# Part 2 Math grades for both schools

```
In [2]:
         import pandas as pd
                                                #Importing the required package
         import numpy as np
         import matplotlib.pyplot as plt
         import seaborn as sns
         from sklearn.preprocessing import MinMaxScaler
         from sklearn.model_selection import KFold, cross_val_predict
         from sklearn.pipeline import Pipeline
         from sklearn.metrics import mean_squared_error
         from sklearn.metrics import r2_score
         from sklearn.model_selection import GridSearchCV
         from timeit import default_timer as timer
         from sklearn.linear_model import LinearRegression
         import warnings
         warnings.simplefilter(action='ignore', category=FutureWarning)
         warnings.simplefilter(action='ignore', category=DeprecationWarning)
         warnings.simplefilter(action='ignore', category=UserWarning)
In [ ]:
         data = pd.read_csv('student-mat.csv',sep=';')
         data.head()
In [ ]:
         data = data.drop(['G1','G2'],axis=1) #dropping columns G1 and G2 in order to get a m
         data.head()
In [ ]:
          print("Number of rows:", len(data)) #Looking for number of rows in total
In [ ]:
         data.isna().sum() #Checking for Null Values
In [ ]:
                                   #checking for Data types of each variables
         data.info()
In [ ]:
         for var in ['traveltime','studytime','Medu','Fedu','famrel','freetime','goout','Dalc
              data[var] = data[var].astype('category')
         data.info()
         cat_col = list(set(data.columns)-set(num_col))
num_col = list(num_col)[0.2]
#Getting the numerical dataset fr
#By removing the numerical dataset
In [ ]:
In [ ]:
         nominal_col = ['higher','Fjob','address','guardian','school','paid','sex','Mjob','ad
         ordinal_col = np.setdiff1d(cat_col,nominal_col)
In [ ]:
         for col in ordinal col:
              data[col] = data[col].astype(int)
                                                                  # Changing Ordinal data to int
In [ ]:
         for col in data[cat_col]:
              print(col,data[col].unique())
                                                                # Checking for erroneous data in
```

### **Exploratory Data Analytics**

#### Based on observation, it is shown the following:

Fedu and Medu has moderate correlation

Walc and Dalc has moderate correlation

Walc and goout has weak correlation

```
In [ ]:
    sns.set(rc={'figure.figsize':(15,5)})
    sns.swarmplot(data=data, x='school', y='age')
```

#### Based on observation, it is shown that:

Students in GP are relatively younger than MS

```
ax = sns.boxplot(data=data, x='school', y='absences')
ax.set_ylim([0, 35])
```

#### Based on observation, it is shown that:

GP has more absentees comapred to MS

#### Based on Observation on the boxplot, it is shown that:

In terms of median, 75th and 25th percentile, students in GP are relatively doing better than students in MS

# Reducing Skewness in numerical data

```
In [ ]:
         skew_limit = 0.75
         skew_cols = (data[num_col].skew()
                      .sort_values(ascending=False)
                       .to frame()
                       .rename(columns={0:'Skew'})
                       .query('abs(Skew) > {}'.format(skew_limit)))
         skew_cols
In [ ]:
        field = "absences"
         fig, (ax_original, ax_log1p) = plt.subplots(1, 2, figsize=(15, 5))
         data[field].hist(ax=ax_original)
         # Apply a log transformation (numpy syntax) to this column
         data[field].apply(np.log1p).hist(ax=ax_log1p)
         # Formatting of titles etc. for each subplot
         ax_original.set(title='before np.log1p', ylabel='frequency', xlabel='value')
         ax_log1p.set(title='after np.log1p', ylabel='frequency', xlabel='value')
         fig.suptitle('Field "{}"'.format(field));
         print('pop_orignial skewness: ',data[field].skew())
         print('pop_log1p skewness: ',data[field].apply(np.log1p).skew())
         # fall in skewness after log1p
In [ ]:
         # apply log1p across all numerical columns
         for col in data[skew_cols.index]:
         # drop famrel as skewness increase after log1p
             data[col] = data[col].apply(np.log1p)
         new_skew_cols = (data[skew_cols.index].skew()
                      .sort_values(ascending=False)
                      .to_frame()
                      .rename(columns={0:'Skew'}))
         skew_cols['New_skew'] = new_skew_cols
         skew_cols['Difference'] = abs(skew_cols['New_skew']) - abs(skew_cols['Skew'])
         skew cols
```

## MiniMax scale numerical data

# One hot encoding of categorical Data

```
data = pd.get_dummies(data, columns=nominal_col,drop_first=True)
    data.head()
    # df is the transformed dataest for ML, df is the orignal dataset
```

## **Polynomial Features**

## **Train Test Split**

```
In [ ]:
         from sklearn.model_selection import train_test_split
         train,test = train_test_split(data,
                                               train_size = 0.7,
                                               test_size = 0.3,
                                               random_state = 42)
         feature_cols = list(filter(lambda x: x!= 'G3', data.columns))
         X_train = train[feature_cols]
         y_train = train['G3']
         X_test = test[feature_cols]
         y_test = test['G3']
In [ ]:
         # splitting of Polynomial feautures
         X_pf_train = X_pf[X_train.index]
         y_pf_train = data['G3'][X_train.index]
         X_pf_{test} = X_pf[X_{test.index}]
         y_pf_test = data['G3'][X_test.index]
```

#### Model Kfold + Evaluation

```
In [ ]:
         # define root mean sq error func
         def rmse(ytrue, ypredicted):
             return np.sqrt(mean_squared_error(ytrue, ypredicted))
         folds = KFold(n splits = 5, shuffle = True, random state = 100)
In [ ]:
         import statsmodels.api as sm
         def diagnostic plot(y test, prediction):
             residual = y_test - prediction
             fig, axs = plt.subplots(2, 2,figsize=(15,10))
             fig.suptitle('Diagnostic plots')
             axs[0,0].plot([0, 15], [0, 20], ls="--", c="red", alpha=0.5)
             axs[0,0].scatter(y test, prediction)
             axs[0,0].set_title('Truth-Prediction plot')
             axs[0,0].set_xlabel("Truth")
             axs[0,0].set ylabel("Predictions")
```

```
sm.qqplot(residual,fit=True,line= 's', ax=axs[0,1])
axs[0,1].set_title('QQ plot')
axs[0,1].set_xlabel("Theoretical Quantiles")
axs[0,1].set ylabel("Sample Quantiles")
sns.residplot(prediction,y_test,
                          lowess=True,
                          scatter_kws={'alpha': 0.5},
                          line_kws={'color': 'red', 'lw': 1, 'alpha': 0.8},
                          ax=axs[1,0]
axs[1,0].set_title('Residual plot')
axs[1,0].set_xlabel("Predicted")
axs[1,0].set_ylabel("Residual")
sqrt_standardized_residual=np.sqrt(np.abs(residual))
sns.regplot(prediction, sqrt_standardized_residual,
              scatter=True,
              lowess=True,
              line_kws={'color': 'red', 'lw': 1, 'alpha': 0.5},
              ax=axs[1,1])
axs[1,1].set_title('Scale-Location plot')
axs[1,1].set_xlabel("Predicted")
axs[1,1].set_ylabel("Sqrt Standarized residuals")
```

## **Dummy Model**

# **Base Linear Regression**

```
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                                                      part2 mat codes
               linear_prediction = linear_cv.predict(X_test)
     In [ ]:
               print("R2 score: ",r2_score(y_test, linear_prediction))
               print('RMSE: ', rmse(y_test,linear_prediction))
     In [ ]:
               diagnostic_plot(y_test, linear_prediction)
     In [ ]:
               print("\u0332".join('Feature_coefficient (Top)'),'\n')
               features = []
               coeffs = []
               coeffs_abs = []
               for feature, coeff in zip(X_test.columns, linear_cv.best_estimator_.coef_):
                   features.append(feature)
                   coeffs.append(coeff)
                   coeffs abs.append(abs(coeff))
               pd.DataFrame({'Features': features, 'Coefficients': coeffs ,'abs_coeffs':coeffs_abs}
     In [ ]:
               print("\u0332".join('Feature_coefficient (Bottom)'),'\n')
               pd.DataFrame({'Features': features, 'Coefficients': coeffs ,'abs_coeffs':coeffs_abs}
     In [ ]:
               lm = LinearRegression()
               lm_pf_cv = GridSearchCV(estimator = lm,
                                       param_grid = {},
                                       cv = folds)
               start = timer()
               lm_pf_cv.fit(X_pf_train, y_pf_train)
               end = timer()
               lm_pf_time = end - start
               print('Time Elapsed (sec):', lm_pf_time)
     In [ ]:
               lm_pf_prediction = lm_pf_cv.predict(X_pf_test)
               print("R2 score: ",r2_score(y_pf_test, lm_pf_prediction))
               print('RMSE: ', rmse(y pf test,lm pf prediction))
     In [ ]:
               diagnostic_plot(y_test, lm_pf_prediction)
```

## **Lasso Regression**

```
print('RMSE: ', rmse(y_test,lasso_prediction))
         end = timer()
         lasso_time = end - start
         print('Time Elapsed (sec):', lasso time)
In [ ]:
         print("Number of non-zero coeff: ", sum(lasso_estimator.coef_ != 0))
         print("Mean coeff: ",sum(abs(lasso_estimator.coef_))/sum(lasso_estimator.coef_ != 0)
In [ ]:
         print("\u0332".join('Feature_coefficient (Top)'),'\n')
         features = []
         coeffs = []
         coeffs abs = []
         for feature, coeff in zip(X test.columns, lasso estimator.coef ):
             features.append(feature)
             coeffs.append(coeff)
             coeffs_abs.append(abs(coeff))
         pd.DataFrame({'Features': features, 'Coefficients': coeffs ,'abs_coeffs':coeffs_abs}
In [ ]:
         diagnostic_plot(y_test, lasso_prediction)
```

### **Ridge Regression**

```
In [ ]:
         alphas = np.geomspace(1, 100, 30)
In [ ]:
         from sklearn.linear model import RidgeCV
         start = timer()
         ridge_estimator = RidgeCV(alphas=alphas,
                                      cv=5).fit(X_train,y_train)
         ridge_prediction = ridge_estimator.predict(X_test)
         print('Alpha param: ', ridge_estimator.alpha_)
         print("R2 score: ",r2 score(y test, ridge prediction))
         print('RMSE: ', rmse(y test, ridge prediction))
         end = timer()
         ridge time = end - start
         print('Time Elapsed (sec):', ridge_time)
In [ ]:
         print("Number of non-zero coeff: ", sum(ridge_estimator.coef_ != 0))
         print("Mean coeff: ",sum(abs(ridge estimator.coef ))/sum(ridge estimator.coef != 0)
In [ ]:
         print("\u0332".join('Feature coefficient (Top)'),'\n')
         features = []
         coeffs = []
         coeffs abs = []
         for feature, coeff in zip(X_test.columns, ridge_estimator.coef_):
             features.append(feature)
             coeffs.append(coeff)
             coeffs abs.append(abs(coeff))
         pd.DataFrame({'Features': features, 'Coefficients': coeffs ,'abs coeffs':coeffs abs}
```

```
In [ ]: diagnostic_plot(y_test, ridge_prediction)
```

### **Elasticnet Regression**

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```
In [ ]:
         from sklearn.linear model import ElasticNetCV
         alphas = np.geomspace(0.01, 1, 30)
         11 ratios = np.linspace(0.1, 0.9, 9)
         start = timer()
         elasticNetCV = ElasticNetCV(alphas=alphas,
                                      l1_ratio=l1_ratios,
                                      cv=5).fit(X_train,y_train)
         elasticNet_prediction = elasticNetCV.predict(X_test)
         print('Alpha param: ', elasticNetCV.alpha_)
         print('l1 ratio param: ', elasticNetCV.l1_ratio_)
         print("R2 score: ",r2_score(y_test, elasticNet_prediction))
         print('RMSE: ', rmse(y_test,elasticNet_prediction))
         end = timer()
         EN\_time = end - start
         print('Time Elapsed (sec):', EN_time)
In [ ]:
         print("Number of non-zero coeff: ", sum(elasticNetCV.coef_ != 0))
         print("Mean coeff: ",sum(abs(elasticNetCV.coef_))/sum(elasticNetCV.coef_ != 0))
In [ ]:
         print("\u0332".join('Feature_coefficient (Top)'),'\n')
         features = []
         coeffs = []
         coeffs_abs = []
         for feature, coeff in zip(X_test.columns, elasticNetCV.coef_):
             features.append(feature)
             coeffs.append(coeff)
             coeffs_abs.append(abs(coeff))
         pd.DataFrame({'Features': features, 'Coefficients': coeffs ,'abs_coeffs':coeffs_abs}
In [ ]:
         diagnostic_plot(y_test, elasticNet_prediction)
```

### **Stepwise regression - Forward**

pval[column] = model.pvalues[column]

```
best = pval.min()
                 if best < threshold in:</pre>
                     best_feature = pval.idxmin()
                      included.append(best feature)
                     changed=True
                     if verbose:
                         print('Add {:17} with p-value {:.6}'.format(best_feature, best))
                 if not changed:
                     break
             return included
         # Starting with all features, features are removed iteratively based on the p-value
         def backward_regression(X, y,
                                     threshold out,
                                     verbose=False):
             included=list(X.columns)
             while True:
                 changed=False
                 model = sm.OLS(y, sm.add_constant(pd.DataFrame(X[included]))).fit()
                 # use all coefs except intercept
                 pvalues = model.pvalues.iloc[1:]
                 worst_pval = pvalues.max() # null if pvalues is empty
                 if worst_pval > threshold_out:
                      changed=True
                     worst_feature = pvalues.idxmax()
                     included.remove(worst_feature)
                     if verbose:
                         print('Drop {:17} with p-value {:.6}'.format(worst feature, worst pv
                 if not changed:
                     break
             return included
In [ ]:
         start = timer()
         forward_regression_features = forward_regression(X_train,y_train, threshold_in=0.05,
         forward_regression_features
In [ ]:
         lm = LinearRegression()
         forward = GridSearchCV(estimator = lm,
                                  param_grid = {},
                                  cv = folds)
         # fit the model
         forward.fit(X_train[forward_regression_features], y_train)
         end = timer()
         forward time = end - start
         print('Time Elapsed (sec):', forward time)
In [ ]:
         forward prediction = forward.predict(X test[forward regression features])
         print("R2 score: ",r2_score(y_test,forward_prediction))
         print('RMSE: ', rmse(y_test,forward_prediction))
In [ ]:
         diagnostic plot(y test, forward prediction)
```

### Stepwise regression - Backward

```
In [ ]:
         start = timer()
         backward_regression_features = backward_regression(X_train,y_train, threshold_out=0.
In [ ]:
        lm = LinearRegression()
         backward = GridSearchCV(estimator = lm,
                                 param_grid = {},
                                 cv = folds)
         # fit the model
         backward.fit(X_train[backward_regression_features], y_train)
         end = timer()
         backward time = end - start
         print('Time Elapsed (sec):', backward time)
In [ ]:
         backwards_prediction = backward.predict(X_test[backward_regression_features])
         print("R2 score: ",r2_score(y_test, backwards_prediction))
         print('RMSE: ', rmse(y_test,backwards_prediction))
In [ ]:
         diagnostic plot(y test, backwards prediction)
       Random Forest Regressor
In [ ]:
         from sklearn.ensemble import RandomForestRegressor
         random_tree = RandomForestRegressor(criterion = 'mse')
```

```
In [ ]:
         n_features = len(data.columns)
         max_features = ['auto', 'sqrt', 'log2']
         max_depth = [3, 4]
         min_samples_split = [2, 5, 10]
         min_samples_leaf = [1, 2, 4]
         param_test = {'max_features': max_features,
          'max_depth': max_depth,
          'min_samples_split': min_samples_split,
          'min_samples_leaf': min_samples_leaf}
         print(param_test)
In [ ]:
         random_tree = GridSearchCV(estimator = random_tree,
                                       param_grid = param_test,
                                       cv = folds)
         start = timer()
         random_tree.fit(X_train, y_train)
         end = timer()
         rf time = end - start
         print('Time Elapsed (sec):', rf_time)
In [ ]:
         random tree prediction = random tree.predict(X test)
         print('Best param: ', random_tree.best_params_)
```

```
print("R2 score: ",r2_score(y_test, random_tree_prediction))
         print('RMSE: ', rmse(y_test,random_tree_prediction))
In [ ]:
         diagnostic_plot(y_test, random_tree_prediction)
In [ ]:
         from sklearn.tree import export graphviz
         tree = random_tree.best_estimator_.estimators_[5]
         export_graphviz(tree, out_file='small.dot',
                         feature names = X train.columns)
In [ ]:
         from IPython.display import Image
         Image(filename = 'small.png')
In [ ]:
         feat_df = pd.DataFrame({'Feature':X_train.columns, 'Importance':random_tree.best_est
         feat_df = feat_df.sort_values(by = 'Importance', ascending=False)
         feat_df.head()
In [ ]:
         g = sns.barplot(data=feat_df, x='Feature', y= 'Importance')
         for item in g.get_xticklabels():
             item.set_rotation(90)
In [ ]:
         models_pred = [dummy_prediction, linear_prediction, lm_pf_prediction, lasso_predicti
         time elapsed = [dummy time, linear time, lm pf time, lasso time, ridge time, EN time
         rmse_vals = []
         for pred in models pred:
             rmse_vals.append(rmse(y_test, pred))
         R2\_score = []
         for pred in models_pred:
             R2_score.append(r2_score(y_test, pred))
         labels = ['Dummy', 'Linear', 'Linear + PF', 'Ridge', 'Lasso', 'ElasticNet','Stepwise
         eval df = pd.DataFrame({'RMSE':rmse vals, 'R2 Score':R2 score, 'Time Elapsed':time e
         eval df
In [ ]:
```