## **Assignment3**

- 1. Perfect Relationship Zero Correlation
- Consider the quadratic relationship:

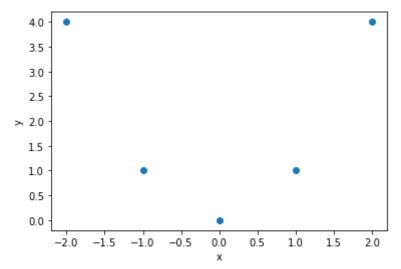
```
i xi yi
```

- 1 24
- 2 11
- 300
- 411
- 524
- a. Produce a graph of this data set;
- b. Calculate the correlation coefficient and chack that it is equal to zero.

```
In [1]: import numpy as np
import matplotlib.pyplot as plt
%matplotlib inline
```

a.

```
In [3]: x = np.array([-2,-1,0,1,2])
y = x**2
plt.scatter(x,y)
plt.xlabel('x')
plt.ylabel('y')
plt.title = 'graph of this data set'
plt.show()
```



b.

In [4]: r = np.corrcoef(x, y)[0][1]
 print('The correlation coefficient is: {}'.format(r))
 print('It is equal to zero')

The correlation coefficient is: 0.0 It is equal to zero

2. Weighted Sums Show that the estimate  $\hat{\beta}_1$  is a weighted sum of the observed response  $y_i$ , ie, that

 $\hat{\beta_1} = \sum_{i=1}^n \omega_i y_i$ 

Hint: Consider the weights

 $\omega_i = \frac{x_i - \bar{x}}{SS_{xx}}$ 

and note that

$$\sum_{i=1}^{n} \omega_i = 0$$

$$\hat{\beta}_{1} = \frac{Cov(X,Y)}{Var(X)} = \frac{\sum_{i=1}^{n} (x_{i} - \bar{x})(y_{i} - \bar{y})}{\sum_{i=1}^{n} (x_{i} - \bar{x})^{2}} = \frac{\sum_{i=1}^{n} (x_{i} - \bar{x})y_{i} - \sum_{i=1}^{n} (x_{i} - \bar{x})\bar{y}}{\sum_{i=1}^{n} (x_{i} - \bar{x})^{2}}$$

$$= \frac{\sum_{i=1}^{n} (x_{i} - \bar{x})y_{i}}{\sum_{i=1}^{n} (x_{i} - \bar{x})^{2}} - \frac{\sum_{i=1}^{n} (x_{i} - \bar{x})\bar{y}}{\sum_{i=1}^{n} (x_{i} - \bar{x})y_{i}} = \frac{\sum_{i=1}^{n} (x_{i} - \bar{x})y_{i}}{\sum_{i=1}^{n} (x_{i} - \bar{x})^{2}} - \bar{y} \sum_{i=1}^{n} \omega_{i}$$

$$= \frac{\sum_{i=1}^{n} (x_{i} - \bar{x})y_{i}}{\sum_{i=1}^{n} (x_{i} - \bar{x})y_{i}} - 0 = \sum_{i=1}^{n} \frac{(x_{i} - \bar{x})y_{i}}{\sum_{i=1}^{n} (x_{i} - \bar{x})^{2}} = \sum_{i=1}^{n} \omega_{i}y_{i}$$

so the estimate  $\overset{\wedge}{eta_1}$  is a weighted sum of the observed response  $y_i$ 

- 3. Simple Linear Regression Model The file thuesen contains the blood glucose and short velocity measurements for 25 patients (note that the short velocity data for patient 17 is missing).
- a. Fit a simple linear regression model to the data (use short velocity as the response);
- b. Test the hypothesis that  $\beta_1 = 0$ ;
- c. Calculate the model's  $\mathbb{R}^2$ ;
- d. Examine the residuals. Do you have confidence the linear regression model's assumptions hold?

a.

```
In [2]: thuesen = np.genfromtxt('thuesen.csv', delimiter = ',')
          x = thuesen[1:, 0]
          y = thuesen[1:, 1]
          x = np.delete(x, 15)
          y = np.delete(y, 15)
          beta_1_hat = np.corrcoef(x, y)[0][1] * (np.std(y)/np.std(x))
          beta 0 hat = np.mean(y) - beta 1 hat * np.mean(x)
          print('y = {}x+{}\}'.format(beta_1_hat, beta_0_hat))
          y = 0.021962522259996755x+1.0978148777723817
 In [3]: def linear_regression_model(x):
               y = beta_0_hat + x * beta_1_hat
               return y
          b.
                                        t = \frac{\hat{\beta_1} - \beta_1}{se(\hat{\beta_1})} = \frac{\hat{\beta_1} - \beta_1}{(\frac{\sqrt{MSE}}{\sqrt{\sum(x - \tilde{x})^2}})}
In [11]: y pred = linear regression model(x)
          beta1 = 0
          denominator = 0.0105 # se(beta 1 hat)
          t = (beta 1 hat - beta1)/denominator
In [15]: |print("t = : ", t)
          t = : 2.0916687866663577
In [16]: from scipy import stats
          crit = 2.0739 \# p = 0.05/2, df=22
          if t>crit:
               print('reject H0')
          else:
               print('do not reject H0')
          reject H0
          C.
 In [9]: | SSR = np.sum(np.square(y pred-np.mean(y)))
          SST = np.sum(np.square(y-np.mean(y)))
          R2 = SSR/SST
          print('R2: ',R2)
          R2: 0.17368439567153518
```

localhost:8888/notebooks/Desktop/2022Spring/Regress and Forecasting model/homework3.ipynb

d.

```
residuals = y-y pred
In [18]:
           plt.scatter(x,y)
           a = np.linspace(0,20,200)
           plt.plot(a, linear_regression_model(a))
           plt.show()
            1.8
            1.6
            1.4
            1.2
            1.0
                                           12.5
                      2.5
                           5.0
                                 7.5
                                     10.0
                                                15.0
                                                      17.5
                                                           20.0
                0.0
In [19]: plt.scatter(x, residuals)
           plt.xlabel("blood glucose")
           plt.ylabel('residuals')
Out[19]: Text(0, 0.5, 'residuals')
               0.4
               0.2
           residuals
               0.0
              -0.2
              -0.4
                         6
                               8
                                                           18
                                          12
                                                     16
                                    10
                                               14
                                                                 20
                                     blood glucose
```

The residual is large and does not satisfy homoscedasticity. therefore, I do not have confidence that the linear regression assumption model holds.