

2. Covariance and Correlation

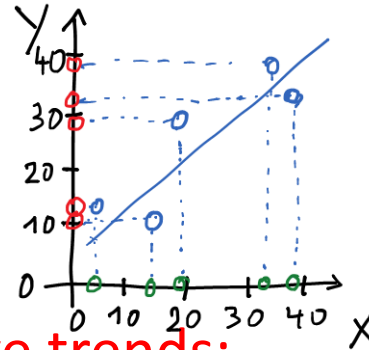
- ✓ Covariance
- ✓ Correlation

Covariance

Covariance can classify three relationships between variables:

1. Relationships with **positive trends**:

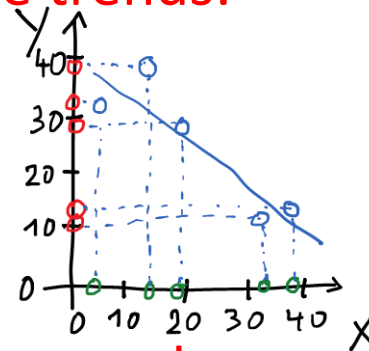
Covariance is POSITIVE



Example:
mRNA of gene X in 5 different cells (X-axis)
and mRNA of gene Y in **SAME** 5 different cells (Y-axis)

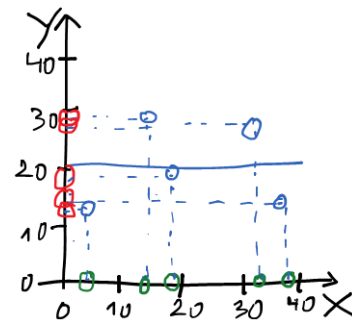
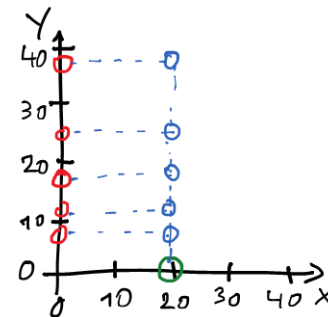
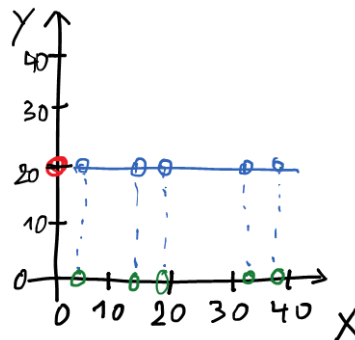
2. Relationships with **negative trends**:

Covariance is NEGATIVE



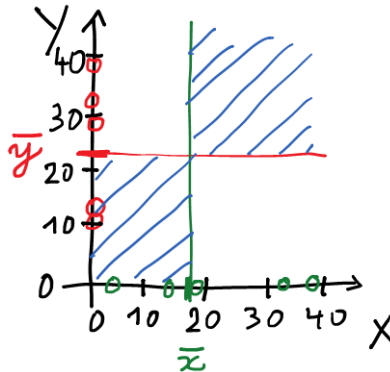
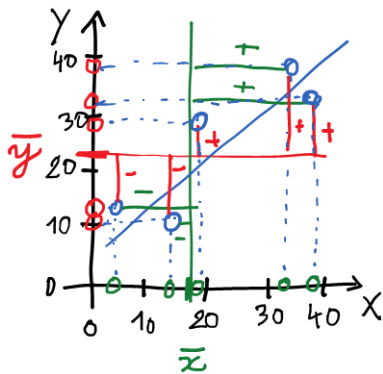
3. No relationship because **no trend**:

Covariance = 0

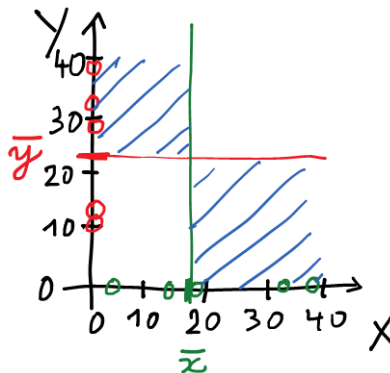
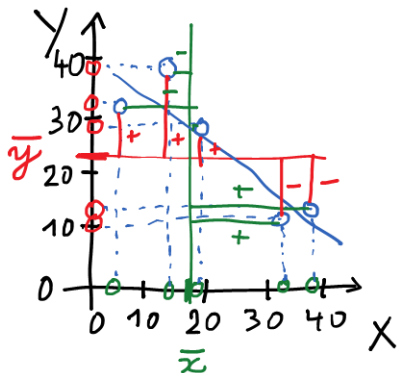


$$\text{Covariance} = \frac{\sum (x - \bar{x})(y - \bar{y})}{n - 1} = \frac{SP_{xy}}{n - 1}$$

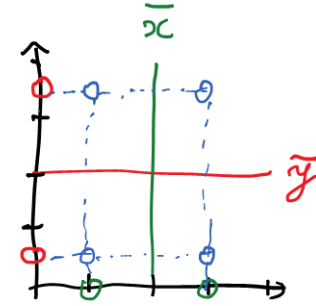
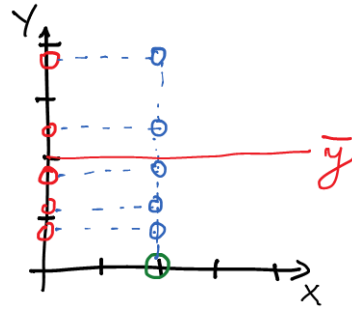
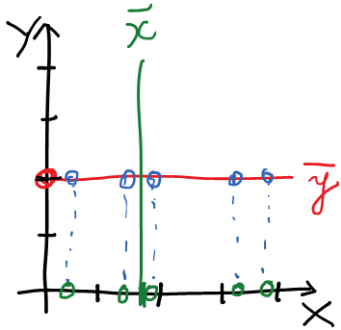
*SP_{xy} refers to the **sum of cross products***



Data in these two quadrants contribute **positive values** to the total covariance.



Data in these two quadrants contribute **negative values** to the total covariance.



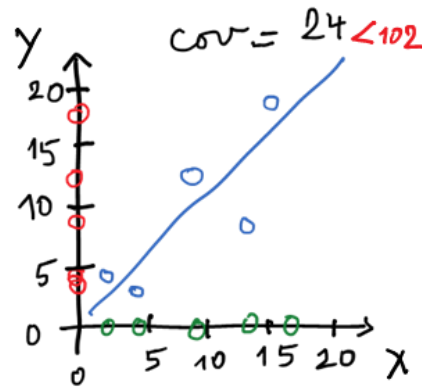
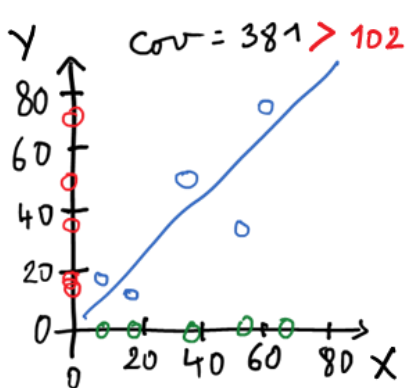
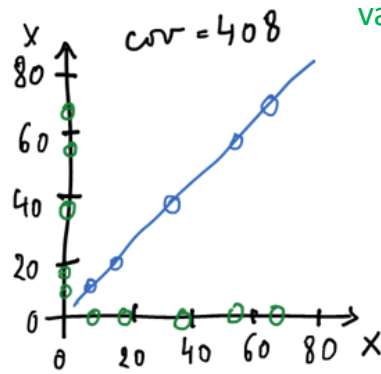
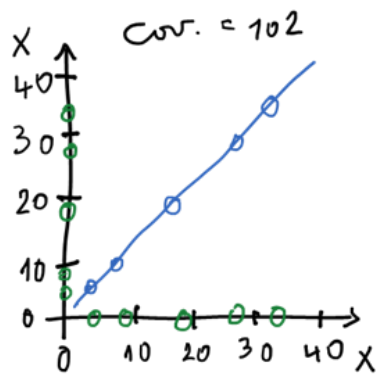
NO RELATIONSHIP between X and Y

$$\text{Covariance} = \frac{\sum (x - \bar{x})(y - \bar{y})}{n - 1} = 0$$

Covariance cannot quantify the strength of a relationship

The covariance for gene X with itself is the same thing as the estimated variance for gene X

$$\frac{\sum (x - \bar{x})(y - \bar{y})}{n - 1} = \frac{\sum (x - \bar{x})^2}{n - 1} = \sigma^2$$



- If you **change the scale** that the data is on (all values x2), the relationship does not change, BUT the **covariance changes!**
- Sensitivity to the scale makes that the covariance **cannot tell if the data are close to the line** representing the relationship, or far from that line.
- Covariance **doesn't tell anything about the slope** of the line representing the relationship.

COVARIANCE JUST TELLS IF THE RELATIONSHIP IS POSITIVE OR NEGATIVE

→ Covariance is a computational stepping stone to something that is interesting, like correlation or PCA.

Correlation

Correlation

- Correlation describes relationships between variables and is **NOT sensitive to the scale** of the data

→ THEREFORE: correlation can **quantify the strength of a relationship.**

- Correlation =
$$\frac{\text{Covariance (X,Y)}}{\sqrt{\text{Variance(X)}}\sqrt{\text{Variance(Y)}}}$$

Numerator: any value between $-\infty$ and $+\infty$, depending on

- 1) whether the slope of the line that represents the relationship is positive or negative
- 2) how far the data are spread around the means
- 3) the scale of the data

$$\text{Correlation} = \frac{\text{Covariance (X,Y)}}{\sqrt{\text{Variance(X)}}\sqrt{\text{Variance(Y)}}}$$

“How much of the variance in the data (X and Y) is accounted for by the covariance?”

Denominator: squeezes the covariance to be a number between

– 1 and +1

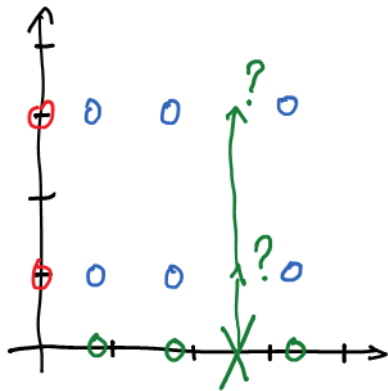
→ ensures that the scale of the data does not affect correlation

→ Correlations are easier to interpret than covariance.

1. When there is NO RELATIONSHIP between X and Y, none of the variance in the data is accounted for by the covariance (covariance is 0), the **correlation is 0**.

Covariance = 0

$$\Leftrightarrow \text{Correlation} = \frac{0}{\sqrt{\text{Variance}(X)}\sqrt{\text{Variance}(Y)}} = 0$$

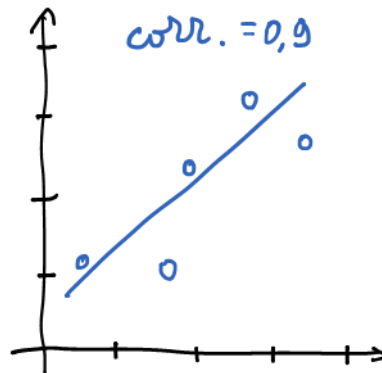


- If correlation = 0
 \Leftrightarrow a value on the X-axis doesn't tell us anything about what to expect on the Y-axis, because there is no reason to choose one value or another.

2. When we observe a trend in the data, BUT the data do not all fall on a straight line, the covariance accounts for some but not all of the variance in the data, and the correlation gets closer to 1 or -1.

$$|\text{Covariance}| < \sqrt{\text{Variance}(X)}\sqrt{\text{Variance}(Y)}$$

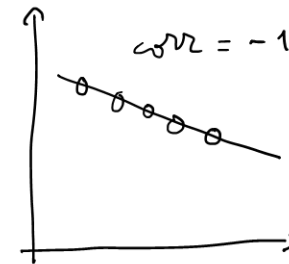
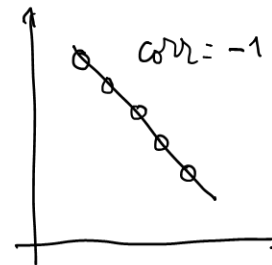
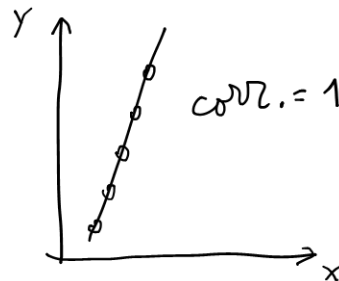
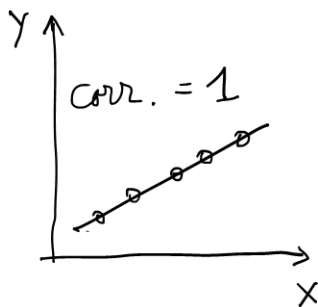
$$\Leftrightarrow -1 < \left[\text{Correlation} = \frac{\text{Covariance}(X,Y)}{\sqrt{\text{Variance}(X)}\sqrt{\text{Variance}(Y)}} \right] < 1$$



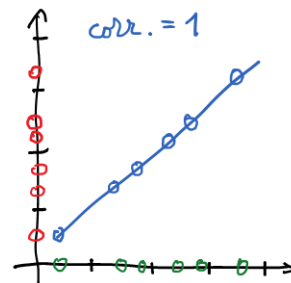
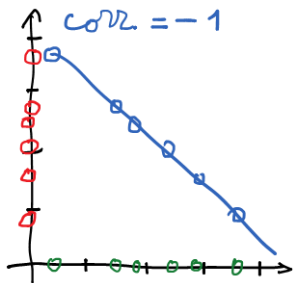
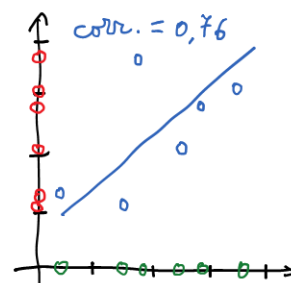
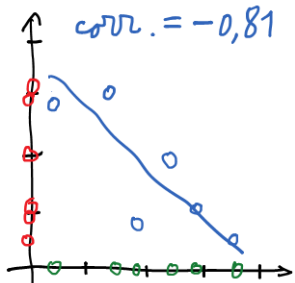
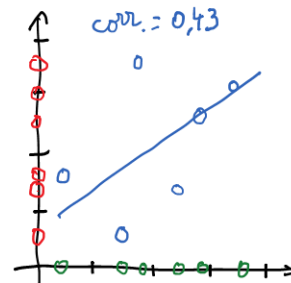
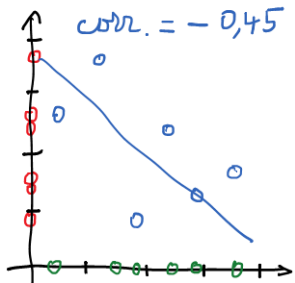
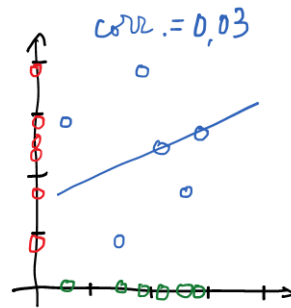
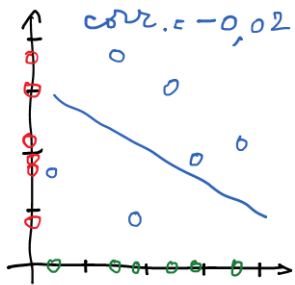
3. When all the data fall on a straight line with positive or negative slope, the covariance accounts for all the variance in the data, and the correlation is 1 or -1.

$$|\text{Covariance}| = \sqrt{\text{Variance}(X)}\sqrt{\text{Variance}(Y)}$$

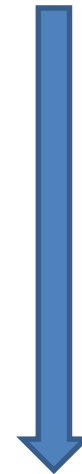
$$\Leftrightarrow \text{Correlation} = \frac{\text{Covariance}(X,Y)}{\sqrt{\text{Variance}(X)}\sqrt{\text{Variance}(Y)}} = \pm 1$$



Slope can be large or small



Correlation can quantify the strength of a relationship.

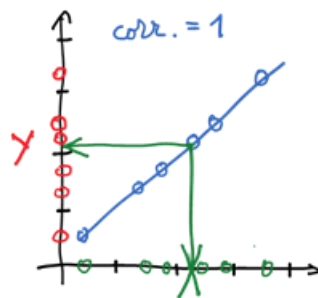
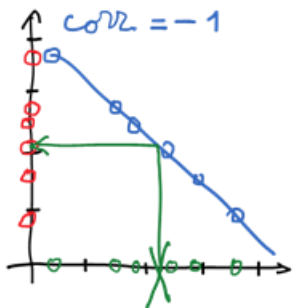
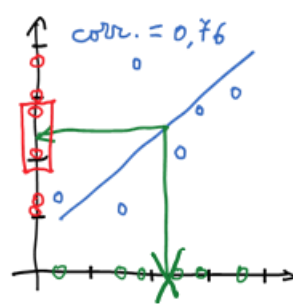
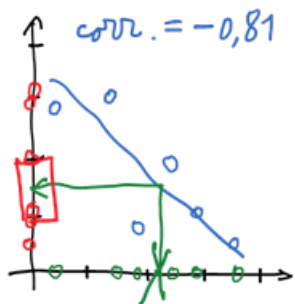
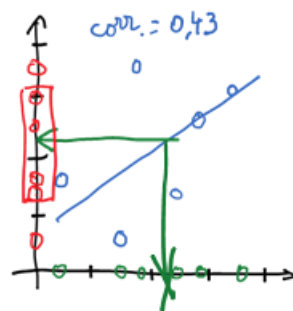
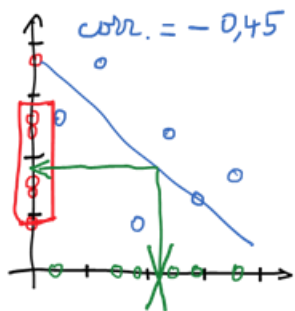
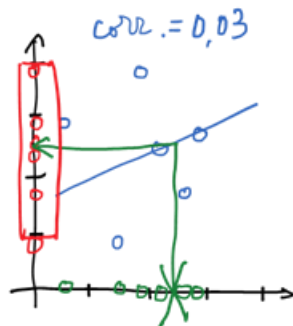
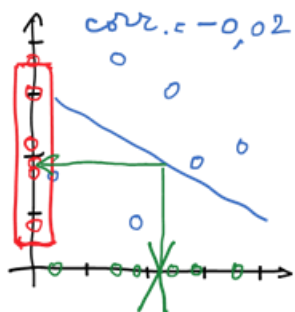


Data closer to the trend line with negative slope

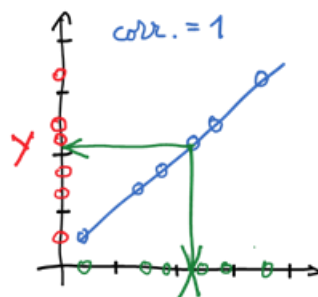
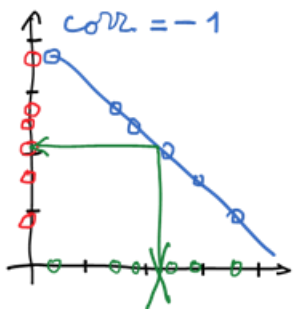
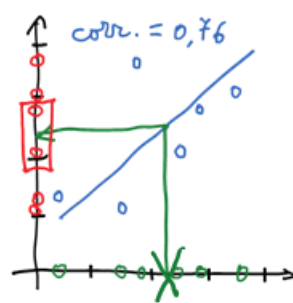
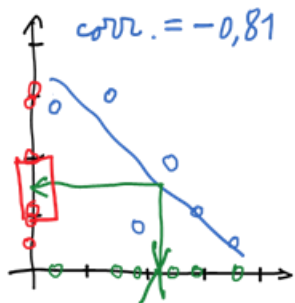
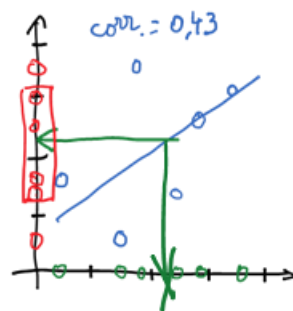
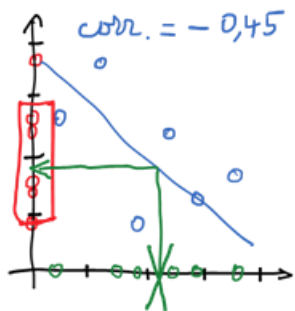
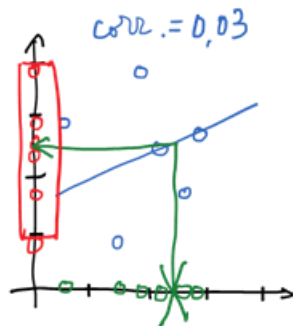
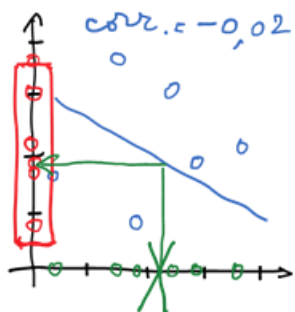
⇔ correlation closer to -1

Data closer to the trend line with positive slope

⇔ correlation closer to 1

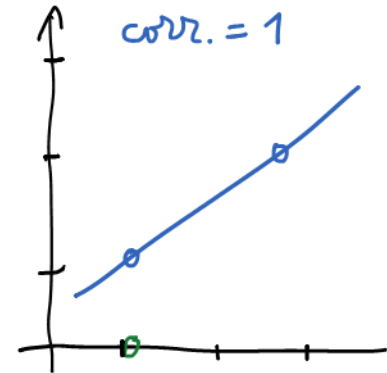
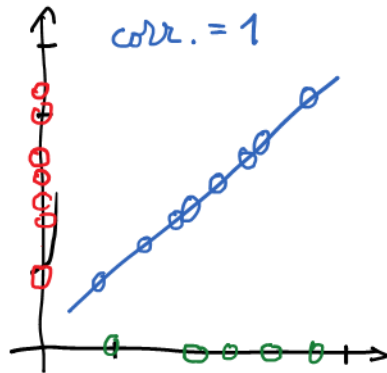


- If correlation $\neq 0$
 \Leftrightarrow the trend can be used to make predictions (educated guesses).
- When correlation values get closer to -1 or 1; the guesses become more refined.



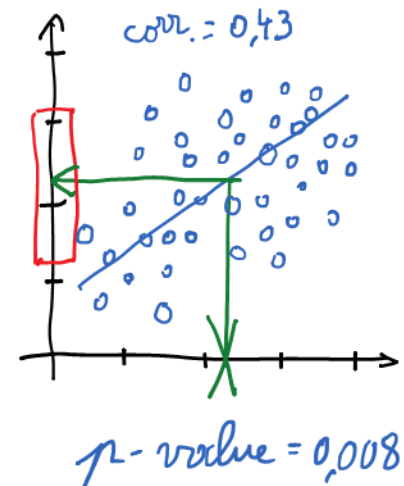
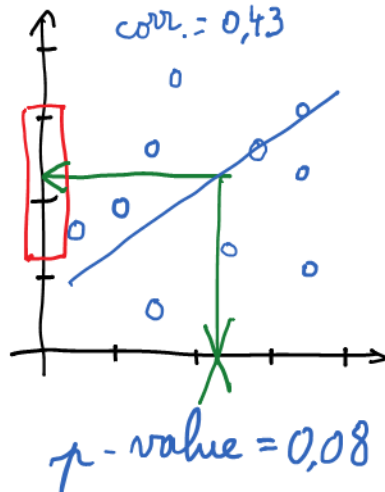
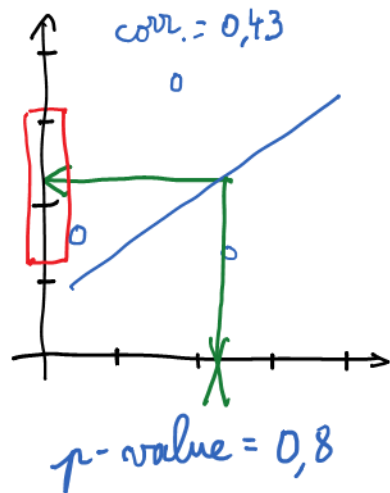
- If we have a new measurement for gene X, then we can use the trend line to **predict a range of values** for gene Y.
- If we have a new measurement for gene Y, then we can use the trend line to **predict a range of values** for gene X.
- **If the data are closer to the trend line, then**
 - there is a stronger relationship between X and Y.
 - values for gene X tell us more about gene Y (and *vice versa*).
 - **given a value for gene X, we predict that gene Y falls in a narrower range.**

The confidence in how useful the relationship is depends on the amount of data



- When all data are on a straight line, correlation is 1 or -1, regardless of how much data we have.
- **Two datapoints** are ALWAYS on a straight line!
 - correlation is 1 or -1
 - This makes the relationship appear strong
 - But we should not have any confidence in predictions made with this line.
- The probability that we can draw a straight line through a number of points gets smaller with each additional point.
 - The more data we have, the more confidence we have in the predictions we make with the line.

- The **p-value for correlation** tells the probability that randomly drawn dots will result in a similarly strong relationship, or stronger.
- Smaller p-value \Leftrightarrow more confidence in the predictions made with the line



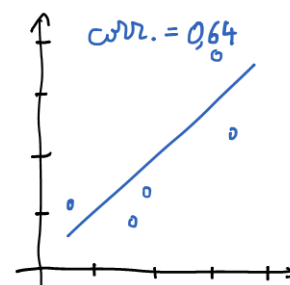
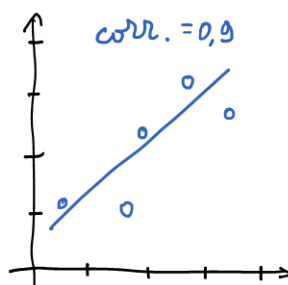
Correlation value 0.43 is small: all three graphs represent BAD GUESSES.
BUT most confidence in the bad guess that came from most data.

Conclusion:

Even if you have a lot of data and therefore a lot of confidence in your guess (low p-value), **if the correlation value is close to 0, your guess will still be bad!**

Correlations are not the most easy to interpret.

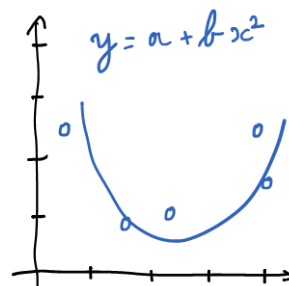
- Not obvious that the relationship with correlation value 0.9 is twice as good at making predictions as the relationship with correlation value 0.64



- R^2 solves this problem

$$> 0.9^2 / 0.64^2$$
$$[1] \quad 1.977539$$

- R^2 can also quantify relationships that are more complicated than straight lines!



(see later!)

Parameters for populations

(→ using Greek letters!)

Covariance between x and y: σ_{xy}

$$\sigma_{xy} = \left(\sum_{i=1}^N (y_i - \mu_y)(x_i - \mu_x) \right) / N$$

can be negative!

Correlation (Pearson's) between two variables, *y and x*: ρ

$$\rho_{xy} = \frac{\sigma_{xy}}{\sqrt{\sigma_x^2 \times \sigma_y^2}}$$

Standardize the covariance

Ranges from -1 to +1; with strong negative correlations near to -1 and strong positive correlations near to +1.

Statistics from the sample

(= estimates of the population parameters!)

Covariance between x and y: s_{xy}

$$s_{xy} = \left(\sum_{i=1}^n (y_i - \bar{y})(x_i - \bar{x}) \right) / (n - 1)$$

Correlation (Pearson's) between two variables, *y* and *x*: r

$$r_{xy} = \frac{s_{xy}}{\sqrt{s_x^2 \times s_y^2}}$$

Ranges from -1 to +1; with strong negative correlations near to -1 and strong positive correlations near to +1.