```
def display_side_by_side(*args):
            html str=''
            for df in args:
                html str+=df.to html()
             display_html(html_str.replace('table','table style="display:inline"'),raw=True)
         class display(object):
             """Display HTML representation of multiple objects"""
             template = """<div style="float: left; padding: 10px;">
             {0}{1}
             </div>"""
            def __init__(self, *args):
                self.args = args
            def repr html (self):
                return '\n'.join(self.template.format(a, eval(a)._repr_html_())
                                 for a in self.args)
            def __repr__(self):
                return '\n\n'.join(a + '\n' + repr(eval(a))
                                   for a in self.args)
In [2]:
         class Master:
             def predectors_select(data_frame, Exception=None):
                predictors = []
                data_type = ['int64', 'float64']
                 for column in data_frame.columns:
                     if data_frame[column].dtype in data_type:
                        if Exception == None:
                            predictors.append(column)
                         elif column not in Exception:
                            predictors.append(column)
                 return set(predictors)
             def correlation(data frame: 'dataframe', min:'decimal'=.5, max:'decimal'=1):
                 data frame temp = data frame.corr().abs().reset index()
                 data_frame_temp.drop(columns=['index'], inplace=True)
                unique_v1 = set()
                unique_v2 = set()
                counter = 0
                 for index, data in data_frame_temp.iterrows():
                     el=[_ for _ in range(data_frame_temp.index.size) if _!= counter]
                     for j in el:
                        if data[j] > min and data[j] < max:</pre>
                             unique_v1.add(data.index[index])
                             unique_v2.add(data.index[j])
                             #print("{0:>15} {1:>15} {2:>10.5f}".format(data.index[index], data.index[j], data[j]))
                     counter +=1
                 return sorted(list(unique_v1.intersection(unique_v2))) #unique_v1, unique_v2
             def vif(dataframe):
                variance inflation factor (VIF)
                Parameters
                data_frame = dataframe
                Returns
                Pandas Series formed by VIF for each value in the dataframe
                 #vif dataframe = pd.DataFrame()
                 #vif_dataframe['VIF Factor'] = [variance_inflation_factor(dataframe.values, i) for i in range(dataframe.shape[1]
                 #vif dataframe['Features'] = dataframe.columns
                 vif = dict()
                 for idx in range(dataframe.shape[1]):
                     vif[ dataframe.columns[idx] ] = variance_inflation_factor(dataframe.values, idx)
                 return pd.Series(vif, index=dataframe.columns)
             def log_trans(data_frame, to_log=None):
                 #data frame log = pd.DataFrame()
                 if to_log == None:
                    data_frame = data_frame.applymap(lambda x: np.log(x) if x > 0 else 0)
                     for feat in to log:
                         data_frame[feat] = data_frame[feat].apply(lambda x: np.log(x))
                 #return data frame
             def sklearn_model(x,y):
                 x_train, x_test, y_train, y_test = train_test_split(x, y, test_size=.3, random_state=7)
                linreg = LinearRegression()
                linreg.fit(x_train, y_train)
                y_hat_train = linreg.predict(x_train)
                y_hat_test = linreg.predict(x_test)
                 train_residuals = y_hat_train - y_train
                 test_residuals = y_hat_test - y_test
```

In [1]:

from IPython.display import display_html

```
train_mse = mean_squared_error(y_train, y_hat_train)
       test_mse = mean_squared_error(y_test, y_hat_test)
       print('R^2 Score {0:>22}'.format(round(linreg.score(x, y),2)))
       print('Train Mean Squarred Error {0:>20}'.format(train_mse))
       print('Test Mean Squarred Error {0:>21}'.format(test_mse))
       return linreg
#
    def metrics(y, y hat):
       r2 = r2_score(y, y_hat)
       mae = mean_absolute_error(y, y_hat)
       mse = mean_squared_error(y, y_hat)
       rmse = np.sqrt(mean_squared_error(y, y_hat))
                       = y scaler.inverse transform(y)
       y original
       y_hat_original = y_scaler.inverse_transform(np.array(y_hat).reshape(-1,1))
       mae_inverse = mean_absolute_error( y_original, y_hat_original)
       rmse_inverse = np.sqrt(mean_squared_error( y_original, y_hat_original))
       print(f"{'R-squared':<25}{r2:>5.3f}")
       print(f"{'Mean Absolute Error':<25}{mae:>5.3f} ${mae_inverse:,.2f}")
       print(f"{'Mean Squared Error':<25}{mse:>5.3f}")
       print(f"{'Root Mean Squared Error':<25}{rmse:>5.3f} ${rmse_inverse:,.2f}")
       return mae, mse, rmse
   def stepwise_selection(X, y,
                           initial list=[],
                           threshold in=0.01,
                           threshold out = 0.05,
                           verbose=True):
        """ Perform a forward-backward feature selection
       based on p-value from statsmodels.api.OLS
       Arguments:
           X - pandas.DataFrame with candidate features
           y - list-like with the target
           initial_list - list of features to start with (column names of X)
           threshold_in - include a feature if its p-value < threshold_in</pre>
           threshold_out - exclude a feature if its p-value > threshold_out
           verbose - whether to print the sequence of inclusions and exclusions
       Returns: list of selected features
       Always set threshold in < threshold out to avoid infinite looping.
       See https://en.wikipedia.org/wiki/Stepwise_regression for the details
       included = list(initial list)
       while True:
           changed=False
            # forward step
           excluded = list(set(X.columns)-set(included))
           new pval = pd.Series(index=excluded, dtype='float64')
            for new column in excluded:
                model = sm.OLS(y, sm.add_constant(pd.DataFrame(X[included+[new column]]))).fit()
                new_pval[new_column] = model.pvalues[new_column]
           best_pval = new_pval.min()
           if best_pval < threshold_in:</pre>
                best feature = new pval.idxmin()
                included.append(best_feature)
                changed=True
                if verbose:
                    print('Add {:30} with p-value {:.6}'.format(best feature, best pval))
            # backward step
           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()
                # worst_feature = pvalues.argmax()
                included.remove(worst_feature)
                if verbose:
                    print('Drop {:30} with p-value {:.6}'.format(worst_feature, worst_pval))
            if not changed:
       return included
```