```
In [88]: import pandas as pd
from scipy.stats import linregress
import seaborn as sns
import matplotlib.pyplot as plt

df = pd.read_csv('activity7.csv').set_index('Country')
```

1. Use Excel to find the summation of x, y, xy, x^2 , and y^2 .

```
In [89]:
         sumX= df['Birth Rate'].sum()
         sumY = df['Life Expectancy'].sum()
         print("Sum of x: " + str(sumX) + '\nSum of y: '+ str(sumY))
         sumX2 = 0
         sumY2 = 0
         for _,row in df.iterrows():
             sumXY += row.iloc[0]*row.iloc[1]
             sumX2 += row.iloc[0] ** 2
             sumY2 += row.iloc[1] ** 2
         print("Sum of xy: "+str(sumXY) + "\nSum of x^2: " + str(sumX2) + "\nSum of y
        Sum of x: 466
        Sum of y: 1788
        Sum of xy: 34382
        Sum of x^2: 9736
        Sum of y^2: 133512
```

2. Use Excel to find the covariance S_{xy} .

```
In [90]: cov = df.cov()['Birth Rate'].iloc[1]
print(cov)
-14.565217391304348
```

Covariance of xy is -14.562

3. Use Excel to calculate $S_{\boldsymbol{x}}$, $S_{\boldsymbol{y}}$, and the correlation coefficient.

```
In [911: sx = df['Birth Rate'].std()
    sy = df['Life Expectancy'].std()
    print('x devation: '+str(sx))
    print("y deviation: "+str(sy))
    corrCoef = df.corr()['Birth Rate'].iloc[1]
    print("Correlation coeffecient: " + str(corrCoef))

x devation: 5.468619304856508
y deviation: 3.647512553245973
Correlation coeffecient: -0.7302011028563675
```

4. Use Excel to calculate the coefficient of determination.

```
In [921: print("Coefficient of determination: " + str(corrCoef**2))
```

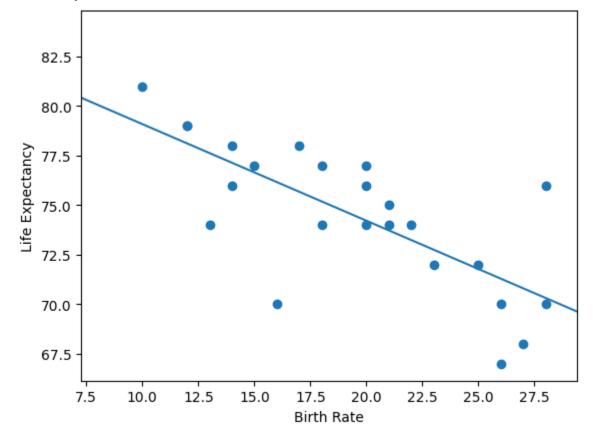
5. Use Excel to calculate b_1 , \boldsymbol{x} , \boldsymbol{y} , and b_0 .

```
In [93]: b1 = cov / sx**2
b0 = df['Life Expectancy'].mean() - (b1*df['Birth Rate'].mean())
print('slope: ' + str(b1) + ' intercept: ' + str(b0))
print(f"Model: ŷ={b0:.3f}+({b1:.3f})x")

fig, ax= plt.subplots()
ax.set_xlabel('Birth Rate')
ax.set_ylabel('Life Expectancy')
ax.axline((0,b0), slope = b1)
scatter = plt.scatter(df['Birth Rate'], df['Life Expectancy'])
ax.set_xlim(df['Birth Rate'].min()-df['Birth Rate'].std()/2)
plt.show()
```

slope: -0.48703658832081403 intercept: 83.9566270898958

Model: $\hat{y}=83.957+(-0.487)x$



6. Given the scatterplot and regression equation you created in Excel and the information you have found up to this point, do you think there is a linear relationship between birthrate and life expectancy?

ANSWER: Yes, it seems linear

7. Given the regression equation above, what is the general trend for life expectancy given a country's birthrate? Is this a positive or negative association?

8. Notice that Paraguay had a birth rate of 28 per 1000 and a life expectancy of 76 years. Find Paraguay's expected life expectancy and the residual. Do you think the model accurately predicted the life expectancy for Paraguay?

```
In [94]: predParaguay = b0 + b1*28
print(f"predicted expectancy: {predParaguay: .3f}, Life expectency standard

predicted expectancy: 70.320, Life expectency standard devation: 3.64751255
3245973
```

ANSWER: The model predicts 70 years of life expentency vs. 76 measured, that is not within a standard devation of what has been measured. Not a good prediction

9. Find another country that the model poorly predicted the life expectancy of. Find the expected life expectancy of the country and the residual

```
In [95]: yStd = df['Life Expectancy'].std()
    print("Bad predictions (outside 1 standard deviation), life expectency, and
    for index,row in df.iterrows():
        y = row.iloc[1]
        pred = b0 + b1*row.iloc[0]
        lowStd = y - yStd
        highStd = y + yStd
        if pred < lowStd or pred > highStd:
            print(f"{index}: predicted life expectency: {round(pred,3)}, residua

Bad predictions (outside 1 standard deviation), life expectency, and residua
l:

Bahamas: predicted life expectency: 76.164, residual: -6.164
Bolivia: predicted life expectency: 71.294, residual: -4.294
Paraguay: predicted life expectency: 70.32, residual: 5.68
```

10. Notice that Nicaragua had a birth rate of 23 per 1000 and a life expectancy of 72 years (23, 72). Find Nicaragua's expected life expectancy and the residual. Do you think the model accurately predicted the life expectancy for Nicaragua?

```
In [96]: pred = pred = b0 + b1*23
    y= 72
    lowStd = pred - yStd
    highStd = pred + yStd
    print(f"Predicted life expectency: {pred:.3f}.\nStandard devation interval f

Predicted life expectency: 72.755.
    Standard devation interval from prediction: (76.4,69.1)
```

ANSWER: Yes, measured expected life expectency is within 1 standard devation of predicted.

11. Find another country that the model accurately predicted the life expectancy of. Find the expected life expectancy of the country and the residual.

```
print("Good predictions (within 1 standard devation of prediction): \n")
In [97]:
         for index,row in df.iterrows():
             y = row.iloc[1]
             pred = b0 + b1*row.iloc[0]
             lowStd = y - yStd
             highStd = y + yStd
             if pred > lowStd and pred < highStd:</pre>
                 print(f"{index}: predicted life expectency: {round(pred,3)}, residual
        Good predictions (within 1 standard devation of prediction):
        Argentina: predicted life expectency: 75.19, residual: 1.81
        Barbados: predicted life expectency: 77.625, residual: -3.625
        Belize: predicted life expectency: 70.807, residual: -2.807
        Canada: predicted life expectency: 79.086, residual: 1.914
        Chile: predicted life expectency: 76.651, residual: 0.349
        Colombia: predicted life expectency: 75.19, residual: -1.19
        Costa Rica: predicted life expectency: 75.677, residual: 2.323
        Dominican Republic: predicted life expectency: 73.242, residual: 0.758
        Ecuador: predicted life expectency: 73.729, residual: 1.271
        El Salvador: predicted life expectency: 71.781, residual: 0.219
        Guatemala: predicted life expectency: 70.32, residual: -0.32
        Honduras: predicted life expectency: 71.294, residual: -1.294
        Jamaica: predicted life expectency: 74.216, residual: -0.216
        Mexico: predicted life expectency: 74.216, residual: 1.784
        Nicaragua: predicted life expectency: 72.755, residual: -0.755
        Panama: predicted life expectency: 74.216, residual: 2.784
        Puerto Rico: predicted life expectency: 78.112, residual: 0.888
        United States: predicted life expectency: 77.138, residual: 0.862
        Uruguay: predicted life expectency: 77.138, residual: -1.138
        Venezuela: predicted life expectency: 73.729, residual: 0.271
        Virgin Islands: predicted life expectency: 78.112, residual: 0.888
```