

Predicting Student Performance in High School Math

This program uses machine learning regression and classification algorithms to predict the final math scores of students from two Portuguese schools, based on various demographic and social factors. Using multiple linear regression and decision tree classification, outputs include the identification of important features, results of linearity testing, and model accuracy in predicting on new data.

Data Source: University of California, Irvine (UCI) Machine Learning Repository.

Please view the README for more details.

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```
In [1]: #Import the necessary packages

import pandas as pd
import numpy as np
import seaborn as sns
import matplotlib.pyplot as plt
import math
import xgboost as xgb
import statsmodels.api as sm

from scipy import stats
from sklearn.linear_model import LinearRegression
from sklearn.model_selection import train_test_split
from sklearn.metrics import r2_score
from sklearn import datasets
from sklearn.ensemble import RandomForestRegressor
from sklearn.metrics import mean_absolute_error
from sklearn.metrics import mean_squared_error
from sklearn import linear_model
from itertools import combinations
from sklearn.svm import SVC
from sklearn.metrics import accuracy_score
from sklearn.ensemble import RandomForestClassifier
from sklearn.decomposition import PCA
from sklearn.metrics import classification_report
from sklearn.metrics import confusion_matrix
```

```
In [2]: # Read in the student mathematics grades csv files as a data frame
math = pd.read_csv('student-mat.csv')
```

```
In [3]: # Display the first five rows of the student grades data frame
print(math.head())

# Display the column names in the student grades data frame
print(math.columns)
```

```
school sex  age address famsize Pstatus  Medu  Fedu  Mjob  Fjob  ...  \
```

0	GP	F	18	U	GT3	A	4	4	at_home	teacher	...
1	GP	F	17	U	GT3	T	1	1	at_home	other	...
2	GP	F	15	U	LE3	T	1	1	at_home	other	...
3	GP	F	15	U	GT3	T	4	2	health	services	...
4	GP	F	16	U	GT3	T	3	3	other	other	...

	famrel	freetime	goout	Dalc	Walc	health	absences	G1	G2	G3
0	4		3	4	1	1	3	6	5	6
1	5		3	3	1	1	3	4	5	5
2	4		3	2	2	3	3	10	7	8
3	3		2	2	1	1	5	2	15	14
4	4		3	2	1	2	5	4	6	10

```
[5 rows x 33 columns]
Index(['school', 'sex', 'age', 'address', 'famsize', 'Pstatus', 'Medu', 'Fedu',
      'Mjob', 'Fjob', 'reason', 'guardian', 'traveltime', 'studytime',
      'failures', 'schoolsup', 'famsup', 'paid', 'activities', 'nursery',
      'higher', 'internet', 'romantic', 'famrel', 'freetime', 'goout', 'Dalc',
      'Walc', 'health', 'absences', 'G1', 'G2', 'G3'],
      dtype='object')
```

```
In [4]: # Display a summary of the data
math.info()

# Data contains 33 columns including both categorical and numerical data. All columns c
# Observe that the data types are not consistent for all variables (object and integer)
```

```
<class 'pandas.core.frame.DataFrame'>
RangeIndex: 395 entries, 0 to 394
Data columns (total 33 columns):
#   Column          Non-Null Count  Dtype
---  -
0   school          395 non-null   object
1   sex             395 non-null   object
2   age            395 non-null   int64
3   address         395 non-null   object
4   famsize         395 non-null   object
5   Pstatus         395 non-null   object
6   Medu            395 non-null   int64
7   Fedu            395 non-null   int64
8   Mjob            395 non-null   object
9   Fjob            395 non-null   object
10  reason          395 non-null   object
11  guardian        395 non-null   object
12  traveltime      395 non-null   int64
13  studytime       395 non-null   int64
14  failures        395 non-null   int64
15  schoolsup       395 non-null   object
16  famsup          395 non-null   object
17  paid            395 non-null   object
18  activities       395 non-null   object
19  nursery         395 non-null   object
20  higher          395 non-null   object
21  internet        395 non-null   object
22  romantic        395 non-null   object
23  famrel          395 non-null   int64
24  freetime        395 non-null   int64
25  goout           395 non-null   int64
26  Dalc            395 non-null   int64
27  Walc            395 non-null   int64
28  health          395 non-null   int64
29  absences        395 non-null   int64
30  G1              395 non-null   int64
31  G2              395 non-null   int64
```

```
32 G3          395 non-null    int64
dtypes: int64(16), object(17)
memory usage: 102.0+ KB
```

Map Categorical Variables to Numeric Values

```
In [5]: # Obtain names of the columns containing objects (categorical data)
cat_names=math.dtypes[math.dtypes=='object'].index

# Create a new dataframe named "cat_df" containing all of the categorical data of the m
cat_df=math[cat_names]

# Display the first five rows of the new "cat_df" data frame
print(cat_df.head())
```

	school	sex	address	famsize	Pstatus	Mjob	Fjob	reason	guardian	\
0	GP	F	U	GT3	A	at_home	teacher	course	mother	
1	GP	F	U	GT3	T	at_home	other	course	father	
2	GP	F	U	LE3	T	at_home	other	other	mother	
3	GP	F	U	GT3	T	health	services	home	mother	
4	GP	F	U	GT3	T	other	other	home	father	

	schoolsup	famsup	paid	activities	nursery	higher	internet	romantic
0	yes	no	no	no	yes	yes	no	no
1	no	yes	no	no	no	yes	yes	no
2	yes	no	yes	no	yes	yes	yes	no
3	no	yes	yes	yes	yes	yes	yes	yes
4	no	yes	yes	no	yes	yes	no	no

```
In [6]: # Initialize an empty list for the names of columns which contain binary categorical da
binary_list=[]

# Initialize a second empty list for the names of columns which contain non-binary cate
non_binary_list=[]

# Loop through all the columns in the "cat_df" data frame and if the column contains on
# add the name of that column to the binary list. Else, add the column name to the non
for i in cat_df.columns:
    if (len(cat_df[i].unique())>2):
        non_binary_list.append(i)

    else:
        binary_list.append(i)
```

```
In [7]: # Map symmetric binary categorical variables to numeric values

# Initialize an empty data frame named "new_num_df" which will store the newly mapped
# numerical data from the original "cat_df" categorical data
new_num_df = pd.DataFrame()
print(new_num_df)
```

```
Empty DataFrame
Columns: []
Index: []
```

```
In [8]: # Loop through the columns in the binary column names list
for i in range(0, len(binary_list)):
```

```

# Obtain the 2 unique values cooresponding which each column
Unique_Values = cat_df[binary_list[i]].unique()

# If parent's cohabitation status is T (living together) then assign a 1 to imply s
if binary_list[i] == 'Pstatus':

    if Unique_Values[0] == 'T':
        new_num_df[binary_list[i]] = cat_df[binary_list[i]].replace([Unique_Values[0]], 1)

    else:
        new_num_df[binary_list[i]] = cat_df[binary_list[i]].replace([Unique_Values[0]], 0)

# For all other columns, if it is a yes/no, assign 1 to yes to imply significance
elif Unique_Values[0] == 'yes':
    new_num_df[binary_list[i]] = cat_df[binary_list[i]].replace([Unique_Values[0]], 1)

# Assign 0 value to "no" observations
elif Unique_Values[0] == 'no':
    new_num_df[binary_list[i]] = cat_df[binary_list[i]].replace([Unique_Values[0]], 0)

# If the observed value is not yes/no, assign 0 and 1 at random (first unique value
else:
    new_num_df[binary_list[i]] = cat_df[binary_list[i]].replace([Unique_Values[0]], 0)

# Display the "new_num_df" to ensure all of the binary categorical data has been mapped
print(new_num_df)

```

	school	sex	address	famsize	Pstatus	schoolsup	famsup	paid	\
0	1	1	1	1	0	1	0	0	
1	1	1	1	1	1	0	1	0	
2	1	1	1	0	1	1	0	1	
3	1	1	1	1	1	0	1	1	
4	1	1	1	1	1	0	1	1	
..	
390	0	0	1	0	0	0	1	1	
391	0	0	1	0	1	0	0	0	
392	0	0	0	1	1	0	0	0	
393	0	0	0	0	1	0	0	0	
394	0	0	1	0	1	0	0	0	

	activities	nursery	higher	internet	romantic
0	0	1	1	0	0
1	0	0	1	1	0
2	0	1	1	1	0
3	1	1	1	1	1
4	0	1	1	0	0
..
390	0	1	1	0	0
391	0	0	1	1	0
392	0	0	1	0	0
393	0	0	1	1	0
394	0	1	1	1	0

[395 rows x 13 columns]

In [9]:

```

# Map multiclass categorical variables to numeric values

# Loop through the columns in the non-binary column names list
for i in range(0, len(non_binary_list)):

```

```

# Obtain multiple unique values corresponding to each column
Unique_Values = cat_df[non_binary_list[i]].unique()

# Assign numeric values from 0 - 4 randomly based on the total number of unique values
if len(Unique_Values) == 3:
    new_num_df[non_binary_list[i]] = cat_df[non_binary_list[i]].replace([Unique_Values[0], Unique_Values[1], Unique_Values[2]], [0, 1, 2])

elif len(Unique_Values) == 4:
    new_num_df[non_binary_list[i]] = cat_df[non_binary_list[i]].replace([Unique_Values[0], Unique_Values[1], Unique_Values[2], Unique_Values[3]], [0, 1, 2, 3])

elif len(Unique_Values) == 5:
    new_num_df[non_binary_list[i]] = cat_df[non_binary_list[i]].replace([Unique_Values[0], Unique_Values[1], Unique_Values[2], Unique_Values[3], Unique_Values[4]], [0, 1, 2, 3, 4])

```

In [10]:

```

# There is a function that can be used to transform categorical data into integers.

# from sklearn.preprocessing import LabelEncoder
# LabelEncoder().fit_transform()

# These functions were not used in this program because of the specific mapping that was required

```

In [11]:

```

# Create a data frame that contains all of the numeric data from the original "math" dataset
num_names = math.dtypes[math.dtypes == 'int64'].index
org_num_df = math[num_names]

# Display the first five rows of the new "org_num_df" data frame
print(org_num_df.head())

```

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

	Dalc	Walc	health	absences	G1	G2	G3
0	1	1	3	6	5	6	6
1	1	1	3	4	5	5	6
2	2	3	3	10	7	8	10
3	1	1	5	2	15	14	15
4	1	2	5	4	6	10	10

In [12]:

```

# Combine the original numeric data with the newly mapped numerical data into a dataframe
# containing all of the columns from the raw imported dataset
combined_table = pd.concat([org_num_df, new_num_df], axis=1)

# Display the first five rows of the new "combined_table" data frame
print(combined_table.head())

# Display the data types of all the columns to ensure all are now integers
print(combined_table.dtypes)

```

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

	Dalc	...	paid	activities	nursery	higher	internet	romantic	Mjob	\
0	1	...	0	0	1	1	0	0	0	
1	1	...	0	0	0	1	1	0	0	
2	2	...	1	0	1	1	1	0	0	
3	1	...	1	1	1	1	1	1	1	
4	1	...	1	0	1	1	0	0	2	

	Fjob	reason	guardian
0	0	0	0
1	1	0	1
2	1	1	0
3	2	2	0
4	1	2	1

[5 rows x 33 columns]

```

age                int64
Medu               int64
Fedu               int64
traveltime         int64
studytime          int64
failures           int64
famrel             int64
freetime           int64
goout              int64
Dalc               int64
Walc               int64
health             int64
absences           int64
G1                 int64
G2                 int64
G3                 int64
school             int64
sex                int64
address            int64
famsize            int64
Pstatus            int64
schoolsup          int64
famsup             int64
paid               int64
activities          int64
nursery            int64
higher             int64
internet           int64
romantic           int64
Mjob               int64
Fjob               int64
reason             int64
guardian           int64
dtype: object

```

Apply Multiple Linear Regression ML Model

```

In [13]: # Randomly split the data into 20% testing and 80% training
train, test = train_test_split(combined_table, test_size = 0.2, random_state = 2022)

# Display the dimensions of the training and testing data
print('Size of the train dataset is', train.shape)
print('Size of the test dataset is', test.shape)

# Split training data into x (independent variables) and y (dependent variable)

# Assign all of the columns except G3 (final mathematics grade from 0-20) to be x

```

```

x_train = train.loc[:, train.columns != 'G3']

# Assign G3 (final mathematics grade from 0-20) to be y
y_train = train['G3']

x_test = test.loc[:, test.columns != 'G3']
y_test = test['G3']

```

Size of the train dataset is (316, 33)
Size of the test dataset is (79, 33)

```

In [14]: #Plot pairwise scatter plot to analyze initial trends and highly coorelated variables

# This command takes a long time to execute. If you have the computing power,
# please uncomment the next line to see the pairplot.

#sns.pairplot(train)

```

```

In [15]: # Define a function for creating a correlation heatmap
def create_correlation_heatmap(data):
    _ , ax = plt.subplots(figsize =(40, 25))
    colormap = sns.diverging_palette(220, 10, as_cmap = True)

    sns.heatmap(
        data.corr(),
        cmap = colormap,
        square=True,
        cbar_kws={'shrink':.9 },
        ax=ax,
        annot=True,
        linewidths=0.1,vmax=1.0, linecolor='white',
        annot_kws={'fontsize':12 }
    )

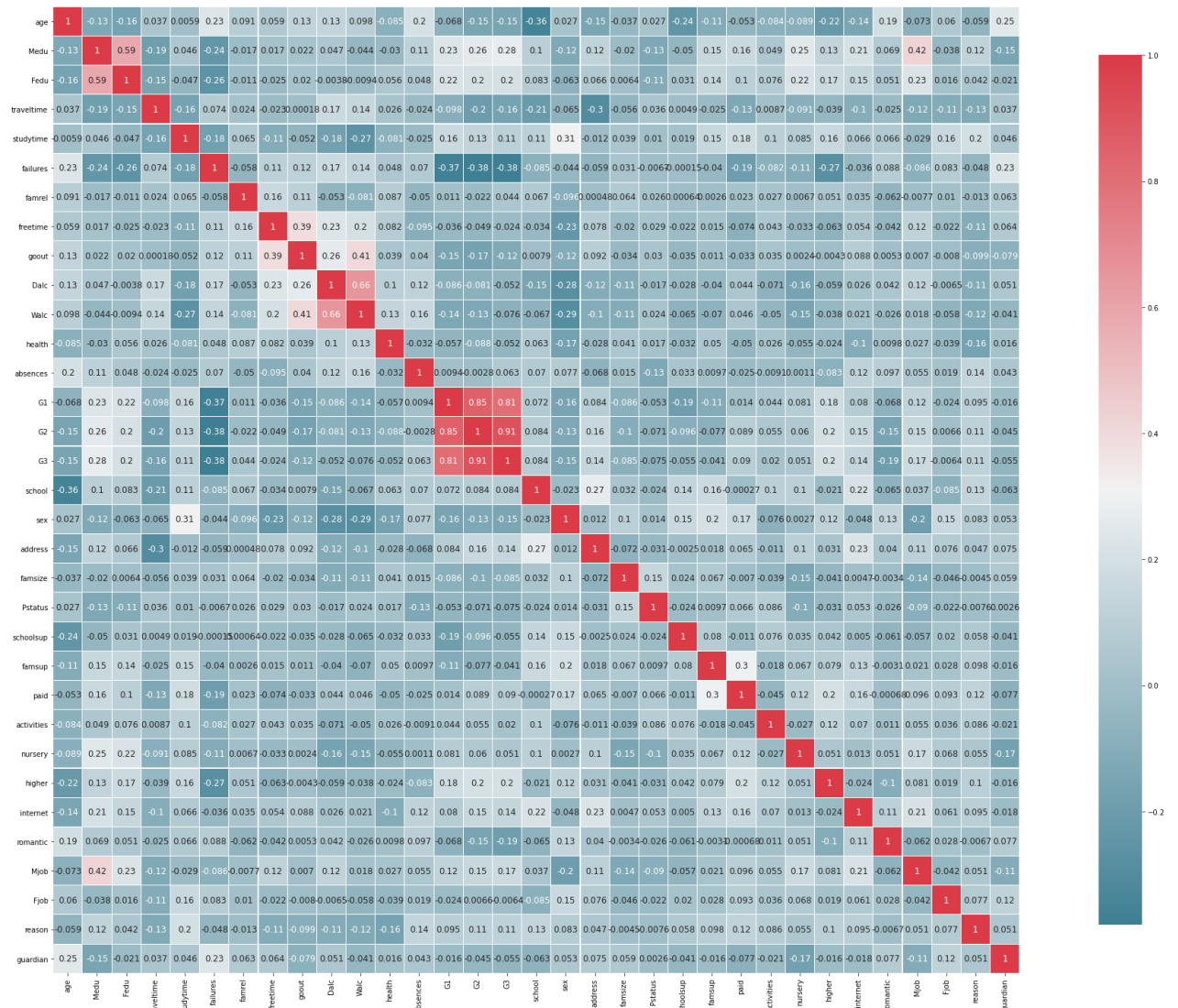
    plt.title('Pearson Correlation of Features', y=1.05, size=35)

# Plot the heatmap using the entire training dataset to analyze feature coorelation
create_correlation_heatmap(train)

# Observe a few instances of highly coorelated variables (coorelation >.5 or >-.5)

```

Pearson Correlation of Features



Mitigate Highly Coorelated Independent Variables

In [16]:

```
# It is important to have mitigate or remove highly coorelated variables from a linear
# highly coorelated variables indicate high model variance and can lead to an unreliable model

# G1 (first period grade) and G2 (second period grade) have a high positive coorelation
# Walc (weekend alcohol consumption) and Dalc (workday alcohol consumption) have a high
# Medu (mother's education) and Fedu (father's education) have a high positive coorelation

# To remove the negative impacts of high coorelation, a weighted average of each of the
# calculated and counted for as a new variable.
# G1 and G2 were combined into "Avg_G1&G2".
# Walc and Dalc were combined into "Avg_ALc".
# Medu and Fedu were combined into "Avg_Par_Edu".

# Create separate data frames containing the data for each highly coorelated variable
G1_df = combined_table['G1']
G2_df = combined_table['G2']
```



```

Walc_df = combined_table['Walc']
Dalc_df = combined_table['Dalc']
Medu_df = combined_table['Medu']
Fedu_df = combined_table['Fedu']

# Initialize empty lists for each new weighted average value to be added to
Avg_G12 = []
Avg_Alc = []
Avg_Par_Edu = []

# Loop through all the rows in the combined data set
for i in range(0, len(G1_df)):

    # Calculated the weighted average for each of the three highly coorelated sets sepa
    Avg_G_Value = int((G1_df[i] + G2_df[i]) / 2)
    Avg_Alc_Value = int(((5*Dalc_df[i]) + (2*Walc_df[i]))) / 7)
    Avg_Par_Edu_Value = int((Medu_df[i] + Fedu_df[i]) / 2)

    # Append the new average value to the respective weighted average list
    Avg_G12.append(Avg_G_Value)
    Avg_Alc.append(Avg_Alc_Value)
    Avg_Par_Edu.append(Avg_Par_Edu_Value)

# Add each list of newly averaged values to the respective empty data frame
Avg_G_Table = pd.DataFrame(Avg_G12)
Avg_Alc_Table = pd.DataFrame(Avg_Alc)
Avg_Par_Edu_Table = pd.DataFrame(Avg_Par_Edu)

# Rename the column of average values to the appropriate name for each separate data fr
Avg_G_Table.rename(columns = {0: 'Avg_G1&G2'}, inplace = True)
Avg_Alc_Table.rename(columns = {0: 'Avg_Alc'}, inplace = True)
Avg_Par_Edu_Table.rename(columns = {0: 'Avg_Par_Edu'}, inplace = True)

# Combine the original "combined_table" data frame used for the first linear regressio
# with the new data frames containing the weighted averages.
Averages_df = pd.concat((Avg_G_Table, Avg_Alc_Table, Avg_Par_Edu_Table, combined_table),

# Display the data types of each column of the new data frame containing the average va
# previously high coorelated variables to ensure they are all integers.
print(Averages_df.dtypes)

```

```

Avg_G1&G2      int64
Avg_Alc        int64
Avg_Par_Edu    int64
age            int64
Medu           int64
Fedu           int64
traveltime     int64
studytime      int64
failures       int64
famrel         int64
freetime       int64
goout          int64
Dalc           int64
Walc           int64
health         int64
absences       int64
G1             int64
G2             int64

```

```

G3          int64
school      int64
sex         int64
address     int64
famsize     int64
Pstatus     int64
schoolsup   int64
famsup      int64
paid        int64
activities  int64
nursery     int64
higher      int64
internet    int64
romantic    int64
Mjob        int64
Fjob        int64
reason      int64
guardian    int64
dtype: object

```

```

In [17]: # Create a new data frame called "Uncoorelated_df" containing only low coorelated varia
# removing the highly coorelated variables from the data frame and keeping the averaged
Uncoorelated_df = Averages_df.drop(['G2', 'G1', 'Fedu', 'Medu', 'Walc', 'Dalc'], axis=1)
print(Uncoorelated_df)

```

```

      Avg_G1&G2  Avg_Alc  Avg_Par_Edu  age  traveltime  studytime  failures  \
0              5        1            4   18           2           2           0
1              5        1            1   17           1           2           0
2              7        2            1   15           1           2           3
3             14        1            3   15           1           3           0
4              8        1            3   16           1           2           0
..          ...      ...          ...   ...          ...          ...          ...
390            9        4            2   20           1           2           2
391           15        3            2   17           2           1           0
392            9        3            1   21           1           1           3
393           11        3            2   18           3           1           0
394            8        3            1   19           1           1           0

```

```

      famrel  freetime  goout  ...  paid  activities  nursery  higher  \
0          4          3      4  ...    0           0          1         1
1          5          3      3  ...    0           0          0         1
2          4          3      2  ...    1           0          1         1
3          3          2      2  ...    1           1          1         1
4          4          3      2  ...    1           0          1         1
..          ...      ...      ...  ...    ...          ...          ...          ...
390         5          5      4  ...    1           0          1         1
391         2          4      5  ...    0           0          0         1
392         5          5      3  ...    0           0          0         1
393         4          4      1  ...    0           0          0         1
394         3          2      3  ...    0           0          1         1

```

```

      internet  romantic  Mjob  Fjob  reason  guardian
0            0          0     0     0       0          0
1            1          0     0     1       0          1
2            1          0     0     1       1          0
3            1          1     1     2       2          0
4            0          0     2     1       2          1
..          ...      ...      ...  ...    ...          ...
390           0          0     3     2       0          2
391           1          0     3     2       0          0
392           0          0     2     1       0          2
393           1          0     3     1       0          0
394           1          0     2     4       0          1

```

[395 rows x 30 columns]

In [18]:

```
# Perform a random split on the new uncoorelated dataset into 20% testing and 80% train
uncoor_train, uncoor_test = train_test_split(Uncoorelated_df, test_size = 0.2, random_s

# Display the dimensions of the new training and new testing data
print('Size of the train dataset is ', uncoor_train.shape)
print('Size of the test dataset is ', uncoor_test.shape)

# Split training data into x (independent variables) and y (dependent variable)

# Assign all of the columns except G3 (final mathematics grade from 0-20) to be the new
uncoor_x_train = uncoor_train.loc[:, uncoor_train.columns != 'G3']

# Assign G3 (final mathematics grade from 0-20) to be y
uncoor_y_train = uncoor_train['G3']

uncoor_x_test = uncoor_test.loc[:, uncoor_test.columns != 'G3']
uncoor_y_test = uncoor_test['G3']
```

Size of the train dataset is (316, 30)

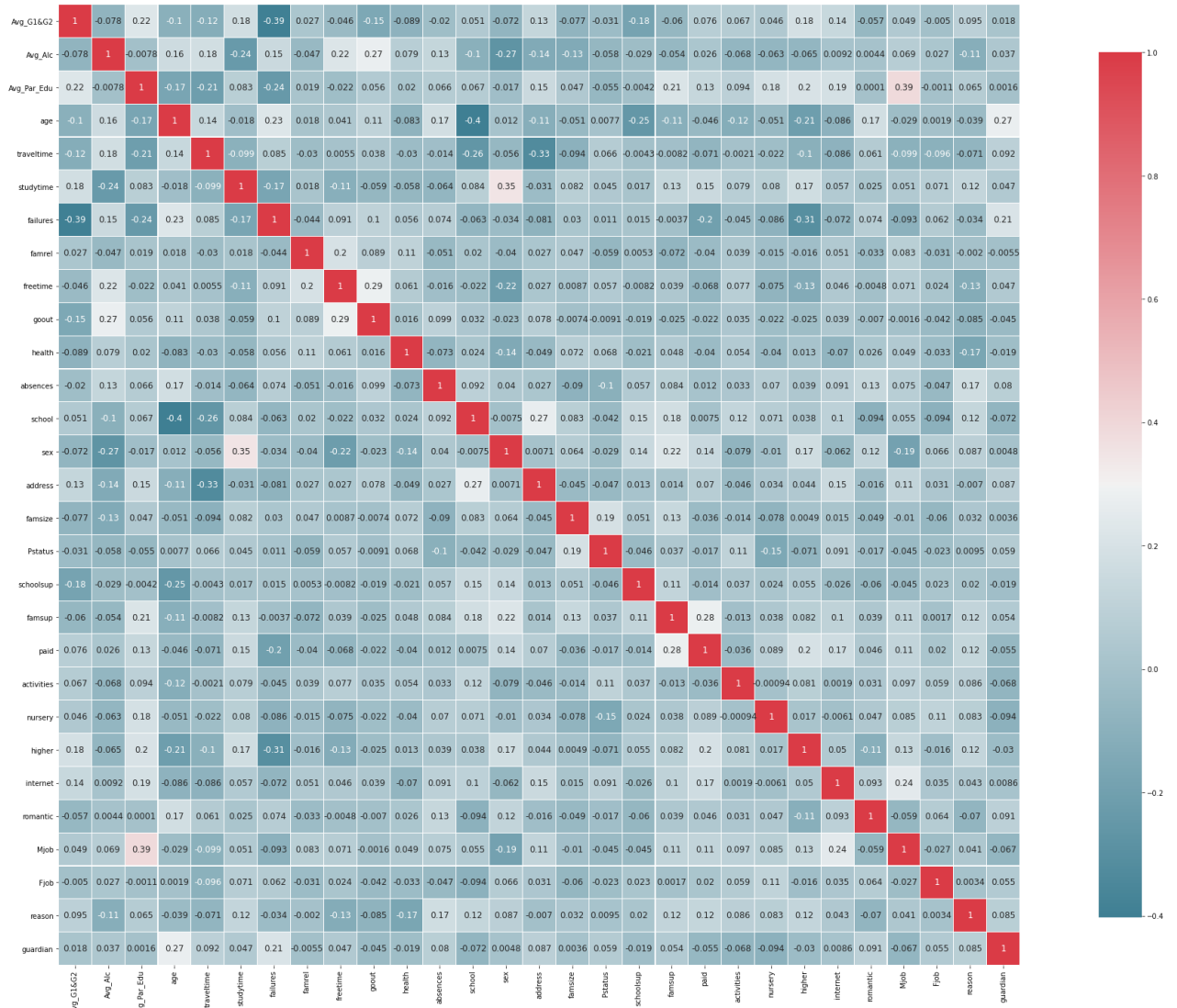
Size of the test dataset is (79, 30)

In [19]:

```
# Plot the new heatmap using the entire uncoorelated training dataset to analyze featur
create_correlation_heatmap(uncoor_x_train)

# Observe that there are no instances of extremly positive or negative coorelation
```

Pearson Correlation of Features



In [20]:

```
# Fit a Multiple Linear Regression (MLR) model to the training data
linear_regression = LinearRegression()
linear_regression.fit(uncoor_x_train, uncoor_y_train)

# Display the y intercept and coefficients of the MLR model
print('The intercept estimated by Sklearn is', round(linear_regression.intercept_,3))
print('The coefficients estimated by Sklearn are:',linear_regression.coef_)
```

The intercept estimated by Sklearn is 1.437
The coefficients estimated by Sklearn are: [1.23563601 0.03068752 -0.02571503 -0.2324223 -0.02530547 -0.24781729 -0.08524729 0.27445552 0.03787983 0.05116432 0.06472663 0.05501262 -0.5935353 -0.09798499 0.11610519 -0.02176417 -0.18447538 0.73082502 0.05520577 0.34549593 -0.31707647 -0.32967602 -0.2797912 -0.30169945 -0.38896107 0.00555177 -0.06748769 0.0661223 -0.23370326]

In [21]:

```
# Alternative method of calculating the model coefficients using the formula
inverse_matrix = np.linalg.inv(np.dot(uncoor_x_train.T, uncoor_x_train))
hat_matrix = np.dot(inverse_matrix, uncoor_x_train.T)
est_coeff = np.dot(hat_matrix, uncoor_y_train)
print('Coefficients estimated by the formula are:', est_coeff)
```

```
# Observe similar results to the Sklearn package estimated coefficients
```

```
Coefficients estimated by the formula are: [ 1.23987952e+00  3.08355865e-02 -1.20445645e-02 -1.76024328e-01
 2.42206348e-04 -2.49367300e-01 -7.67945158e-02  2.84753953e-01
 4.87940688e-02  4.80013977e-02  7.36685699e-02  5.33974715e-02
-4.88694869e-01 -9.14223008e-02  1.34767115e-01 -1.60982175e-02
-1.39264391e-01  7.74892439e-01  5.39200508e-02  3.44336574e-01
-3.18175775e-01 -3.15749372e-01 -1.68666193e-01 -2.83910625e-01
-3.94183815e-01  1.47245747e-03 -5.67397129e-02  6.92190640e-02
-2.63802418e-01]
```

In [22]:

```
#Check the significance of parameters by applying ordinary Least squares (OLS) regressi
lr = sm.OLS(uncoor_y_train, uncoor_x_train)
lr = lr.fit()

# Display the summary to check for model linearity and significant features
print(lr.summary())

# Observe an R^2 value of .969 and adjusted R^2 value of .966 (adjusting for non-signif
# Since R^2 is very close to 1, this indicates that the data fits one of the linearity
# Since the data is linear, the MLR model should predict new data with a fairly high ac

# Variables with a low p-value <.05 (on a 95% confidence interval) are classified as si
# Observing the output table, the significant variables include Avg_G1&G2(.000), age(.0
# famrel(.046), absences(.002), schoolsup(.033).

# The least significant variables from observing the table are Avg_Par_Edu (p = .931),
# famsize (p =.954), and Mjob (p =.99).
```

OLS Regression Results

```
=====
Dep. Variable:          G3      R-squared (uncentered):          0.969
Model:                  OLS      Adj. R-squared (uncentered):    0.966
Method:                 Least Squares      F-statistic:          308.9
Date:                  Sun, 13 Nov 2022      Prob (F-statistic):      2.83e-198
Time:                  10:52:38      Log-Likelihood:         -667.55
No. Observations:      316      AIC:                      1393.
Df Residuals:          287      BIC:                      1502.
Df Model:              29
Covariance Type:       nonrobust
=====
```

	coef	std err	t	P> t	[0.025	0.975]
Avg_G1&G2	1.2399	0.041	30.169	0.000	1.159	1.321
Avg_Alc	0.0308	0.156	0.198	0.843	-0.276	0.338
Avg_Par_Edu	-0.0120	0.139	-0.087	0.931	-0.285	0.261
age	-0.1760	0.063	-2.801	0.005	-0.300	-0.052
traveltime	0.0002	0.189	0.001	0.999	-0.372	0.372
studytime	-0.2494	0.163	-1.530	0.127	-0.570	0.071
failures	-0.0768	0.191	-0.403	0.687	-0.452	0.298
famrel	0.2848	0.142	2.006	0.046	0.005	0.564
freetime	0.0488	0.135	0.363	0.717	-0.216	0.314
goout	0.0480	0.121	0.398	0.691	-0.190	0.286
health	0.0737	0.087	0.843	0.400	-0.098	0.246
absences	0.0534	0.017	3.096	0.002	0.019	0.087
school	-0.4887	0.398	-1.226	0.221	-1.273	0.296
sex	-0.0914	0.294	-0.311	0.756	-0.669	0.487
address	0.1348	0.327	0.412	0.681	-0.509	0.779
famsize	-0.0161	0.279	-0.058	0.954	-0.565	0.533
Pstatus	-0.1393	0.453	-0.308	0.759	-1.031	0.752

schoolsup	0.7749	0.362	2.140	0.033	0.062	1.488
famsup	0.0539	0.285	0.189	0.850	-0.507	0.614
paid	0.3443	0.265	1.301	0.194	-0.176	0.865
activities	-0.3182	0.252	-1.261	0.208	-0.815	0.179
nursery	-0.3157	0.311	-1.017	0.310	-0.927	0.296
higher	-0.1687	0.561	-0.301	0.764	-1.272	0.935
internet	-0.2839	0.353	-0.803	0.423	-0.980	0.412
romantic	-0.3942	0.270	-1.462	0.145	-0.925	0.137
Mjob	0.0015	0.115	0.013	0.990	-0.224	0.227
Fjob	-0.0567	0.139	-0.409	0.683	-0.330	0.216
reason	0.0692	0.104	0.667	0.505	-0.135	0.273
guardian	-0.2638	0.205	-1.286	0.200	-0.668	0.140

Omnibus:	80.037	Durbin-Watson:	2.018
Prob(Omnibus):	0.000	Jarque-Bera (JB):	176.692
Skew:	-1.260	Prob(JB):	4.28e-39
Kurtosis:	5.659	Cond. No.	109.

Notes:

[1] R^2 is computed without centering (uncentered) since the model does not contain a constant.

[2] Standard Errors assume that the covariance matrix of the errors is correctly specified.

```
In [23]: # Predict new instances of student's grades in math (y) using the x testing data and a
y_predict = linear_regression.predict(uncoor_x_test)
```

```
In [24]: # Create a quantile-quantile plot to assess the second linearity assumption
stats.probplot(uncoor_y_train - linear_regression.predict(uncoor_x_train), dist="norm",

# Observe that the data falls closely to the red line (with some variability on the low
# indicating the normality assumption for linear regression is met.
```

```
Out[24]: ((array([-2.84925316, -2.55442126, -2.38761145, -2.26880689, -2.1753769 ,
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```

```

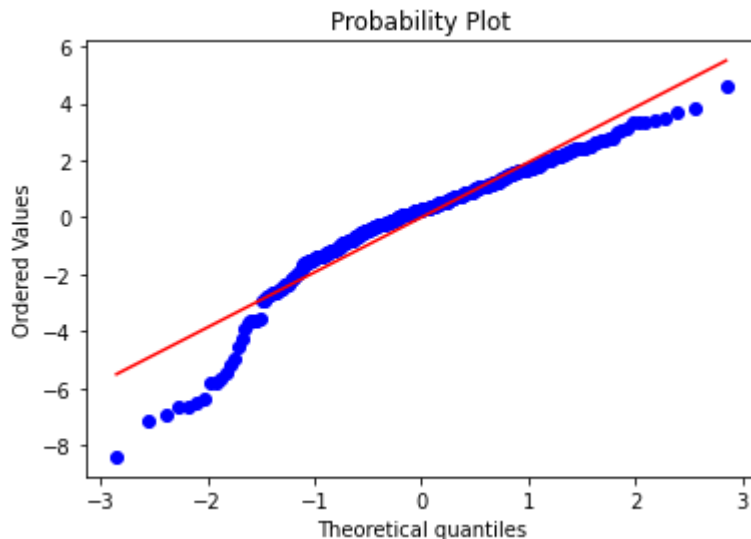
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```



```

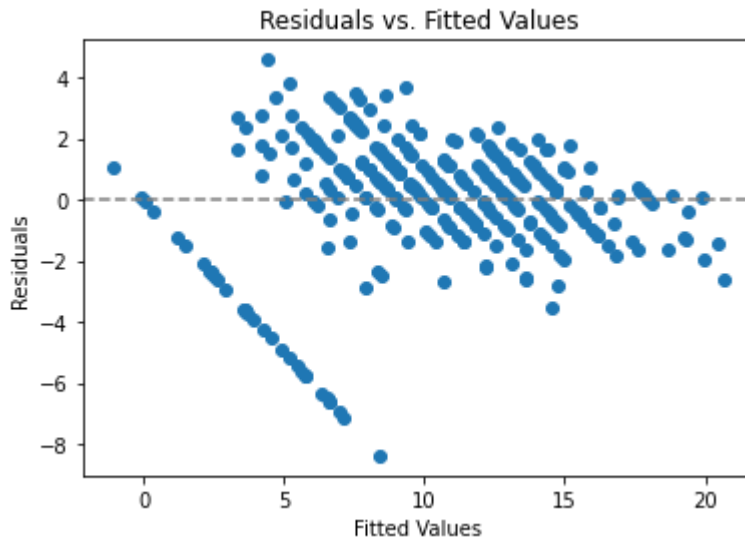
In [25]: # Plot the residuals vs fitted values to analyze the linear model assumption
residuals = uncoor_y_train - linear_regression.predict(uncoor_x_train)
plt.scatter(linear_regression.predict(uncoor_x_train), residuals)
plt.xlabel('Fitted Values')
plt.ylabel('Residuals')
plt.title('Residuals vs. Fitted Values')
plt.axhline(y = 0, color = 'grey', ls = '--')

```



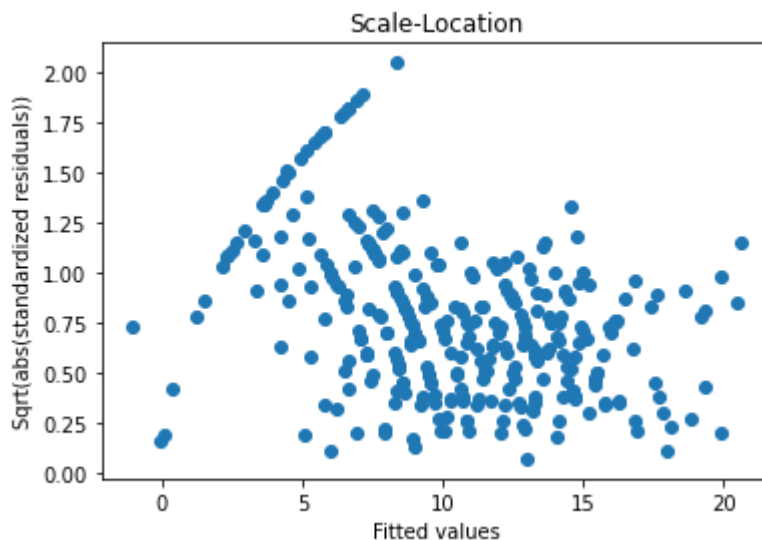
```
# Observe that the residuals are mostly spread along the horizontal line and no distinct  
# linear model assumption is met.
```

Out[25]: <matplotlib.lines.Line2D at 0x15a20fe8700>



```
In [26]: # Create a scale-location plot to test the equal variance assumption  
standardized_residuais = (residuals - residuals.mean()) / residuals.std()  
plt.scatter(linear_regression.predict(uncoor_x_train), np.sqrt(abs(standardized_residuais)))  
plt.xlabel('Fitted values')  
plt.ylabel('Sqrt(abs(standardized residuals))')  
plt.title('Scale-Location')  
  
# Observe that the majority of the standardized residuals are equally spread around .75  
# variance assumption is met.
```

Out[26]: Text(0.5, 1.0, 'Scale-Location')



```
In [27]: # Create a data frame with two columns (true values from the testing dataset and predicted values)  
compare = pd.DataFrame({'True Value': uncoor_y_test, 'Predicted Value': y_predict})  
  
# Reset the index and drop the 'index' column because the true y test value has a row i  
compare.reset_index(inplace = True)
```

```
compare = compare.drop(['index'], axis = 1)

# Round the predicted values to the nearest thousandth
compare['Predicted Value'] = round(compare['Predicted Value'], 3)

# Display the first 5 rows of the comparison data frame
compare.head()

# Observe that the predicted values are similar to the true values. MSE calculated Lat
```

Out[27]:

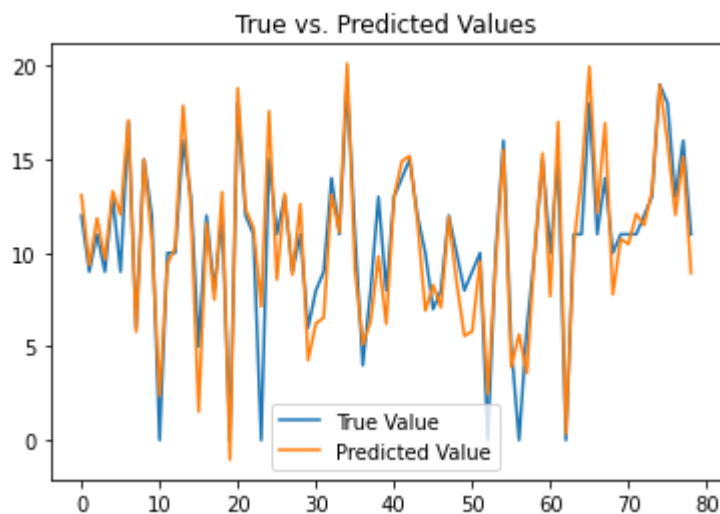
	True Value	Predicted Value
0	12	13.085
1	9	9.351
2	11	11.851
3	9	9.650
4	13	13.298

In [28]:

```
# Plot the true vs predicted values to visualize differences
plt.figure(figsize=(8,6))
compare.plot()
plt.title('True vs. Predicted Values')
```

Out[28]: Text(0.5, 1.0, 'True vs. Predicted Values')

<Figure size 576x432 with 0 Axes>



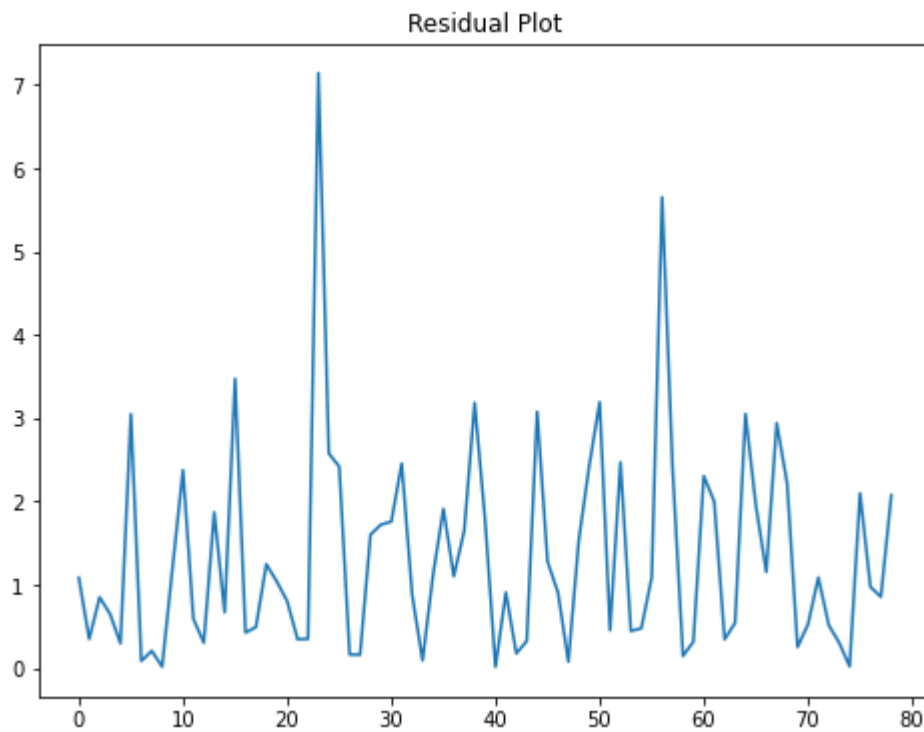
Plot residuals in testing dataset

In [29]:

```
# Plot the residuals (difference between predicted values and true values) to analyze m
plt.figure(figsize=(8,6))
plt.plot(abs(compare['Predicted Value'] - compare['True Value']))
plt.title('Residual Plot')

# Observe that the majority of the error is centered around 2-3 points from the true va
```

Out[29]: Text(0.5, 1.0, 'Residual Plot')



In [30]:

```
# Assess model performance using the mean squared error (MSE)

#Calculate MSE for the linear regression model
MSE_Linear_Reg = mean_squared_error(uncoor_y_test, y_predict)
print('MSE for the linear regression model is', round(MSE_Linear_Reg,3))

# Observing a low MSE of 3.39, it can be accepted that the linear regression model fits
# (given the data contained a low number of total observations (rows))
```

MSE for the linear regression model is 3.39

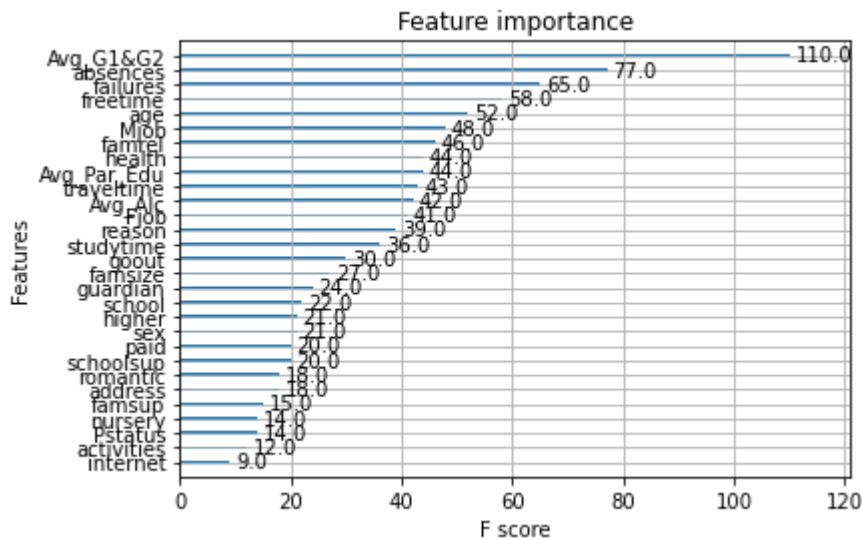
Using XGBoost for Important Feature Identification

In [31]:

```
# Fit an XGBoost regressor with 100 estimators to the training data
xg_reg = xgb.XGBRegressor(objective='reg:squarederror', colsample_bytree = 0.3, learning_rate=0.1)
xg_reg.fit(uncoor_x_train, uncoor_y_train)

# Plot the significant features that contribute the most to the prediction of the dependent variable
xgb.plot_importance(xg_reg)
plt.rcParams['figure.figsize'] = [40, 40]
plt.show()

# A higher F score indicates a higher importance.
# Observe that the most significant features from this model are Avg_G1&G2, absences, and f
# The least significant factors are internet, activites, and Pstatus.
```



Random Forest Regression for Prediction and Feature Importance

In [32]:

```
# Fit a random forest regression model to the training data
Random_F_Reg = RandomForestRegressor(n_estimators=100)
Random_F_Reg.fit(uncoor_x_train, uncoor_y_train)

# Set the features equal to the column names of the training data
features = uncoor_x_train.columns

# Identify the important features using the feature_importances_ built in function
importances = Random_F_Reg.feature_importances_

# Sort the important features from most important to least important
indices = np.argsort(importances)
sorted_indices = np.argsort(importances)[::-1]

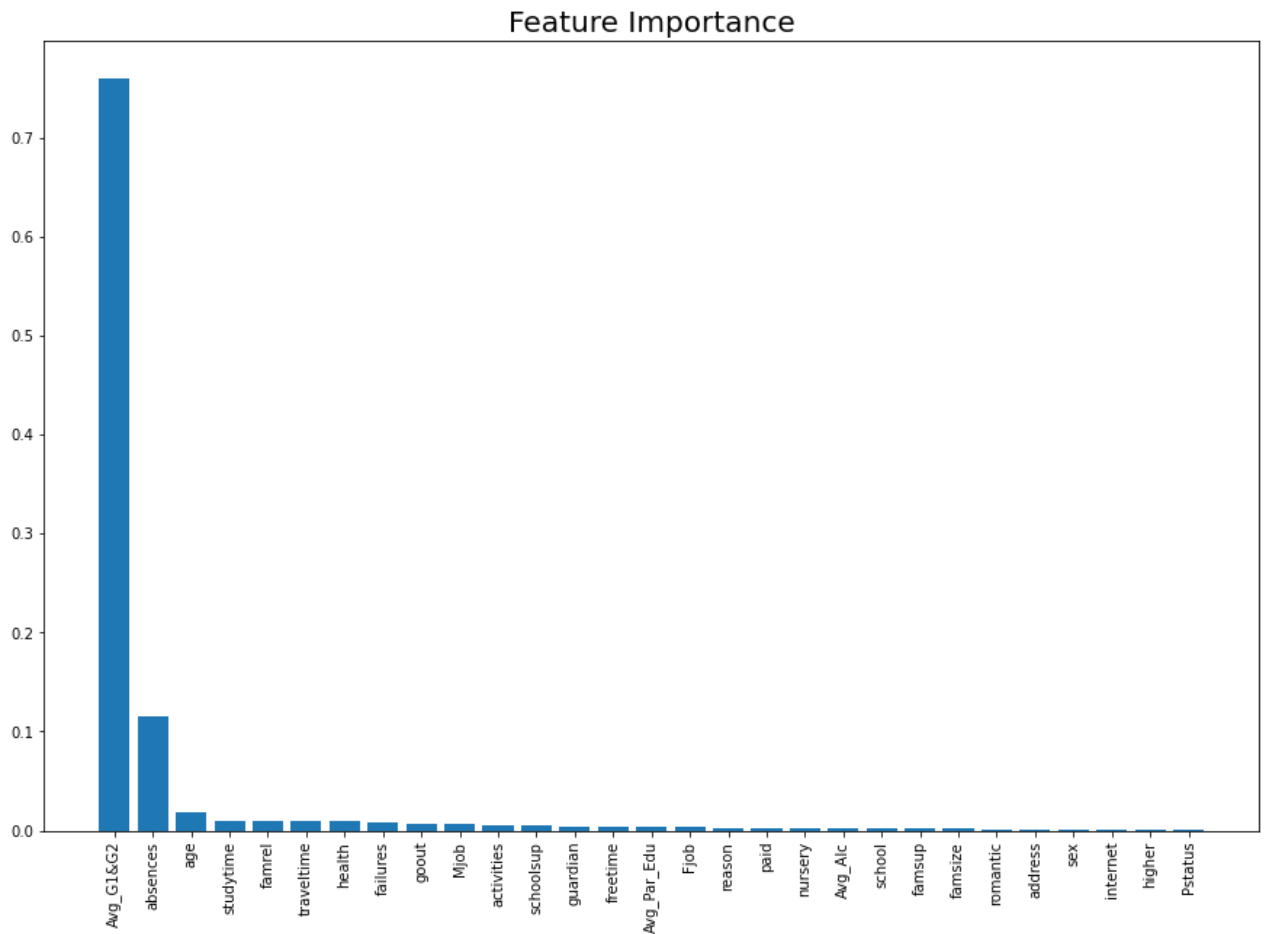
# Print the important features in order from greatest to least
for f in range(uncoor_x_train.shape[1]):
    print("%2d) %-*s %f" % (f + 1, 20, features[sorted_indices[f]], importances[sorted_

# Plot the important features using a vertical bar chart
plt.figure(figsize=(15,10))
plt.title('Feature Importance', size = 20)
plt.bar(range(uncoor_x_train.shape[1]), importances[sorted_indices], align = 'center')
plt.xticks(range(uncoor_x_train.shape[1]), uncoor_x_train.columns[sorted_indices], rota
plt.show()

# Observing the output, the most important features in the prediction of y are Avg_G1&G
# The least important are Pstatus, romantic, internet, and higher.
```

1) Avg_G1&G2	0.760005
2) absences	0.115165
3) age	0.017810
4) studytime	0.010224
5) famrel	0.010173
6) traveltime	0.010044
7) health	0.009456
8) failures	0.007628

9)	goout	0.006282
10)	Mjob	0.006179
11)	activities	0.005531
12)	schoolsup	0.005242
13)	guardian	0.004424
14)	freetime	0.004357
15)	Avg_Par_Edu	0.004079
16)	Fjob	0.003813
17)	reason	0.002723
18)	paid	0.002176
19)	nursery	0.002131
20)	Avg_Alc	0.002131
21)	school	0.002061
22)	famsup	0.001653
23)	famsize	0.001556
24)	romantic	0.001267
25)	address	0.001174
26)	sex	0.000933
27)	internet	0.000789
28)	higher	0.000607
29)	Pstatus	0.000389



In [33]:

```
# Predict course grades using the random forest regressor on the test data
y_pred_RFR = Random_F_Reg.predict(uncoor_x_test)

# Calculate and display the mean absolute error and mean squared error for the model
MAE = mean_absolute_error(uncoor_y_test, y_pred_RFR)
MSE = mean_squared_error(uncoor_y_test, y_pred_RFR)

print('MAE: ', round(MAE,3))
print('MSE: ', round(MSE, 3))
```

```
# Create a dataframe that contains the true vs predicted values
compare_RFR = pd.DataFrame({'Real Value': uncoor_y_test, 'Predict Value': y_pred_RFR})

# Reset the index and drop the 'index' column because the true y test value has a row i
compare_RFR.reset_index(inplace = True)
compare_RFR = compare_RFR.drop(['index'], axis = 1)
compare_RFR.head()
```

MAE: 1.108

MSE: 2.278

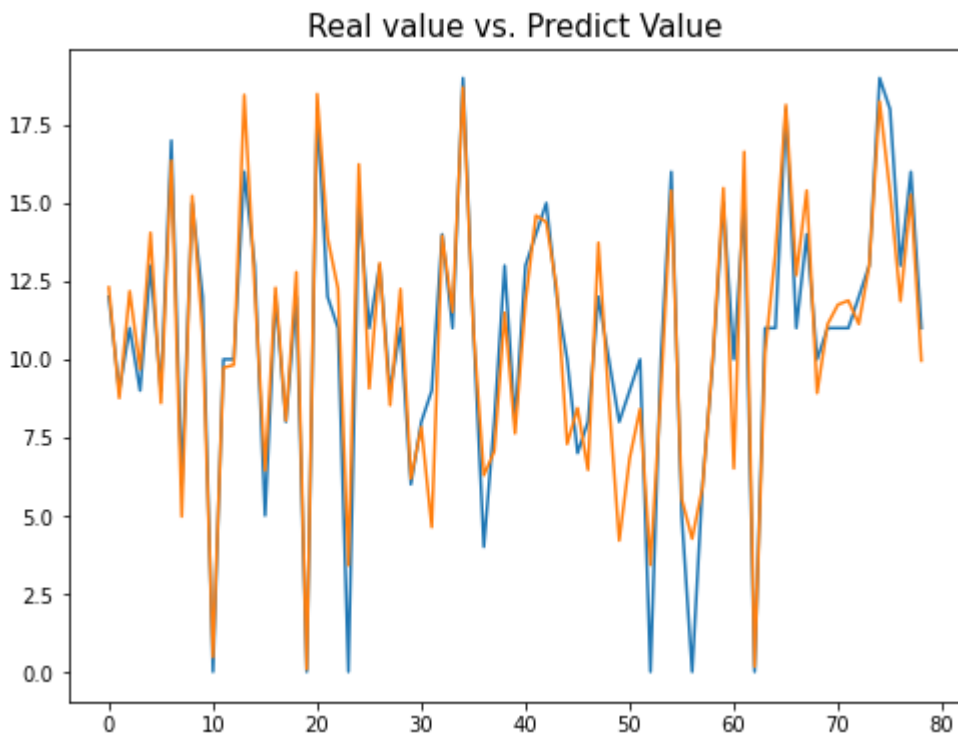
Out[33]:

	Real Value	Predict Value
0	12	12.30
1	9	8.76
2	11	12.18
3	9	9.68
4	13	14.05

In [34]:

```
# Plot the true vs predicted values to visualize differences
plt.figure(figsize=(8,6))
plt.plot(compare_RFR)
plt.title('Real value vs. Predict Value', size = 15)
```

Out[34]: Text(0.5, 1.0, 'Real value vs. Predict Value')



Predicting using ML Classification Methods

Convert continuous data to discrete by mapping grade value integers to letter grade objects. Set a threshold for assigning letter grades. Class 1: 0-5 -> D, Class 2: 6-10 -> C, Class 3: 11-15 -> B, Class

4: 16-20 -> A

In [35]:

```
# Create a data frame containing only the G3 (final math grade) data
G3_df = Uncoorelated_df['G3']

# Initialize an empty list for the new Letter grade strings to be added to
Letter_Grades_List = []

# Loop through all the rows in the subsetted G3 dataframe
for i in range(0, len(G3_df)):

    # For grades between 0-5 assign the Letter D
    if (G3_df[i] <= 5):
        Letter_Grade = 'D'

    # For grades between 6-10 assign the Letter C
    elif (G3_df[i] >= 6 and G3_df[i] <= 10):
        Letter_Grade = 'C'

    # For grades between 11-15 assign the Letter B
    elif (G3_df[i] >= 11 and G3_df[i] <= 15):
        Letter_Grade = 'B'

    # For grades between 16-20 assign the Letter A
    elif (G3_df[i] >= 16 and G3_df[i] <= 20):
        Letter_Grade = 'A'

    # Append the Letter grade to the List on each iteration
    Letter_Grades_List.append(Letter_Grade)

# Add the Letter grade List to the Letter grades data frame
Letter_Grade_Table = pd.DataFrame(Letter_Grades_List)

# Rename the column of Letter grades to the appropriate name
Letter_Grade_Table.rename(columns = {0: 'LetterGrade'}, inplace = True)

# Combine the original "Uncoorelated_df" data frame used for the regression analysis
# with the new Letter grade dataframe.
Classification_df = pd.concat((Letter_Grade_Table, Uncoorelated_df), axis=1)

# Display the G3 grade and Letter Grade to verify the mapping was done correctly
print(Classification_df['G3'])
print(Classification_df['LetterGrade'])
```

```
0      6
1      6
2     10
3     15
4     10
..
390    9
391    16
392    7
393    10
394    9
Name: G3, Length: 395, dtype: int64
```

```

0      C
1      C
2      C
3      B
4      C
..
390    C
391    A
392    C
393    C
394    C
Name: LetterGrade, Length: 395, dtype: object

```

```

In [36]: # Create a new data frame called "Classification_df_Updated" to update the "Classification_df"
Classification_df_Updated = Classification_df.drop(['G3'], axis=1)
print(Classification_df_Updated)

```

```

LetterGrade  Avg_G1&G2  Avg_Alc  Avg_Par_Edu  age  traveltime  studytime  \
0           C           5         1           4   18           2           2
1           C           5         1           1   17           1           2
2           C           7         2           1   15           1           2
3           B          14         1           3   15           1           3
4           C           8         1           3   16           1           2
..          ...          ...          ...          ...   ...          ...          ...
390          C           9         4           2   20           1           2
391          A          15         3           2   17           2           1
392          C           9         3           1   21           1           1
393          C          11         3           2   18           3           1
394          C           8         3           1   19           1           1

```

```

failures  famrel  freetime  ...  paid  activities  nursery  higher  \
0         0       4         3  ...   0           0         1         1
1         0       5         3  ...   0           0         0         1
2         3       4         3  ...   1           0         1         1
3         0       3         2  ...   1           1         1         1
4         0       4         3  ...   1           0         1         1
..          ...          ...          ...   ...          ...          ...          ...
390        2       5         5  ...   1           0         1         1
391        0       2         4  ...   0           0         0         1
392        3       5         5  ...   0           0         0         1
393        0       4         4  ...   0           0         0         1
394        0       3         2  ...   0           0         1         1

```

```

internet  romantic  Mjob  Fjob  reason  guardian
0         0         0     0     0         0         0
1         1         0     0     1         0         1
2         1         0     0     1         1         0
3         1         1     1     2         2         0
4         0         0     2     1         2         1
..          ...          ...          ...   ...          ...
390        0         0     3     2         0         2
391        1         0     3     2         0         0
392        0         0     2     1         0         2
393        1         0     3     1         0         0
394        1         0     2     4         0         1

```

[395 rows x 30 columns]

```

In [37]: # Perform a random split on the new dataset containing Letter grades into 20% testing and 80% training
LG_train, LG_test = train_test_split(Classification_df_Updated, test_size = 0.2, random_state = 42)

# Display the dimensions of the new training and new testing data
print('Size of the train dataset is', LG_train.shape)

```



```

print('Size of the test dataset is', LG_test.shape)

# Split training data into x (independent variables) and y (dependent variable)

# Assign all of the columns except LetterGrade to be the new x
LG_x_train = LG_train.loc[:, LG_train.columns != 'LetterGrade']

# Assign LetterGrade to be y
LG_y_train = LG_train['LetterGrade']

LG_x_test = LG_test.loc[:, LG_test.columns != 'LetterGrade']
LG_y_test = LG_test['LetterGrade']

```

Size of the train dataset is (316, 30)
Size of the test dataset is (79, 30)

```

In [38]: # Fit the SVM algorithm to the training data
SVM_classifier = SVC(kernel='poly')
SVM_classifier.fit(LG_x_train, LG_y_train)

# Test the model by predicting the letter grade of the test data
y_pred_SVM = SVM_classifier.predict(LG_x_test)

# Display the SVM model accuracy
print('SVM Model Accuracy:' , round(accuracy_score(LG_y_test, y_pred_SVM),4))

```

SVM Model Accuracy: 0.8101

```

In [39]: # Fit the random forest algorithm to the training data
RF_classifier = RandomForestClassifier(n_estimators = 200)
RF_classifier.fit(LG_x_train, LG_y_train)

# Test the model by predicting the letter grade of the test data
y_pred_RF = RF_classifier.predict(LG_x_test)

# Display the Random Forest model accuracy
print('Random Forest Model Accuracy:' , round(accuracy_score(LG_y_test, y_pred_RF),4))

```

Random Forest Model Accuracy: 0.8101

PCA for Dimension Reduction

```

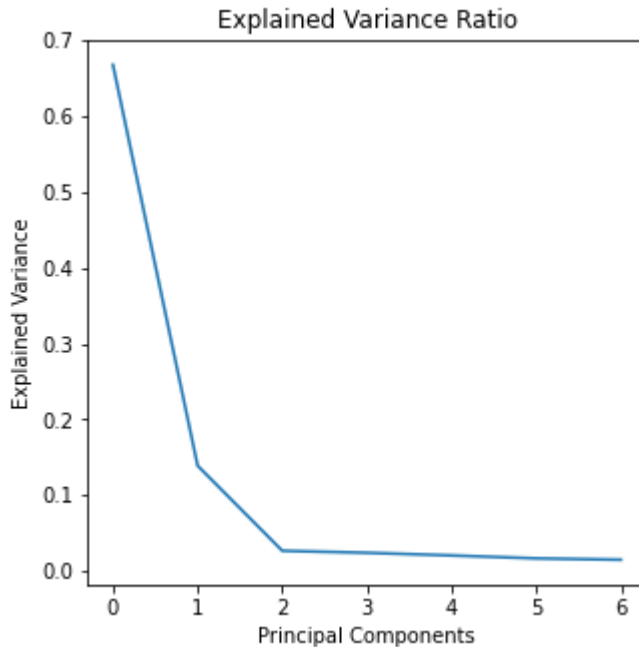
In [40]: # Set the number of PCA components (adjust as necessary)
num_components = 7

# Fit PCA to the training data
pca = PCA(n_components = num_components)
train_features = pca.fit_transform(LG_x_train)

# Plot the explained variance ratio
plt.subplots(figsize=(5, 5))
plt.plot(range(0,num_components), pca.explained_variance_ratio_)
plt.ylabel('Explained Variance')
plt.xlabel('Principal Components')
plt.title('Explained Variance Ratio')
plt.show()

```

```
# Observe that about 7 components explains 100% of the model variance
```



```
In [41]: # Create new x training and testing data based on the reduced number of PCA components
X_train_PCA = pca.fit_transform(LG_x_train)
X_test_PCA = pca.transform(LG_x_test)

# Apply random forest model to newly reduced data
New_RF_classifier = RandomForestClassifier(n_estimators = 200)
New_RF_classifier.fit(X_train_PCA, LG_y_train)

# Test the model by predicting the letter grade of the test data with PCA reduction applied
y_pred_PCA = New_RF_classifier.predict(X_test_PCA)

# Display the new RF accuracy score
print('Random Forest Model Accuracy with PCA reduction: ', round(accuracy_score(LG_y_test, y_pred_PCA), 4))

# Observe that the model accuracy improves by ~ 10% when using PCA reduction.
```

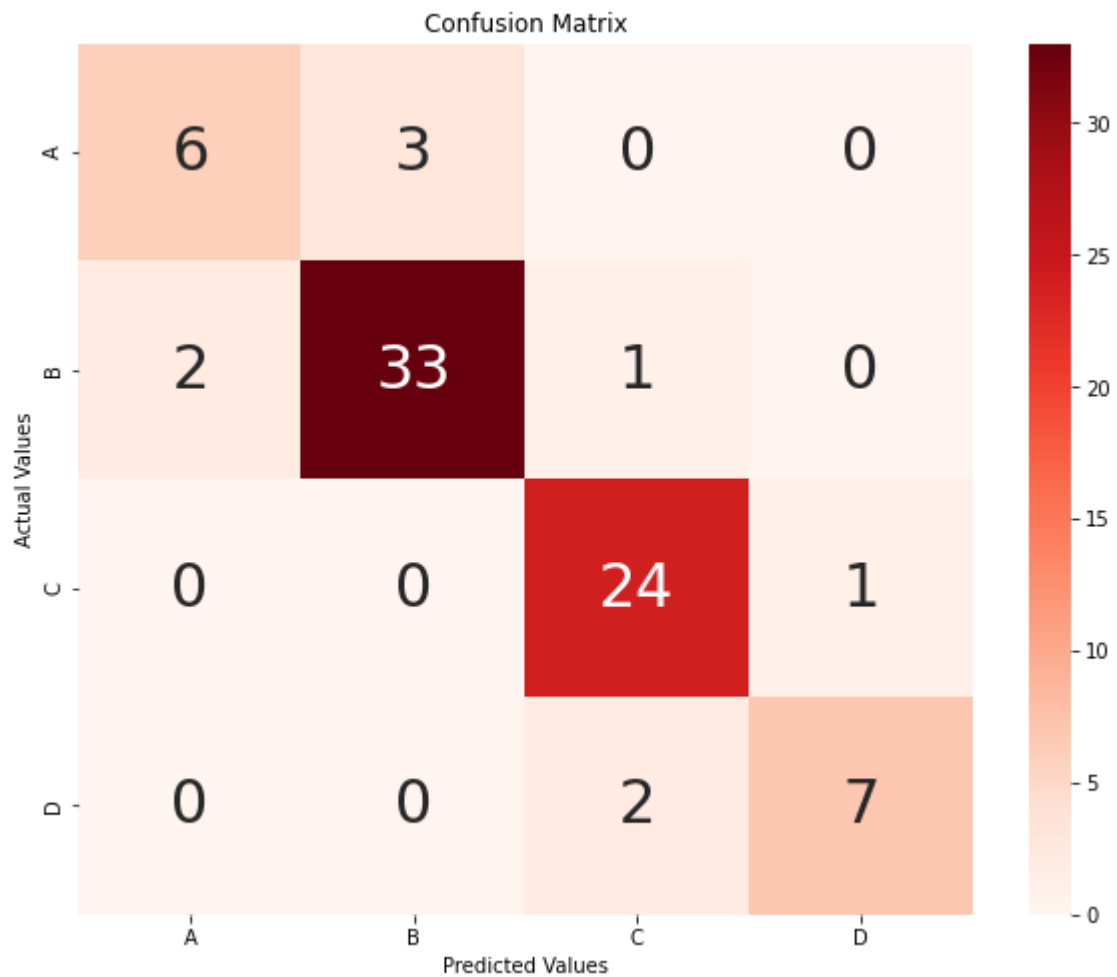
Random Forest Model Accuracy with PCA reduction: 0.8861

```
In [42]: # Create a confusion matrix comparing the true to predicted values
conf_mx = confusion_matrix(LG_y_test, y_pred_PCA)

# Create the labels for the x and y ticks of the confusion matrix
Tick_labels = ['A', 'B', 'C', 'D']

# Use the seaborn heatmap function to create a visual of the confusion matrix
plt.figure(figsize = (10,8))
Conf_Mx_Plot = sns.heatmap(conf_mx, annot = True, annot_kws = {"size": 30}, cmap = 'Reds')
Conf_Mx_Plot.set(title='Confusion Matrix')
Conf_Mx_Plot.set(xlabel='Predicted Values', ylabel='Actual Values')
plt.show()

# Observe that the classes are imbalanced. The data contains more instances of B and C
# To truly evaluate the model accuracy, use the classification report to analyze precision, recall, and f1 score.
```



In [43]:

```
print("RF Classification Report:\n")
print(classification_report(LG_y_test, y_pred_PCA))

# Observe that the precision and recall is the highest for the B class, giving it the h
# The lowest F-1 score occurs for class A.
```

RF Classification Report:

	precision	recall	f1-score	support
A	0.75	0.67	0.71	9
B	0.92	0.92	0.92	36
C	0.89	0.96	0.92	25
D	0.88	0.78	0.82	9
accuracy			0.89	79
macro avg	0.86	0.83	0.84	79
weighted avg	0.88	0.89	0.88	79