.

Thus, four different configurations were tested for this experiment, based on whether the bounded features were included or excluded:

- Including both upper-bound" and lower-bound" features.
- Including upper-bound" but excluding lower-bound" features.
- Excluding upper-bound" but including lower-bound" features.
- Excluding both upper-bound" and lower-bound" features.

2) Machine Learning Techniques

For this experiment, the chosen machine learning technique was Ensemble Learning, using a strategy called Stacking. With stacking, multiple base models are trained on the data, and a final meta-model is then used to combine their predictions and give different weight to each depending on the situation.

The base models used were:

- 1) Random Forest: an ensemble technique that uses a bagging approach. The model uses and combines multiple decision trees that have been created using a random subset of the data. Because of this, Random Forest is particularly useful when dealing with noisy data, as the multiple decision trees help to deal with the noise by splitting it up, and then averaging out the results.
- 2) Gradient Boosting: another ensemble technique that uses a boosting approach. The model also makes use of decision trees, but trains them in a sequential order where the next tree learns the errors done by the previous one and corrects them. Gradient Boosting is useful for learning complex patterns in data, but sometimes requires extensive fine tuning.

For the meta-model, a simple Logistic Regression model was selected. Logistic Regression was chosen for its efficiency and ability to provide probabilistic predictions, making it a suitable choice for aggregating the predictions from the base models.

3) Results

As previously mentioned, four tests were conducted for this experiment, each with different configurations of the upper

and lower bounds.

The best performing model was, as expected, Case 2. This model utilized both G1 and G2, which naturally led to highly accurate predictions, while excluding features that acted as noise. The final accuracy of the model was slightly above 48%. In Figure 2, the predicted values of G3 are plotted against the actual values, with most points lying close to the dotted red line, indicating correct predictions.

The second-best performing model was Case 1, which also included G1 and G2 but did not exclude the lower-bound features. This model achieved an accuracy slightly above 46%. Cases 3 and 4 performed significantly worse, with accuracies just above 21% and just below 24%, respectively.

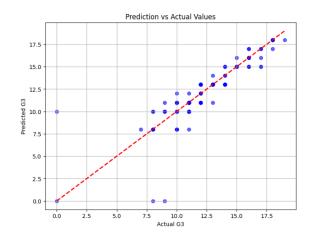


FIGURE 2. Model predictions vs ground truth for best performing model.

The results further emphasize the significant influence of G1 and G2 on predicting G3. The test cases incorporating these features achieved approximately twice the accuracy of the best-performing model that excluded them. However, it is important to highlight that this over-reliance on G1 and G2 may lead the model to underweight other features, potentially hindering its ability to learn diverse patterns in the data. This could negatively impact the model's generalization capability, particularly when handling unseen data.

B. TAILORED FINAL GRADE PREDICTION

1) Overview

The goal of this experiment was to train a model to predict a student's final grade, but with the added complexity of using predefined feature "sub-groups" for training. This approach complements the previously discussed General Final Grade Prediction experiment by examining how well the model performs when trained on different feature subsets.

The three chosen feature sub-groups were as follows:

- **Group 1**: All features.
- **Group 2**: Only the grade features, G1 and G2.
- Group 3: All features excluding the grade features.

These groupings were selected based on the findings from the General Final Grade Prediction experiment, where G1



and G2 were shown to have high correlation with G3. By including or excluding these features, the experiment explores whether the data relies too heavily on these two features.

2) Machine Learning Techniques

For this experiment, a Linear Regression model was chosen. One of the simplest and most widely used regression techniques, Linear Regression models the relationship between a dependent variable and one or more independent variables by fitting a linear equation to the data. Linear Regression is highly interpretable and works well for datasets with linear relationships, but it assumes that the input variables are independent and that there is minimal multicollinearity.

It should be noted that for the experiment, the model's hyperparameters were left as their default values, with no changes made between the different tests.

3) Results

The results clearly demonstrated a distinct performance difference between the models trained with the grade features and those that did not include them. The model trained without the grade features achieved an accuracy of approximately 34%. In contrast, the model trained exclusively on the two grade features (G1 and G2) reached an accuracy of nearly 85%, more than double that of the previous model. The third test, which used all features, resulted in a slight increase in accuracy, with the model achieving just under 86

Although these findings were not entirely unexpected, they reinforce the conclusions of the General Final Grade Prediction experiment, further highlighting the significant dependence of final grade prediction on the G1 and G2 features.

C. DIMENSIONALITY REDUCTION

1) Overview

The goal of this experiment was to reduce the number of features in the dataset and investigate how this reduction would impact the model's accuracy when predicting the final grade (represented by the feature G3).

The experiment was conducted using three versions of the dataset, each with different dimensions. One version used the full dataset, while the other two employed datasets with 95% and 90% explained variance, respectively. Figure 3 illustrates the explained variance as a function of the number of components, which helps to visualize the dimensionality reduction.

2) Machine Learning Techniques

PCA was used for dimensionality reduction. This method combines the original features into new variables, called principal components, which capture the most important patterns in the data. PCA reduces the number of dimensions while keeping the key information, making it useful for simplifying data, reducing noise, and preparing for visualization or analysis.

With the resulting reduced data, Linear Regression models were used to make predictions on G3. Linear regression

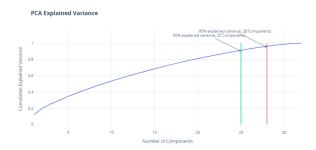


FIGURE 3. Graph illustrating the reduction in cumulative explained variance as the number of components decreases.

is a simple statistical method that models the relationship between a dependent variable and one or more independent variables by fitting a straight line to the data. The line is determined by minimizing the sum of squared differences between the observed and predicted values, making it useful for predicting outcomes and analysing variable relationships.

3) Results

The reduction in model size was notable: from 1.25 KB for the original dataset to 0.85 KB and 0.80 KB for the 95% and 90% explained variance versions, respectively. The dimensionality reduction removed only up to six features from the original dataset.

In terms of accuracy, the Logistic Regression model exhibited the following results:

The original dataset achieved an accuracy of 31The 95% explained variance dataset resulted in an accuracy of 28%. The 90% explained variance dataset yielded an accuracy of 29%. The worst-performing model was the 95% explained variance dataset, which was nearly 10% less accurate than the original. Although this decrease may seem relatively small, it is worth noting that the accuracy was already low, and thus, the change should be interpreted with caution.

D. STUDENT SEGMENTATION

1) Overview

The goal of this experiment was to segment the students into distinct clusters to uncover specific characteristics.

First, the dataset was analyzed for its suitability for K-Means Clustering. Since this algorithm is not ideal for categorical data, a subset of numerical features was selected for clustering. These features were then standardized to ensure that variables with larger scales did not dominate the clustering process.

Next, the number of clusters, k, was determined. The elbow method is a technique that can be used to identify the optimal k clusters that should be generated from a dataset. While other methods exist, such as the silhouette coefficient and gap statistic methods, the elbow method was chosen in particular for its straightforward approach to visualising the optimal number of clusters. Additionally, since the elbow point could be distinctly identified easily (as can be seen in Figure 4),



there was no reason to opt for another method. In the context of this dataset, it was determined that the optimal number is 2 by visualising the point on the curve where the distortion value starts to decrease at a lower rate. K-means was then performed on the standardised data.

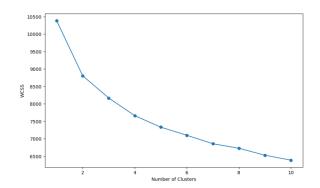


FIGURE 4. Graph showing the relationship between the number of clusters (k) and the Within-Cluster Sum of Squares (WCSS) for the dataset. WCSS represents the total variance within each cluster, and it decreases as the number of clusters increases.

2) Machine Learning Techniques

K-Means Clustering was used for this experiment. K-Means Clustering is an unsupervised machine learning algorithm used to partition data into a specified number, k, of clusters based on feature similarity. It is used for tasks that involve segmentation and works by minimising the distance between data points from a dataset using their centroids, which are essentially the centres of the clusters, to group them together into clusters.

3) Results

Upon analysing the centroids for each cluster, it was found that Cluster 1 (C1) contained slightly older students than Cluster 0 (C0). Students in C0 generally exhibited higher academic performance, as reflected by their "failures" and "G3" scores. This difference may be attributed to several factors, such as higher levels of absenteeism, more frequent social activities, greater alcohol consumption, and less time spent studying in C1 compared to C0. Additionally, the parents of C1 students typically had lower educational levels than those of students in C0, and the students in C1 appeared to have somewhat poorer relationships with their families.

Since the dataset is high-dimensional, with more than two features, Principal Component Analysis (PCA) was used to reduce the dimensionality for visualization purposes. The reduced data was then plotted in a scatter plot, as shown in Figure 5.

V. ETHICAL REVIEW

Ethical considerations must underpin the use of datasets in machine learning to avoid bias and ensure fairness, transparency, and respect for privacy, especially within the domain of education. This analysis examines the ethical implications

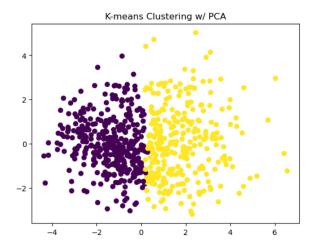


FIGURE 5. Scatter plot visualizing clusters identified by K-Means Clustering after applying dimensionality reduction with Principal Component Analysis (PCA). Each point represents a student, and the two clusters are differentiated by color.

brought by the "Student Performance" dataset and the results generated from applying machine learning techniques to it.

Foremost, it is important to note that the dataset was taken from Kaggle, which is known for its reputable datasets, and compiles student data gathered from a limited number of classes across two Portuguese school involving the subjects of mathematics and Portuguese. C. Paulo [12] claims that the data was collected through a combination of school reports and questionnaires. Therefore, it should be assumed that the data may not accurately reflect the entire student body of these schools, or any school in general for that matter, and may also be subject to outliers.

Furthermore, while the dataset includes information on a total of 649 students, it raises bias concerns due to its sparse geographic representation which might indirectly favour certain groups of people when extending it beyond the scope of Portuguese schools [13], [14]. Additionally, variables in the dataset such as "Medu" and "Fedu" which represent parental education, among other family-related data, pose extensive socioeconomic and cultural concerns that falsely indicate a correlation between familial statuses and student performance. In the context of this dataset, it serves to provide better context towards the specific students in the regional area of these two schools, however, if it were to be used as training data for a machine learning model and extended beyond this region, it might show signs of bias towards those students who come from larger families or whose parents have higher levels of education than others.

As discussed previously, four machine learning experiments were conducted on this dataset. The "General Final Grade Prediction" and "Tailored Final Grade Prediction" experiments both work synonymously to predict what the final grade of a student is. While they make use of the variables "G1" and "G2", which represent a student's first and second period grades, respectively, to predict the final

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grade, they also investigate the use of other features within the dataset to accomplish this task. It should be noted that the best accuracies were mostly correlated to the two grade features, signifying that the models performed better when using statistical performance-related data rather than demographic or familial data. This shows that the models do not exhibit any inherent signs of bias based on a student's background. On the other hand, the "Dimensionality Reduction" experiment may pose ethical issues since it is not transparent with the features it is utilising. The accuracies of this model being low further prove that it is not suitable for such a task since it is most likely making use of demographic data to predict grades.

The "Student Segmentation" experiment involved segmenting students into different clusters to identify specific characteristics. In this case, the model detected certain biases existing within the dataset. For instance, a correlation between age and student performance was identified, with younger students generally performing better in school. Analysing the resulting clusters further revealed an additional correlation between parental education and student performance, as mentioned previously. This is indicative that models trained on this data to predict student grades could inaccurately favour younger students whose parents have higher levels of education, which is reflected in the three previously discussed experiments that performed poorly when taking into consideration such attributes.

Addressing such biases is an important step when applying machine learning models to datasets as it could lead to discriminatory outcomes, such as penalizing students from disadvantaged family backgrounds. The societal impact of deploying such models could unintentionally harm vulnerable groups and enhance existing stigmatization. However, if the biases are rooted from the model, the result might aid struggling students by identifying those who are most likely to fare poorly in school through carefully selected, unbiased features. These issues heavily affect the long-term societal implications of such a model. At a large scale, it could influence educational policies, teacher evaluations, or student tracking systems. With biased data, this would risk institutionalising inequities and perpetuating harmful stereotypes. Furthermore, if the model were to continuously improve itself based upon the gathering of more student data, the chance of there being biased data might increase. Therefore, continuous monitoring would be necessary to ensure that the long-term impact remains positive and aligned with societal values.

VI. WEB PORTAL USAGE GUIDE

Web portal is available at https://mkenely.com/ics5110/.

• Home Page

8

-- Compilation of links to all relevant subpages

• Data Visualisation

-- Feature Reference

- * Table of all features in the dataset we used.
- * Feature: string name of the feature

- * Type: object = categorical feature, int64 = numerical feature
- * Description: textual description of the feature and its categorical labels (if any)
- * Encoding Mappings: labels and their numerical encodings for categorical features

-- Feature Distributions

* Distribution plots for each feature in the database

-- Feature Correlation Matrix

* Shows the correlation between all features. Larger data points indicate greater absolute correlation. Blue points indicate positive correlation, and red points indicate inverse correlation.

-- Feature vs G3 Scatter Plots

* Scatter plots between all features and G3 which we are predicting. The features with the greatest absolute correlation with G3 are shown first. Data points are jittered to get a rough idea of the quantity of individuals at each point due to the large amount of overlap due to G3 being a discrete numerical variable.

• ML Techniques

-- Links to gradio implementations of our 4 ML models. In each implementation, the user is prompted to input their own student features and will be presented with predictions made by the models.

* PCA

 Makes predictions for G3 using 3 different Linear Regression models, each trained on datasets with different numbers of components according to retained variance compared to the original dataset (original, 95% variance, 90% variance). Also shows the runtime and size of each model.

* Ensemble Model

 Makes predictions for G3 using the stacking ensemble model we trained. Also shows the runtime.

* K-Means Clustering

Predicts the cluster to which the inputted student belongs (0 or 1). Also shows the runtime and the indices of the top 5 performing students in the same cluster as the inputted student.

* Linear Regression

 Makes predictions for G3 using 3 different Linear Regression models, each trained on datasets with different combinations of features: all features, all features except grades (G1, G2), and only grades (G1, G2).



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