

# Machine Learning Lab Project Report

Course: SWE344

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## Introduction

### Problem Statement

This project addresses a critical educational challenge: predicting student academic performance early in their academic journey. The ability to accurately forecast final grades (G3) enables targeted interventions for at-risk students, optimized resource allocation, and personalized learning strategies. This predictive modeling task has significant implications for improving educational outcomes in secondary schools.

### Data Overview

The dataset represents student achievements from two Portuguese secondary schools, combining performance in Mathematics and Portuguese language subjects. Collected through school reports and questionnaires, it contains:

- 1044 student records (649 from Portuguese, 395 from Mathematics)
- 33 attributes covering demographic, social, school-related features
- Grade components: G1 (first period), G2 (second period), G3 (final grade)

**Data Source:** UCI Machine Learning Repository

[Student Performance Dataset](#)

## Methodology

### Overall Approach

We follow a structured machine learning workflow:

1. **Data Preprocessing:** Cleaning, transformation, and feature engineering
2. **Exploratory Analysis:** Understanding distributions and relationships

3. **Feature Selection:** Identifying predictive features
4. **Model Development:** Training and tuning multiple algorithms
5. **Evaluation:** Rigorous validation using multiple metrics
6. **Deployment Preparation:** Final model selection and testing

## Prediction Problem Formulation

- **Target Variable:** Final grade (G3) - continuous numerical value (range 0-20)
- **Problem Type:** Regression
- **Evaluation Metrics:**
  - MAE (Mean Absolute Error): Primary metric for interpretability
  - MSE (Mean Squared Error): Emphasizes larger errors
  - RMSE (Root Mean Squared Error): Standard deviation of prediction errors

## Evaluation Metrics and Performance Estimation

Since our target variable is continuous data (G3), this is going to be a regression problem. For this, we are going to use MSE, RMSE and MAE for evaluation metrics. For the performance estimation we will train with 4 different models: linear regression model (lm), SVM, regression trees and random forest. Since the dataset contains more than 500 cases (1044 cases), we will use 10 fold cross validation as the estimation method for internal evaluation. We will use 70:30 holdout for external evaluation.

## Technical Setup

### Software & Tools:

- **Development Environment:** Jupyter Notebook
- **Programming Language:** Python 3.9+
- **Key Python Libraries:**
  - `pandas` (Data manipulation)
  - `numpy` (Numerical operations)
  - `scikit-learn` (Machine learning models & evaluation)
  - `matplotlib` & `seaborn` (Data visualization)
  - `scipy` (Statistical analysis)

### Hardware Specifications:

- **Machine:** MacBook Air (M1 Chip)
- **CPU:** Apple M1 (8-core)
- **RAM:** 16GB Unified Memory

## Data Pre-processing

### Combining Maths dataset with Portuguese dataset

```
import pandas as pd

dfm = pd.read_csv("student/student-mat.csv", sep=";")
dfp = pd.read_csv("student/student-por.csv", sep=";")

dfm['subject'] = 'M'
dfp['subject'] = 'P'

df = pd.concat([dfm, dfp], ignore_index=True)
print(f"Total rows: {len(df)}")

Total rows: 1044
```

### Checking for data quality issues

#### Data Types:

```
df.info()

<class 'pandas.core.frame.DataFrame'>
RangeIndex: 1044 entries, 0 to 1043
Data columns (total 34 columns):
 #   Column          Non-Null Count  Dtype
---  -
 0   school          1044 non-null   object
 1   sex              1044 non-null   object
 2   age              1044 non-null   int64
 3   address         1044 non-null   object
 4   famsize         1044 non-null   object
 5   Pstatus         1044 non-null   object
 6   Medu            1044 non-null   int64
 7   Fedu            1044 non-null   int64
 8   Mjob            1044 non-null   object
 9   Fjob            1044 non-null   object
10  reason          1044 non-null   object
11  guardian        1044 non-null   object
12  traveltime      1044 non-null   int64
13  studytime       1044 non-null   int64
```

14	failures	1044	non-null	int64
15	schoolsup	1044	non-null	object
16	famsup	1044	non-null	object
17	paid	1044	non-null	object
18	activities	1044	non-null	object
19	nursery	1044	non-null	object
20	higher	1044	non-null	object
21	internet	1044	non-null	object
22	romantic	1044	non-null	object
23	famrel	1044	non-null	int64
24	freetime	1044	non-null	int64
25	goout	1044	non-null	int64
26	Dalc	1044	non-null	int64
27	Walc	1044	non-null	int64
28	health	1044	non-null	int64
29	absences	1044	non-null	int64
30	G1	1044	non-null	int64
31	G2	1044	non-null	int64
32	G3	1044	non-null	int64
33	subject	1044	non-null	object

dtypes: int64(16), object(18)  
memory usage: 277.4+ KB

### Checking for missing values

```
print(f"NA values: {df.isna().sum().sum()}")
print(f"Null values: {df.isnull().sum().sum()}")
print(f"Empty strings: {(df == '').sum().sum()}")
```

NA values: 0  
Null values: 0  
Empty strings: 0

### Checking for unique values for each column

```
for col in df.columns:
    unique_vals = sorted(df[col].unique())
    print(f"{col}: {unique_vals}")
```

school: ['GP', 'MS']  
sex: ['F', 'M']  
age: [np.int64(15), np.int64(16), np.int64(17), np.int64(18),  
np.int64(19), np.int64(20), np.int64(21), np.int64(22)]  
address: ['R', 'U']  
famsize: ['GT3', 'LE3']  
Pstatus: ['A', 'T']  
Medu: [np.int64(0), np.int64(1), np.int64(2), np.int64(3),  
np.int64(4)]  
Fedu: [np.int64(0), np.int64(1), np.int64(2), np.int64(3),

```
np.int64(4)]
Mjob: ['at_home', 'health', 'other', 'services', 'teacher']
Fjob: ['at_home', 'health', 'other', 'services', 'teacher']
reason: ['course', 'home', 'other', 'reputation']
guardian: ['father', 'mother', 'other']
traveltime: [np.int64(1), np.int64(2), np.int64(3), np.int64(4)]
studytime: [np.int64(1), np.int64(2), np.int64(3), np.int64(4)]
failures: [np.int64(0), np.int64(1), np.int64(2), np.int64(3)]
schoolsup: ['no', 'yes']
famsup: ['no', 'yes']
paid: ['no', 'yes']
activities: ['no', 'yes']
nursery: ['no', 'yes']
higher: ['no', 'yes']
internet: ['no', 'yes']
romantic: ['no', 'yes']
famrel: [np.int64(1), np.int64(2), np.int64(3), np.int64(4),
np.int64(5)]
freetime: [np.int64(1), np.int64(2), np.int64(3), np.int64(4),
np.int64(5)]
goout: [np.int64(1), np.int64(2), np.int64(3), np.int64(4),
np.int64(5)]
Dalc: [np.int64(1), np.int64(2), np.int64(3), np.int64(4),
np.int64(5)]
Walc: [np.int64(1), np.int64(2), np.int64(3), np.int64(4),
np.int64(5)]
health: [np.int64(1), np.int64(2), np.int64(3), np.int64(4),
np.int64(5)]
absences: [np.int64(0), np.int64(1), np.int64(2), np.int64(3),
np.int64(4), np.int64(5), np.int64(6), np.int64(7), np.int64(8),
np.int64(9), np.int64(10), np.int64(11), np.int64(12), np.int64(13),
np.int64(14), np.int64(15), np.int64(16), np.int64(17), np.int64(18),
np.int64(19), np.int64(20), np.int64(21), np.int64(22), np.int64(23),
np.int64(24), np.int64(25), np.int64(26), np.int64(28), np.int64(30),
np.int64(32), np.int64(38), np.int64(40), np.int64(54), np.int64(56),
np.int64(75)]
G1: [np.int64(0), np.int64(3), np.int64(4), np.int64(5), np.int64(6),
np.int64(7), np.int64(8), np.int64(9), np.int64(10), np.int64(11),
np.int64(12), np.int64(13), np.int64(14), np.int64(15), np.int64(16),
np.int64(17), np.int64(18), np.int64(19)]
G2: [np.int64(0), np.int64(4), np.int64(5), np.int64(6), np.int64(7),
np.int64(8), np.int64(9), np.int64(10), np.int64(11), np.int64(12),
np.int64(13), np.int64(14), np.int64(15), np.int64(16), np.int64(17),
np.int64(18), np.int64(19)]
G3: [np.int64(0), np.int64(1), np.int64(4), np.int64(5), np.int64(6),
np.int64(7), np.int64(8), np.int64(9), np.int64(10), np.int64(11),
np.int64(12), np.int64(13), np.int64(14), np.int64(15), np.int64(16),
np.int64(17), np.int64(18), np.int64(19), np.int64(20)]
subject: ['M', 'P']
```

**Conclusion:** There seems to be no missing data, or incorrect data types. Boolean variables like: schoolsup, internet, etc could be changed to 0 and 1, but we are going to keep them like this for now.

## Visualising the Target Distribution

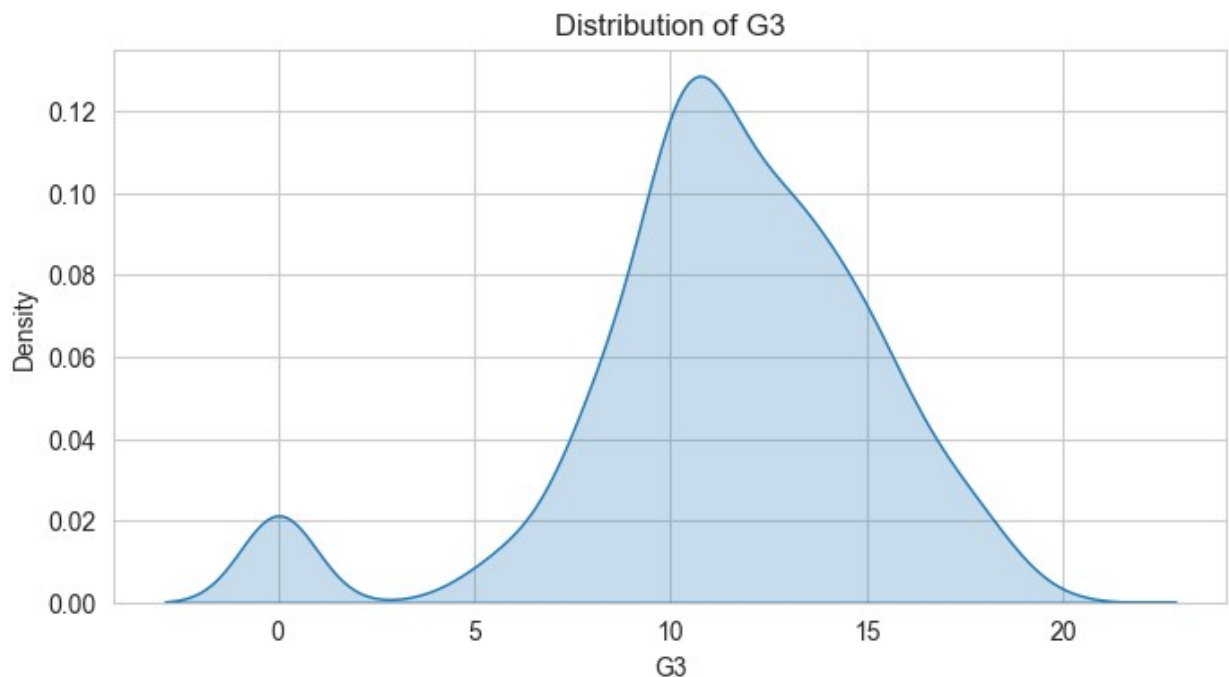
The boxplot shows a few outliers. This suggests that using any of MSE, RMSE and MAE should be fine for our evaluation metrics.

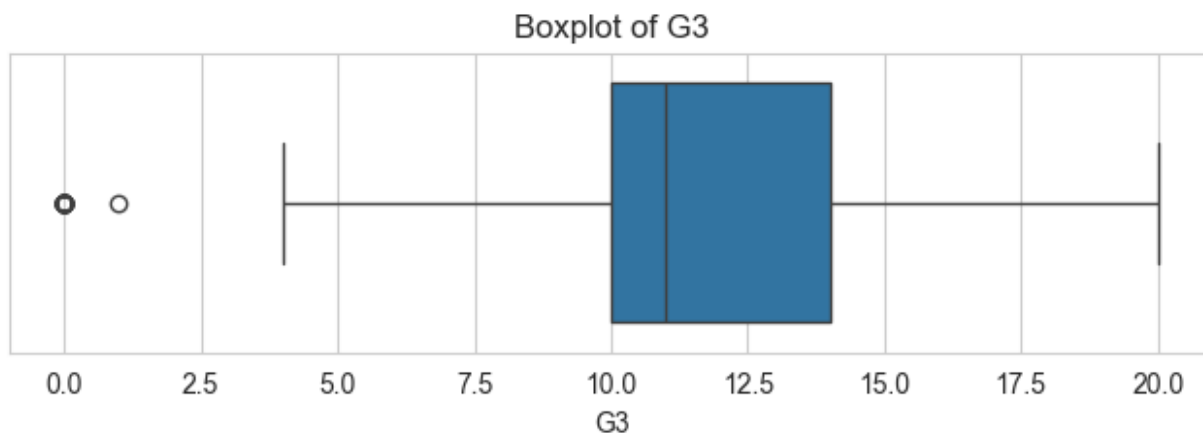
```
import matplotlib.pyplot as plt
import seaborn as sns

sns.set_style("whitegrid")

plt.figure(figsize=(8, 4))
sns.kdeplot(df['G3'], fill=True)
plt.title("Distribution of G3")
plt.show()

plt.figure(figsize=(8, 2))
sns.boxplot(x=df['G3'])
plt.title("Boxplot of G3")
plt.show()
```





## Data Encoding

We transformed our categorical data into labels with label encoding method and one hot encoding to select features to train our svm model.

```
from sklearn.preprocessing import LabelEncoder
```

```
df_lbEnc = df.copy()
```

```
label_encoders = {}
```

```
for col in df_lbEnc.select_dtypes(include=['object']).columns:
```

```
    le = LabelEncoder()
```

```
    df_lbEnc[col] = le.fit_transform(df_lbEnc[col])
```

```
    label_encoders[col] = le
```

```
df_lbEnc.head()
```

	school	sex	age	address	famsize	Pstatus	Medu	Fedu	Mjob	Fjob
...	\									
0	0	0	18	1	0	0	4	4	0	4
...										
1	0	0	17	1	0	1	1	1	0	2
...										
2	0	0	15	1	1	1	1	1	0	2
...										
3	0	0	15	1	0	1	4	2	1	3
...										
4	0	0	16	1	0	1	3	3	2	2
...										

	freetime	goout	Dalc	Walc	health	absences	G1	G2	G3	subject
0	3	4	1	1	3	6	5	6	6	0
1	3	3	1	1	3	4	5	5	6	0

2	3	2	2	3	3	10	7	8	10	0
3	2	2	1	1	5	2	15	14	15	0
4	3	2	1	2	5	4	6	10	10	0

[5 rows x 34 columns]

## One Hot Encoding

```
df_oneHotEnc = df.copy()
df_oneHotEnc = pd.get_dummies(df, drop_first=True)
df_oneHotEnc.head()
```

	age	Medu	Fedu	traveltime	studytime	failures	famrel	freetime
0	18	4	4	2	2	0	4	3
1	17	1	1	1	2	0	5	3
2	15	1	1	1	2	3	4	3
3	15	4	2	1	3	0	3	2
4	16	3	3	1	2	0	4	3

	Dalc	...	guardian_other	schoolsup_yes	famsup_yes	paid_yes
0	1	...	False	True	False	False
1	1	...	False	False	True	False
2	2	...	False	True	False	True
3	1	...	False	False	True	True
4	1	...	False	False	True	True

	activities_yes	nursery_yes	higher_yes	internet_yes	romantic_yes
0	False	True	True	False	False
1	False	False	True	True	False
2	False	True	True	True	False
3	True	True	True	True	True
4	False	True	True	False	False

	subject_P
0	False
1	False



```
2      False
3      False
4      False
```

```
[5 rows x 43 columns]
```

## Feature Selection

We analysed the pearson's correlation of G3 with other features, and picked features with correlation greater than 0.3. We tried it out for both one hot encoding and label encoding and found three features G1, G2 and failures with high correlation to G3.

```
# Calculate correlations for label encoded data (excluding G3 itself)
corr_lbEnc =
df_lbEnc.drop(columns=['G3']).corrwith(df_lbEnc['G3']).abs().sort_valu
es(ascending=False)
print("Label Encoding - Features with |correlation| >= 0.3:")
print(corr_lbEnc[corr_lbEnc >= 0.3])

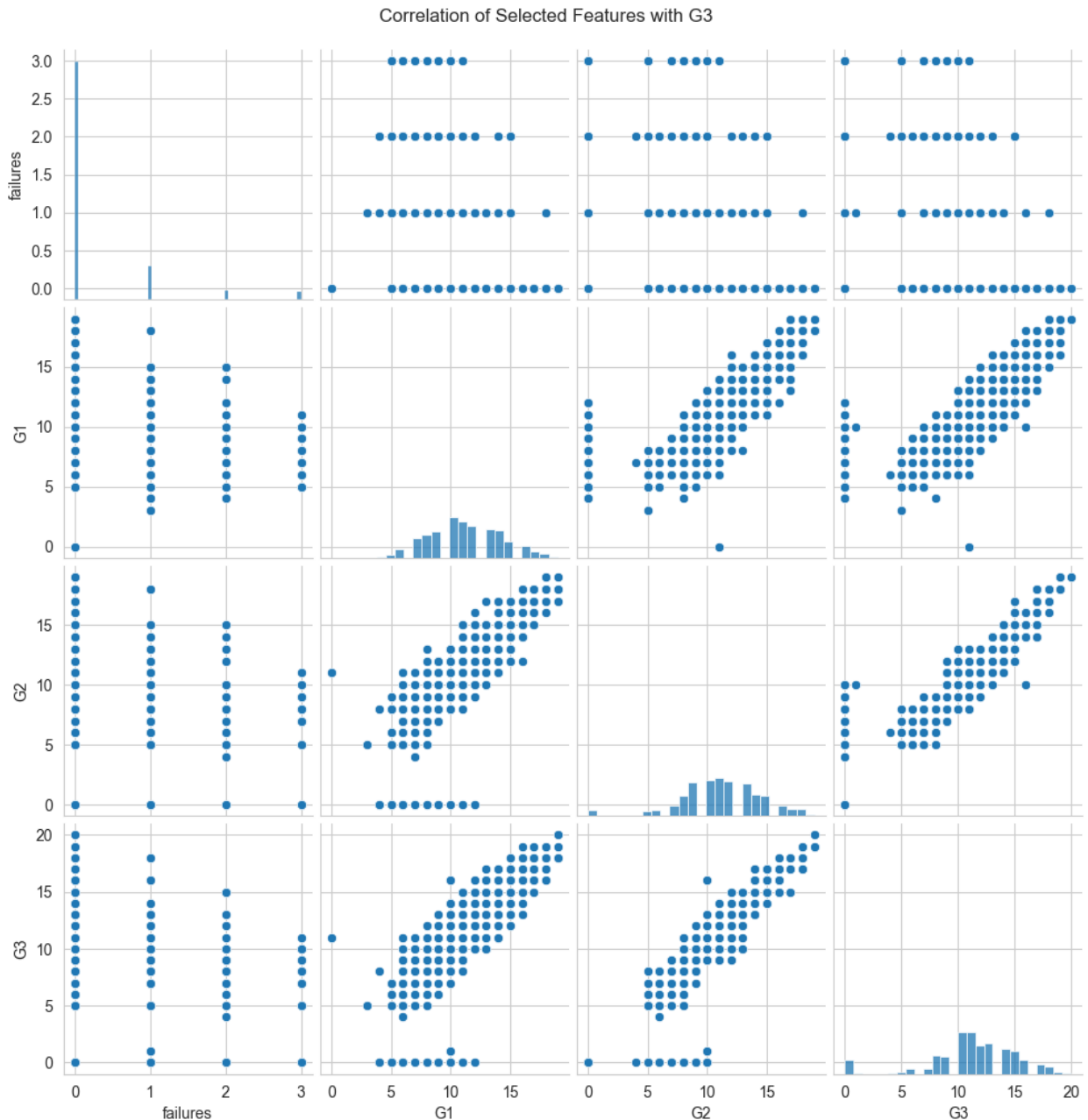
# Calculate correlations for one-hot encoded data (excluding G3
itself)
corr_oneHotEnc =
df_oneHotEnc.drop(columns=['G3']).corrwith(df_oneHotEnc['G3']).abs().s
ort_values(ascending=False)
print("\nOne-Hot Encoding - Features with |correlation| >= 0.3:")
print(corr_oneHotEnc[corr_oneHotEnc >= 0.3])

Label Encoding - Features with |correlation| >= 0.3:
G2          0.910743
G1          0.809142
failures    0.383145
dtype: float64

One-Hot Encoding - Features with |correlation| >= 0.3:
G2          0.910743
G1          0.809142
failures    0.383145
dtype: float64
```

This plot shows the correlation of the three features with G3.

```
sns.pairplot(df_lbEnc[['failures', 'G1', 'G2', 'G3']])
plt.suptitle("Correlation of Selected Features with G3", y=1.02)
plt.show()
```



## Filtering out the important features

We select those 3 feature columns (G1, G2, and failures) and scale the feature values.

### Column Filtering

```
df_lbEnc_top3corr = df_lbEnc[['failures', 'G1', 'G2', 'G3']]
df_lbEnc_top3corr.head()
```

	failures	G1	G2	G3
0	0	5	6	6

1	0	5	5	6
2	3	7	8	10
3	0	15	14	15
4	0	6	10	10

## Scaling

```
from sklearn.preprocessing import StandardScaler

X = df_lbEnc_top3corr[['failures', 'G1', 'G2']]
y = df_lbEnc_top3corr['G3']

scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)

df_scaled = pd.DataFrame(X_scaled, columns=['failures', 'G1', 'G2'])
df_scaled['G3'] = y.values
df_scaled.head()
```

	failures	G1	G2	G3
0	-0.403106	-2.083727	-1.597738	6
1	-0.403106	-2.083727	-1.902291	6
2	4.171268	-1.413029	-0.988631	10
3	-0.403106	1.269766	0.838688	15
4	-0.403106	-1.748378	-0.379525	10

## Train test split

We split out dataset into 70% training and 30% testing.

```
from sklearn.model_selection import train_test_split

SEED = 1234

X_train, X_test, y_train, y_test = train_test_split(
    df_scaled[['failures', 'G1', 'G2']],
    df_scaled['G3'],
    test_size=0.3,
    random_state=SEED
)

print(f"Training set size: {len(X_train)}")
print(f"Testing set size: {len(X_test)}")
```

```
Training set size: 730
Testing set size: 314
```

# Internal Evaluation

## Performance Estimation for Internal Evaluation

For our internal evaluation and finding the best parameters we first check how the models perform on the training dataset. We used lm, svm, random forest and decision tree with various hyperparameters. We used MAE, RMSE and MSE for evaluation, 10 fold cross validation with 2 repetitions.

```
import numpy as np
from sklearn.model_selection import RepeatedKFold, cross_validate
from sklearn.linear_model import LinearRegression
from sklearn.svm import SVR
from sklearn.tree import DecisionTreeRegressor
from sklearn.ensemble import RandomForestRegressor

models = {
    'LinearRegression': {
        'model': LinearRegression(),
        'params': {}
    },
    'SVR': {
        'model': SVR(),
        'params': {
            'C': [2, 4],
            'gamma': [0.01, 0.02, 0.04],
            'kernel': ['linear', 'poly', 'rbf']
        }
    },
    'DecisionTree': {
        'model': DecisionTreeRegressor(random_state=SEED),
        'params': {
            'min_samples_split': [10, 50],
            'ccp_alpha': [0.01, 0.03]
        }
    },
    'RandomForest': {
        'model': RandomForestRegressor(random_state=SEED),
        'params': {
            'n_estimators': [250, 500],
            'max_features': [1, 3]
        }
    }
}

scoring = {
    'mae': 'neg_mean_absolute_error',
    'mse': 'neg_mean_squared_error',
    'rmse': 'neg_root_mean_squared_error'
}
```

```

}

rkf = RepeatedKfold(n_splits=10, n_repeats=2, random_state=SEED)

results = {}
for name, config in models.items():
    model = config['model']
    param_grid = config['params']

    print(f"\nEvaluating {name}...")

    if name == 'LinearRegression':
        cv_results = cross_validate(
            model, X_train, y_train,
            scoring=scoring,
            cv=rkf,
            return_train_score=False
        )
        results[name] = {
            'mae': -np.mean(cv_results['test_mae']),
            'mse': -np.mean(cv_results['test_mse']),
            'rmse': -np.mean(cv_results['test_rmse'])
        }
    else:
        from sklearn.model_selection import GridSearchCV

        grid = GridSearchCV(
            estimator=model,
            param_grid=param_grid,
            scoring=scoring,
            refit='mae',
            cv=rkf,
            return_train_score=False
        )
        grid.fit(X_train, y_train)

        results[name] = {
            'mae': -grid.best_score_,
            'mse': -grid.cv_results_['mean_test_mse']
[grid.best_index_],
            'rmse': -grid.cv_results_['mean_test_rmse']
[grid.best_index_],
            'best_params': grid.best_params_
        }

print("\nInternal Evaluation Results:")
for model, scores in results.items():
    print(f"\n{model}:")
    for metric, value in scores.items():
        if metric != 'best_params':

```

```

        print(f"    {metric.upper()}: {value:.4f}")
    else:
        print(f"    Best Parameters: {value}")

# Alternative more sophisticated visualization
plt.figure(figsize=(12, 8))

# Prepare data in long format for seaborn
plot_data = []
for model in results:
    for metric in ['mae', 'mse', 'rmse']:
        plot_data.append({
            'Model': model,
            'Metric': metric.upper(),
            'Score': results[model][metric]
        })
plot_df = pd.DataFrame(plot_data)

# Create plot
sns.set_style("whitegrid")
ax = sns.barplot(x='Model', y='Score', hue='Metric', data=plot_df,
palette='tab10')

# Add value labels
for p in ax.patches:
    ax.annotate(f"{p.get_height():.3f}",
                (p.get_x() + p.get_width() / 2., p.get_height()),
                ha='center', va='center', xytext=(0, 10),
                textcoords='offset points')

# Customize plot
plt.title('Model Performance Comparison', pad=20)
plt.xlabel('')
plt.ylabel('Score')
plt.legend(title='Metric', bbox_to_anchor=(1.05, 1), loc='upper left')
plt.xticks(rotation=45)
plt.tight_layout()
plt.show()

```

Evaluating LinearRegression...

Evaluating SVR...

Evaluating DecisionTree...

Evaluating RandomForest...

Internal Evaluation Results:

#### LinearRegression:

MAE: 0.9483

MSE: 2.6489

RMSE: 1.5936

#### SVR:

MAE: 0.8890

MSE: 2.7490

RMSE: 1.6193

Best Parameters: {'C': 2, 'gamma': 0.01, 'kernel': 'linear'}

#### DecisionTree:

MAE: 0.9968

MSE: 2.8841

RMSE: 1.6556

Best Parameters: {'ccp\_alpha': 0.03, 'min\_samples\_split': 10}

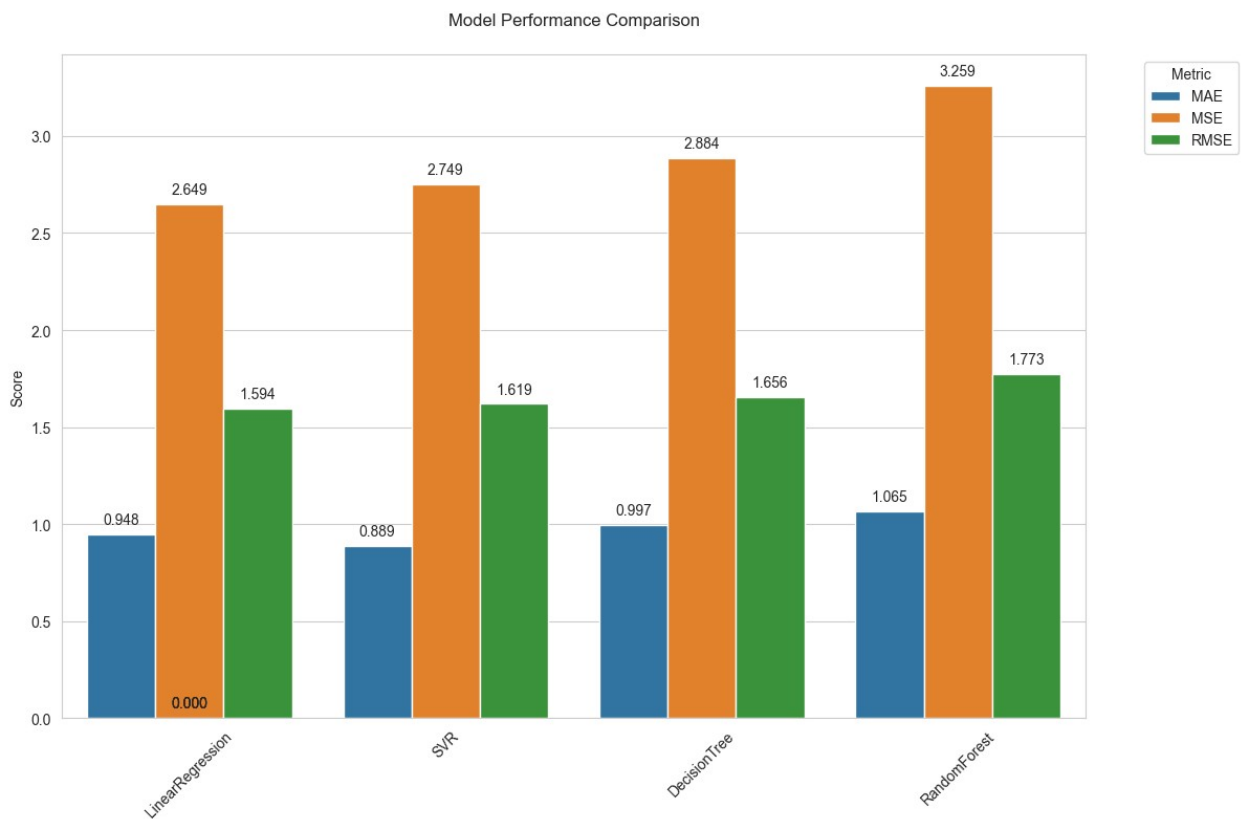
#### RandomForest:

MAE: 1.0649

MSE: 3.2588

RMSE: 1.7726

Best Parameters: {'max\_features': 3, 'n\_estimators': 500}



## Top Performer for Internal Evaluation

Our top performer for the internal evaluation was the linear regression model (lm) for all the three evaluation metrics.

```
best_model = min(results, key=lambda x: results[x]['rmse'])
print(f"Top performer: {best_model} with RMSE: {results[best_model]
['rmse']:.4f}")
```

Top performer: LinearRegression with RMSE: 1.5936

## External Evaluation

We finally train and test our linear regression model.

```
final_model = LinearRegression()
final_model.fit(X_train, y_train)

from sklearn.metrics import mean_absolute_error

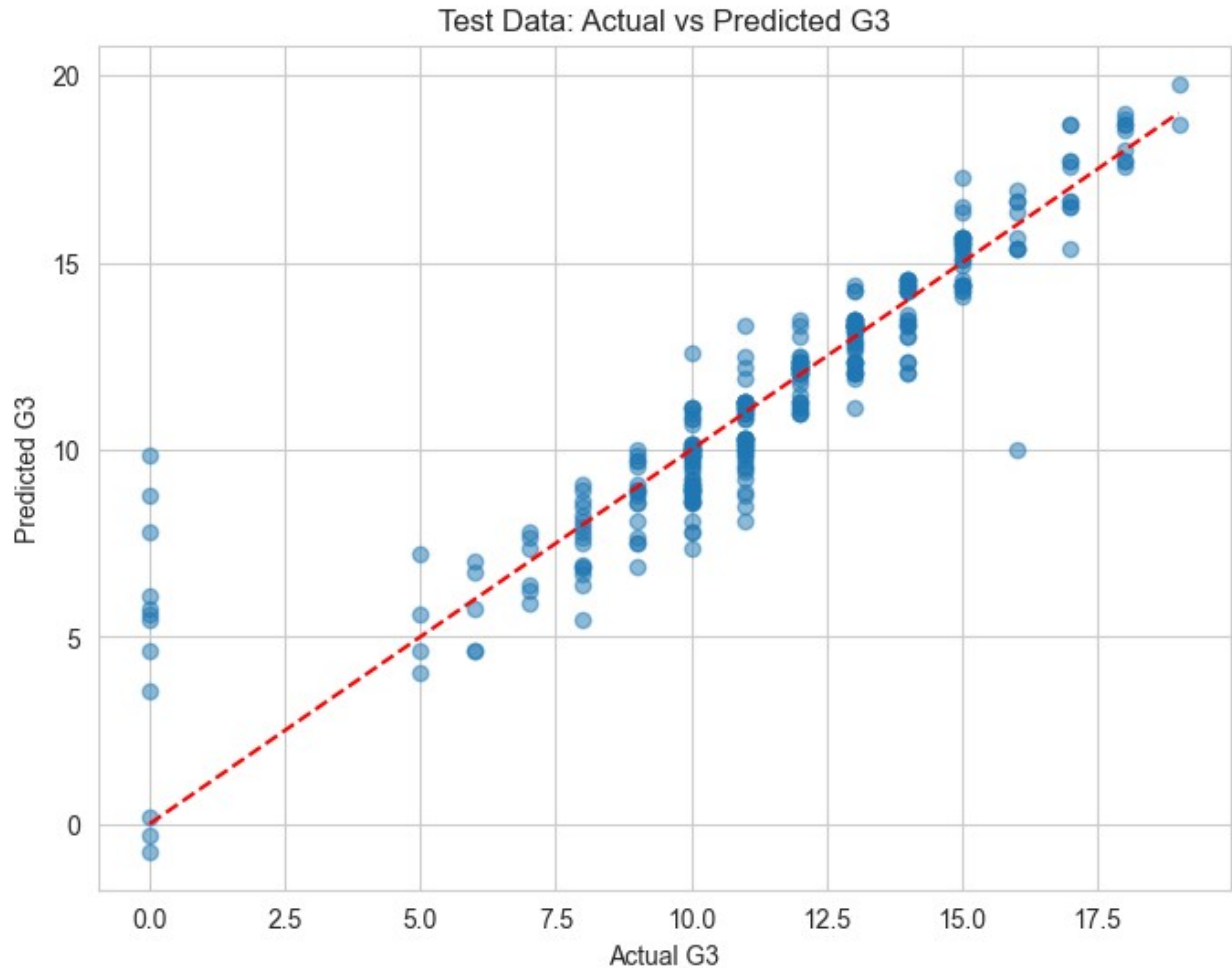
y_pred = final_model.predict(X_test)
mae_test = mean_absolute_error(y_test, y_pred)
print(f"Test MAE: {mae_test:.4f}")
```

Test MAE: 0.8861

The graph shows the relation between the predicted G3 and actual G3 for test data.

```
plt.figure(figsize=(8, 6))
plt.scatter(y_test, y_pred, alpha=0.5)
plt.plot([y_test.min(), y_test.max()], [y_test.min(), y_test.max()],
'r--')
plt.xlabel('Actual G3')
plt.ylabel('Predicted G3')
plt.title('Test Data: Actual vs Predicted G3')
plt.show()
```





## Conclusion

In conclusion, our **linear regression model** demonstrates strong accuracy in predicting a student's G3 grade when provided with G1, G2, and failure data.

### Key Findings:

- **MAE (Mean Absolute Error):**
  - Train: **0.9483**
  - Test: **0.8861**
- Since G3 ranges from **0 to 20**, an error of **~0.9** is negligible.
- Consistent performance across train/test sets suggests **no overfitting or underfitting**.

## Final Thoughts:

- Linear regression provides the best **balance of simplicity and accuracy** for this task.
- Alternative models (e.g., SVR) could improve with **more data or feature engineering**.
- Future work: Test robustness on larger datasets or with additional features.