# **Multiple Linear Regression**

#### **libraries**

```
In [1]: import numpy as np
   import pandas as pd
   import matplotlib.pyplot as plt
   import seaborn as sns
   sns.set()

# regression module
from sklearn import linear_model
```

#### Load the data

```
In [2]: data = pd.read_csv('Multiple_linear regression.csv')
         # Let's explore the top 5 rows of the df
        data.head()
           SAT Rand 1,2,3 GPA
Out[2]:
        0 1714
                       1 2.40
        1 1664
                       3 2.52
                       3 2.54
        2 1760
        3 1685
                       3 2.74
                       2 2.83
        4 1693
       # descriptive statistics
In [3]:
```

```
data.describe()
                        SAT Rand 1,2,3
                                              GPA
Out[3]:
                               84.000000 84.000000
          count
                   84.000000
                 1845.273810
                                2.059524
                                          3.330238
                  104.530661
                                0.855192
                                          0.271617
            std
            min
                 1634.000000
                                1.000000
                                           2.400000
```

 mean
 1845.273810
 2.059524
 3.330238

 std
 104.530661
 0.855192
 0.271617

 min
 1634.000000
 1.000000
 2.400000

 25%
 1772.000000
 1.000000
 3.190000

 50%
 1846.000000
 2.000000
 3.380000

 75%
 1934.000000
 3.000000
 3.810000

 max
 2050.000000
 3.000000
 3.810000

## multiple regression

### Declare the dependent and independent variables

```
In [4]: # There are two independent variables: 'SAT' and 'Rand 1,2,3'
x = data[['SAT','Rand 1,2,3']]

# and a single depended variable: 'GPA'
y = data['GPA']
```

### p-values in sklearn

```
In [5]: # 'stat' module from scipy.stats
       import scipy.stats as stat
        # Here's the full source code of the ORIGINAL class: https://github.com/scikit-learn/scikit-learn/blob/7b136e9/
        class LinearRegression(linear_model.LinearRegression):
            LinearRegression class after sklearn's, but calculate t-statistics
            and p-values for model coefficients (betas).
            Additional attributes available after .fit()
            are `t` and `p` which are of the shape (y.shape[1], X.shape[1])
            which is (n_features, n coefs)
            This class sets the intercept to 0 by default, since usually we include it
            in X.
            11 11 11
            # nothing changes in init
            def __init__(self, fit_intercept=True, normalize=False, copy_X=True,
                        n jobs=1):
                self.fit intercept = fit intercept
               self.normalize = normalize
                self.copy X = copy X
                self.n_jobs = n_jobs
            def fit(self, X, y, n jobs=1):
                self = super(LinearRegression, self).fit(X, y, n_jobs)
                # Calculate SSE (sum of squared errors)
                # and SE (standard error)
                sse = np.sum((self.predict(X) - y) ** 2, axis=0) / float(X.shape[0] - X.shape[1])
                se = np.array([np.sqrt(np.diagonal(sse * np.linalg.inv(np.dot(X.T, X))))])
                # compute the t-statistic for each feature
                self.t = self.coef / se
                # find the p-value for each feature
                self.p = np.squeeze(2 * (1 - stat.t.cdf(np.abs(self.t), y.shape[0] - X.shape[1])))
                return self
```

```
In [6]: # create the regression everything is the same
  reg_with_pvalues = LinearRegression()
  reg_with_pvalues.fit(x,y)
```

Out[6]: LinearRegression(copy\_X=True, fit\_intercept=True, n\_jobs=1, normalize=False)

In [7]: # The difference is that we can check what's contained in the local variable 'p' in an instance of the LinearRe reg\_with\_pvalues.p

Out[7]: array([0. , 0.75717067])

Out[7]: ", 0.73717067]

```
In [8]: # Let's create a new data frame with the names of the features
    reg_summary = pd.DataFrame([['SAT'],['Rand 1,2,3']],columns =['Features'])
# Then we create and fill a second column, called 'Coefficients' with the coefficients of the regression
    reg_summary['Coefficients'] = reg_with_pvalues.coef_
# Finally, we add the p-values we just calculated
    reg_summary['p-values'] = reg_with_pvalues.p.round(3)
```

In [9]: # This result is identical to the one from StatsModels
reg\_summary

Out[9]:		Features	Coefficients	p-values
	0	SAT	0.001654	0.000
	1	Rand 1.2.3	-0.008270	0.757