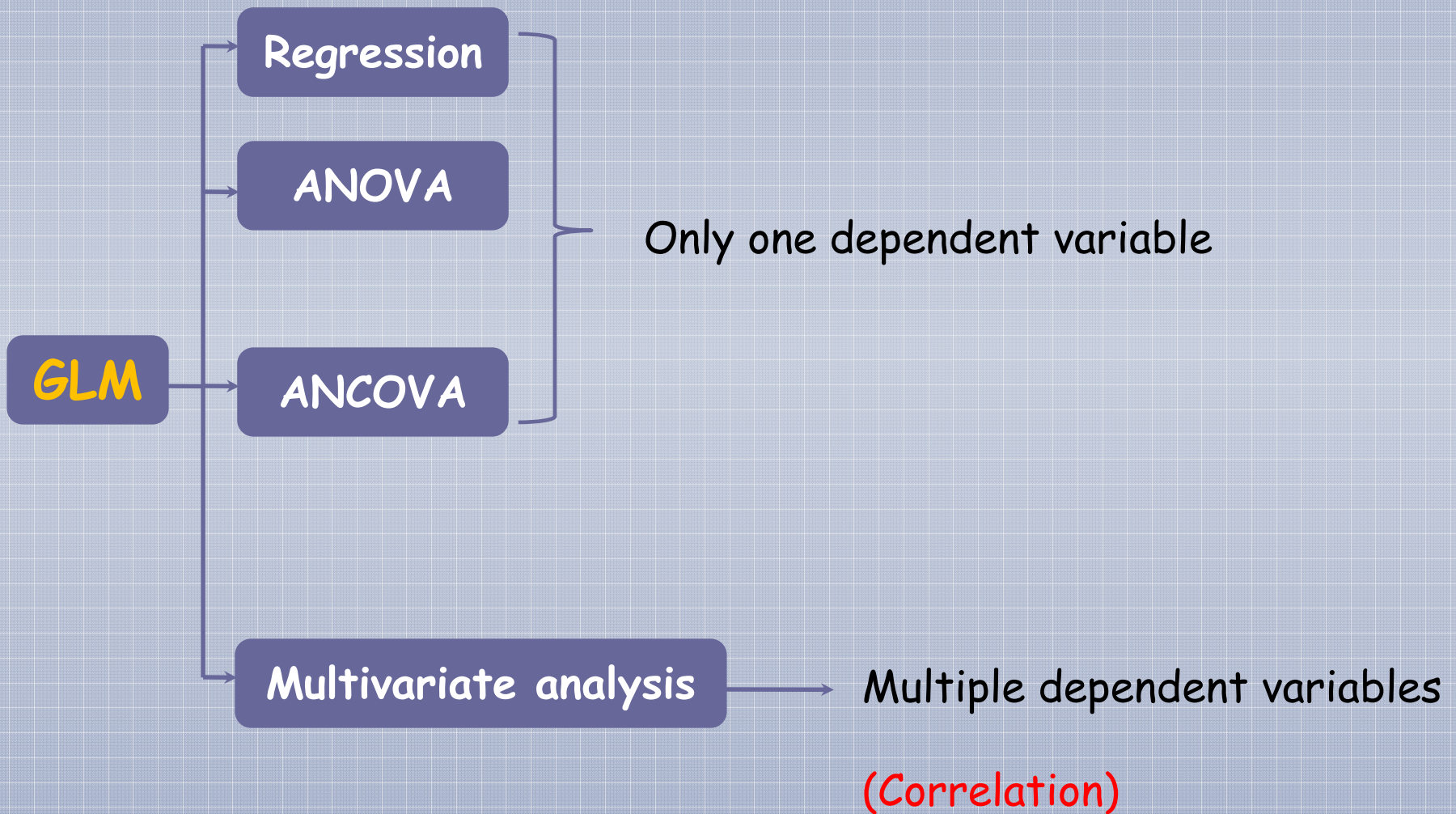


BIOL 4605/7220
CH 20.1 Correlation

GPT Lectures
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November 9, 2011

GLM: correlation



Correlation

- ❖ Two variables associated with each other?
- ❖ No casual ordering (i.e., NEITHER is a function of the other)

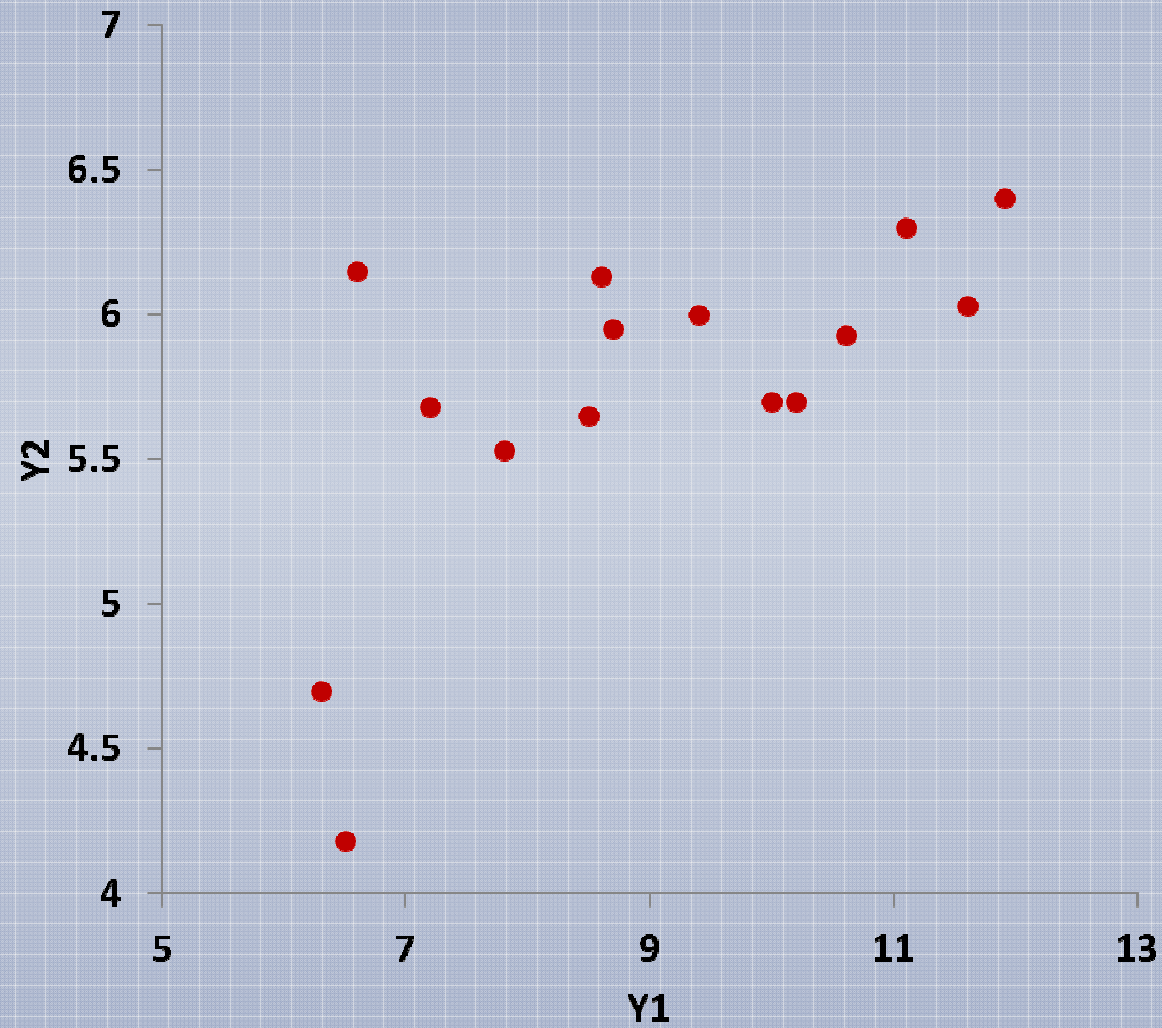
Y_1 – Total length of aphid stem mothers

Y_2 – Mean thorax length of their parthenogenetic offspring

[Data](#) from Box 15.4 Sokal and Rohlf 2012

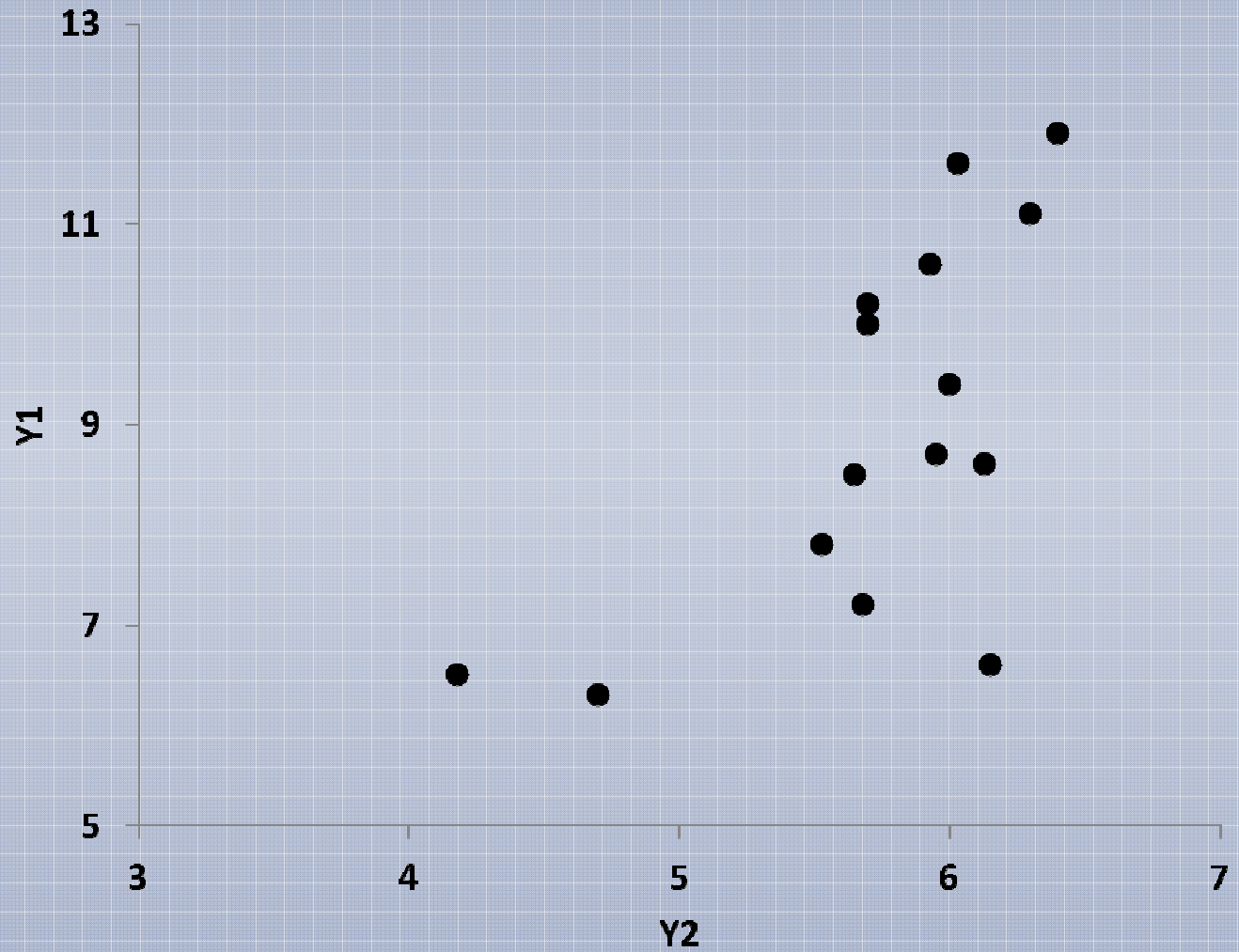
Correlation

Y_2 vs. Y_1



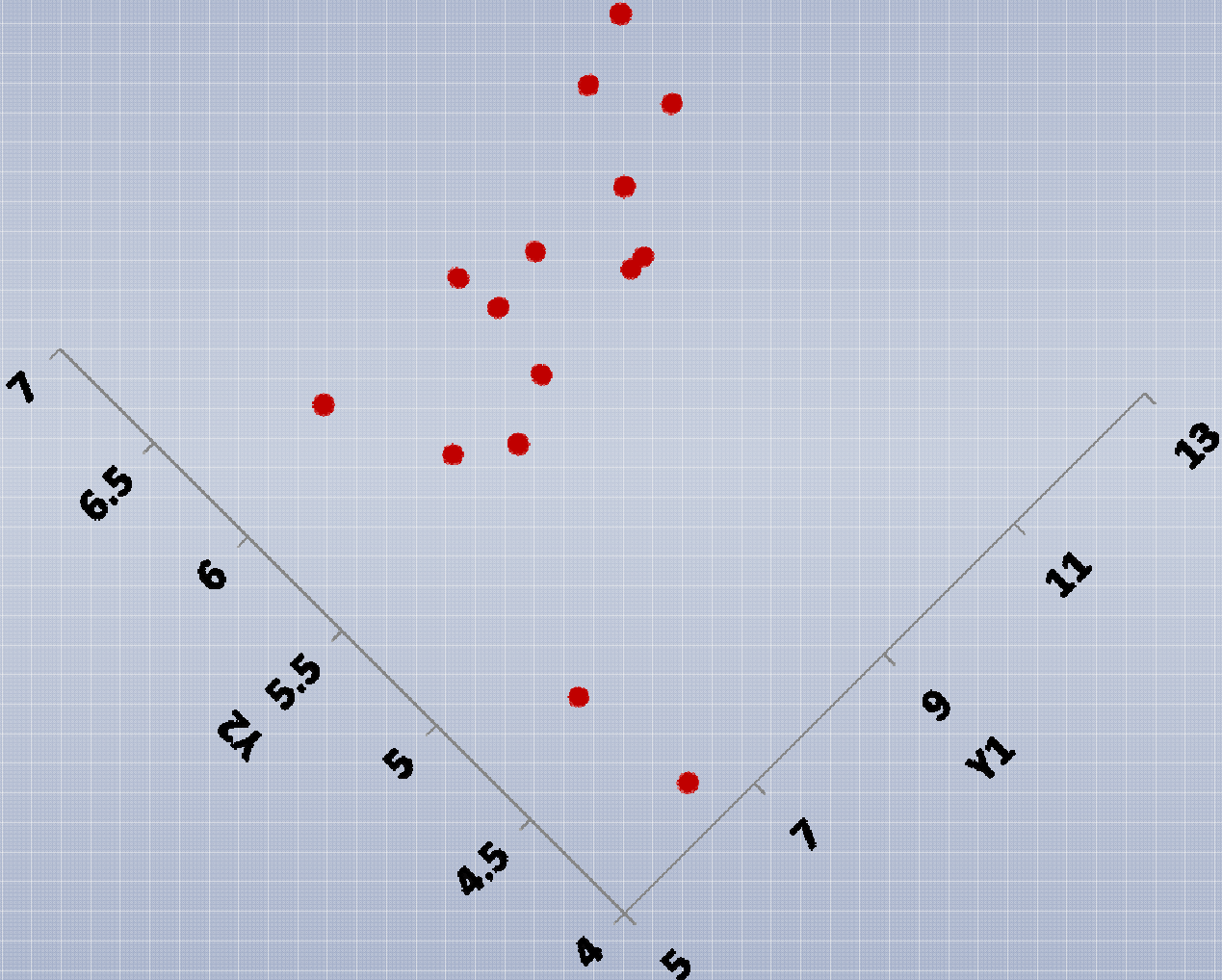
Correlation

Y_1 vs. Y_2



Correlation

Rotate



Regression vs. Correlation

Regression

- Does Y depend on X ?
(describe func. relationship/predict)
- Usually, X is manipulated & Y is a random variable
- Casual ordering $Y=f(X)$

Correlation

- Are Y_1 and Y_2 related?
- Both Y_1 & Y_2 are random variables
- No casual ordering

Correlation: parametric vs. non-parametric

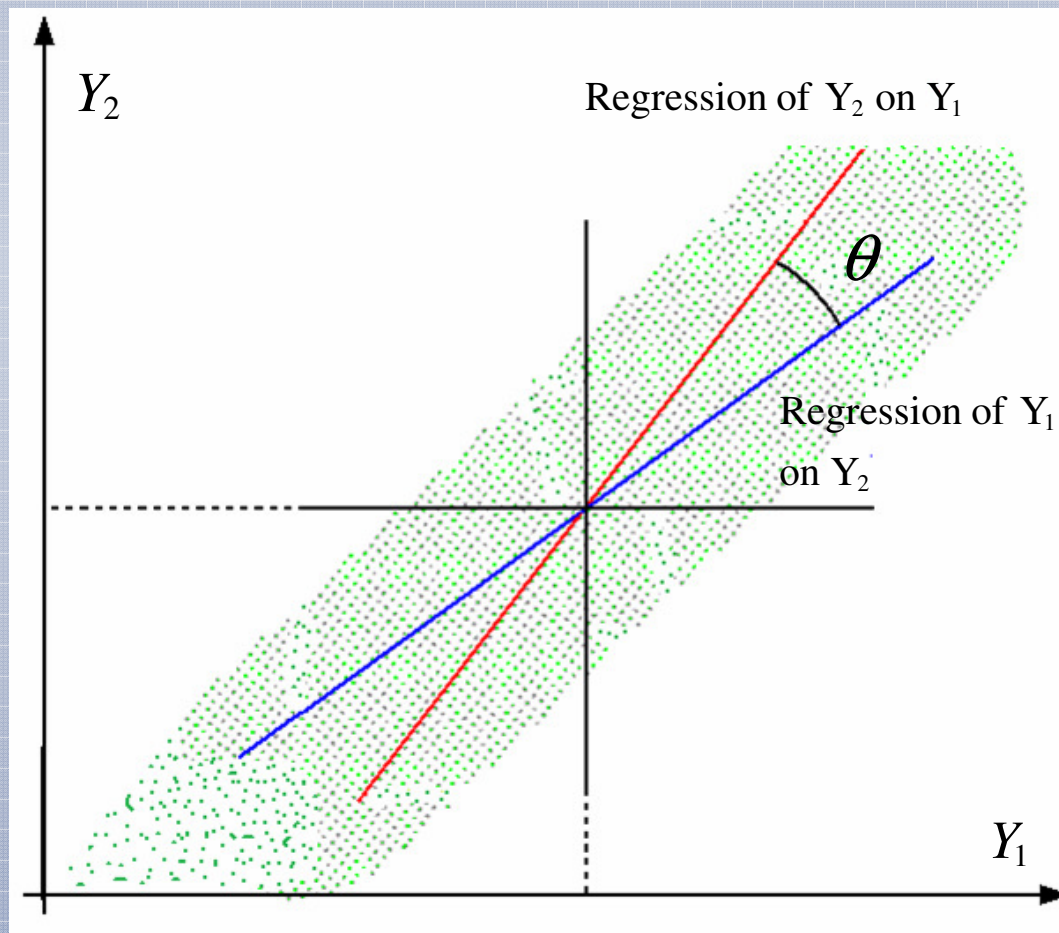
Parametric measures: Pearson's correlation

Nonparametric measures: Spearman's Rho, Kendall's Tau

Type of data	Measures of correlation
Measurements (from Normal/Gaussian Population)	<u>Parametric</u> : Pearson's correlation
Ranks, Scores, or Data that do not meet assumptions for sampling distribution (t , F , χ^2)	<u>Nonparametric</u> : Spearman's Rho, Kendall's Tau

Pearson's Correlation Coefficient (ρ)

- Strength of relation between two variables Y_1 & Y_2
- Geometric interpretation



$$\rho = \cos(\theta)$$

- **Perfect positive association:**
 $\theta = 0^\circ \quad \rho = 1$
- **No association:**
 $\theta = 90^\circ \quad \rho = 0$
- **Perfect negative association:**
 $\theta = 180^\circ \quad \rho = -1$

$-1 \leq \rho \leq 1$, true relation

Pearson's Correlation Coefficient (ρ)

- Strength of relation between two variables Y_1 & Y_2
- Geometric interpretation
- Definition

$$\rho_{Y_1, Y_2} = \frac{\text{cov}(Y_1, Y_2)}{\sigma_{Y_1} \sigma_{Y_2}} = \frac{E[(Y_1 - \mu_{Y_1})(Y_2 - \mu_{Y_2})]}{\sigma_{Y_1} \sigma_{Y_2}}$$

Covariance of the two variables divided by the product of their standard deviations

Pearson's Correlation Coefficient (ρ)

- Strength of relation between two variables Y_1 & Y_2
- Geometric interpretation
- Definition
- Estimate ($\hat{\rho} = r$) from a sample

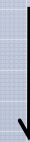
Parameter		Estimate
Name	Symbol	
<i>Mean of Y_1</i>	μ_{Y_1}	\bar{Y}_1
<i>Mean of Y_2</i>	μ_{Y_2}	\bar{Y}_2
<i>Variance of Y_1</i>	$\sigma_{Y_1}^2$	$s_{Y_1}^2$
<i>Variance of Y_2</i>	$\sigma_{Y_2}^2$	$s_{Y_2}^2$

Pearson's Correlation Coefficient (ρ)

- Strength of relation between two variables Y_1 & Y_2
- Geometric interpretation
- Definition
- Estimate ($\hat{\rho} = r$) from a sample

Parameter	Estimate
μ_{Y_1}	\bar{Y}_1
μ_{Y_2}	\bar{Y}_2
$\sigma_{Y_1}^2$	$s_{Y_1}^2$
$\sigma_{Y_2}^2$	$s_{Y_2}^2$

$$\rho_{Y_1, Y_2} = \frac{\text{cov}(Y_1, Y_2)}{\sigma_{Y_1} \sigma_{Y_2}} = \frac{E[(Y_1 - \mu_{Y_1})(Y_2 - \mu_{Y_2})]}{\sigma_{Y_1} \sigma_{Y_2}}$$



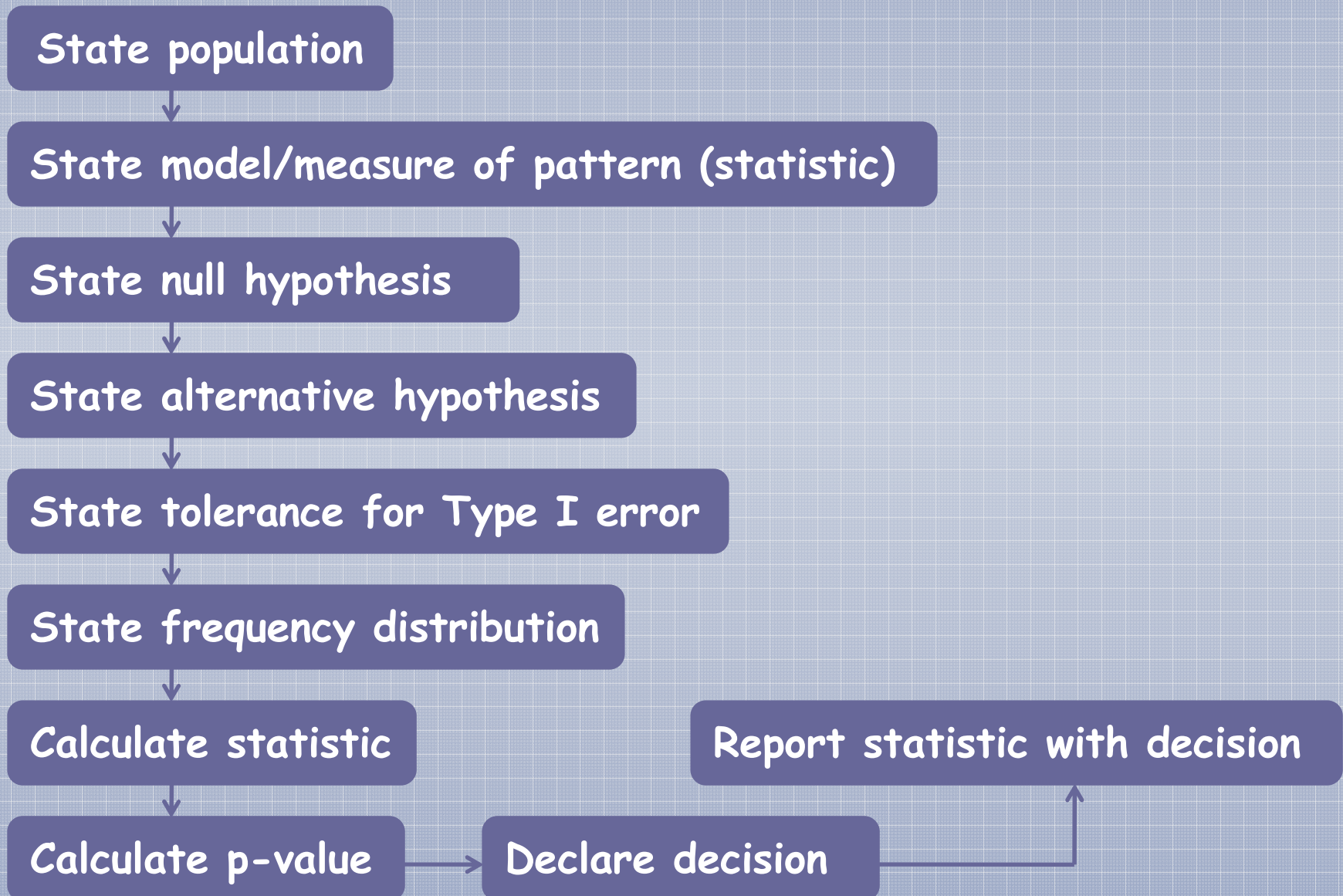
$$r = \hat{\rho} = \frac{1}{n-1} \cdot \frac{\sum_i (Y_{1i} - \bar{Y}_1)(Y_{2i} - \bar{Y}_2)}{s_{Y_1} s_{Y_2}} = \frac{\sum_i (Y_{1i} - \bar{Y}_1)(Y_{2i} - \bar{Y}_2)}{\sqrt{\sum_i (Y_{1i} - \bar{Y}_1)^2 \sum_i (Y_{2i} - \bar{Y}_2)^2}}$$



Pearson's Correlation: Significance Test

- Determine whether a sample correlation coefficient could have come from a population with a parametric correlation coefficient of ZERO
- Determine whether a sample correlation coefficient could have come from a population with a parametric correlation coefficient of CERTAIN VALUE $\neq 0$
- Generic recipe for Hypothesis Testing

Hypothesis Testing --- Generic Recipe



Hypothesis Testing --- Generic Recipe

State population

All measurements on total length of aphid stem mothers & mean thorax length of their parthenogenetic offspring made by the same experimental protocol

- 1). Randomly sampled
- 2). Same environmental conditions

Hypothesis Testing --- Generic Recipe

State population

State model/measure of pattern (statistic)

- Correlation of the two variables, ρ
- In the case $H_0 : \rho = 0$

$$t = \frac{r - \rho}{\sqrt{\frac{1-r^2}{n-2}}} \quad \left(r = \hat{\rho}, \sim \begin{cases} 1) N\left(0, \frac{1-r^2}{n-2}\right), \text{if } n \text{ LARGE} \\ 2) t\text{-distribution, } df = n-2, \text{ otherwise} \end{cases} \right)$$

- In the case $H_0 : \rho = \rho_1$ ($\rho_1 \neq 0$)

$$t = \frac{z - \eta}{1/\sqrt{n-3}} \quad \left(\eta = \frac{1}{2} \ln \left(\frac{1+\rho_1}{1-\rho_1} \right), \text{ where } z = \frac{1}{2} \ln \left(\frac{1+r}{1-r} \right), E(z) = \eta, \text{ var}(z) = \frac{1}{n-3} \right)$$

z : Normal/tends to normal rapidly as n increases for $\rho \neq 0$
t-statistic: $N(0, 1)$ or t ($df = \infty$)

Hypothesis Testing --- Generic Recipe

State population

State model/measure of pattern (statistic)

- Correlation of the two variables, ρ
- In the case $H_0 : \rho = 0$

$$t = \frac{r - \rho}{\sqrt{\frac{1 - r^2}{n - 2}}}$$

$$\left(r = \hat{\rho}, \sim \begin{cases} 1) N\left(0, \frac{1 - r^2}{n - 2}\right), \text{ if } n \text{ LARGE} \\ 2) t\text{-distribution, } df = n - 2 \end{cases} \right)$$

Hypothesis Testing --- Generic Recipe

State population



State model/measure of pattern (statistic)



State null hypothesis



$$H_0 : \rho = 0$$

Hypothesis Testing --- Generic Recipe

State population

State model/measure of pattern (statistic)

State null hypothesis

State alternative hypothesis

$$H_A : \rho \neq 0$$

Hypothesis Testing --- Generic Recipe

State population

State model/measure of pattern (statistic)

State null hypothesis

State alternative hypothesis

State tolerance for Type I error

$\alpha = 5%$ (*conventional level*)

Hypothesis Testing --- Generic Recipe

State population



State model/measure of pattern (statistic)



State null hypothesis



State alternative hypothesis



State tolerance for Type I error



State frequency distribution



t-distribution

Hypothesis Testing --- Generic Recipe

State population

State model/measure of pattern (statistic)

State null hypothesis

State alternative hypothesis

State tolerance for Type I error

State frequency distribution

Calculate statistic

- t-statistic
- correlation coefficient estimate, $r = 0.65$
- $t = (0.65 - 0)/0.21076 = 3.084$

Hypothesis Testing --- Generic Recipe

State population



State model/measure of pattern (statistic)



State null hypothesis



State alternative hypothesis



State tolerance for Type I error



State frequency distribution



Calculate statistic



Calculate p-value

- $t = 3.084, df = 13$
- $p = 0.0044$ (one-tail) & 0.0088 (two-tail)

Hypothesis Testing --- Generic Recipe

State population

State model/measure of pattern (statistic)

State null hypothesis

State alternative hypothesis

State tolerance for Type I error

State frequency distribution

Calculate statistic

Calculate p-value

Declare decision

- $p = 0.0088 < \alpha = 0.05$
- reject H_0
- accept $H_A : \rho \neq 0$

Hypothesis Testing --- Generic Recipe

State population

State model/measure of pattern (statistic)

State null hypothesis

State alternative hypothesis

State tolerance for Type I error

State frequency distribution

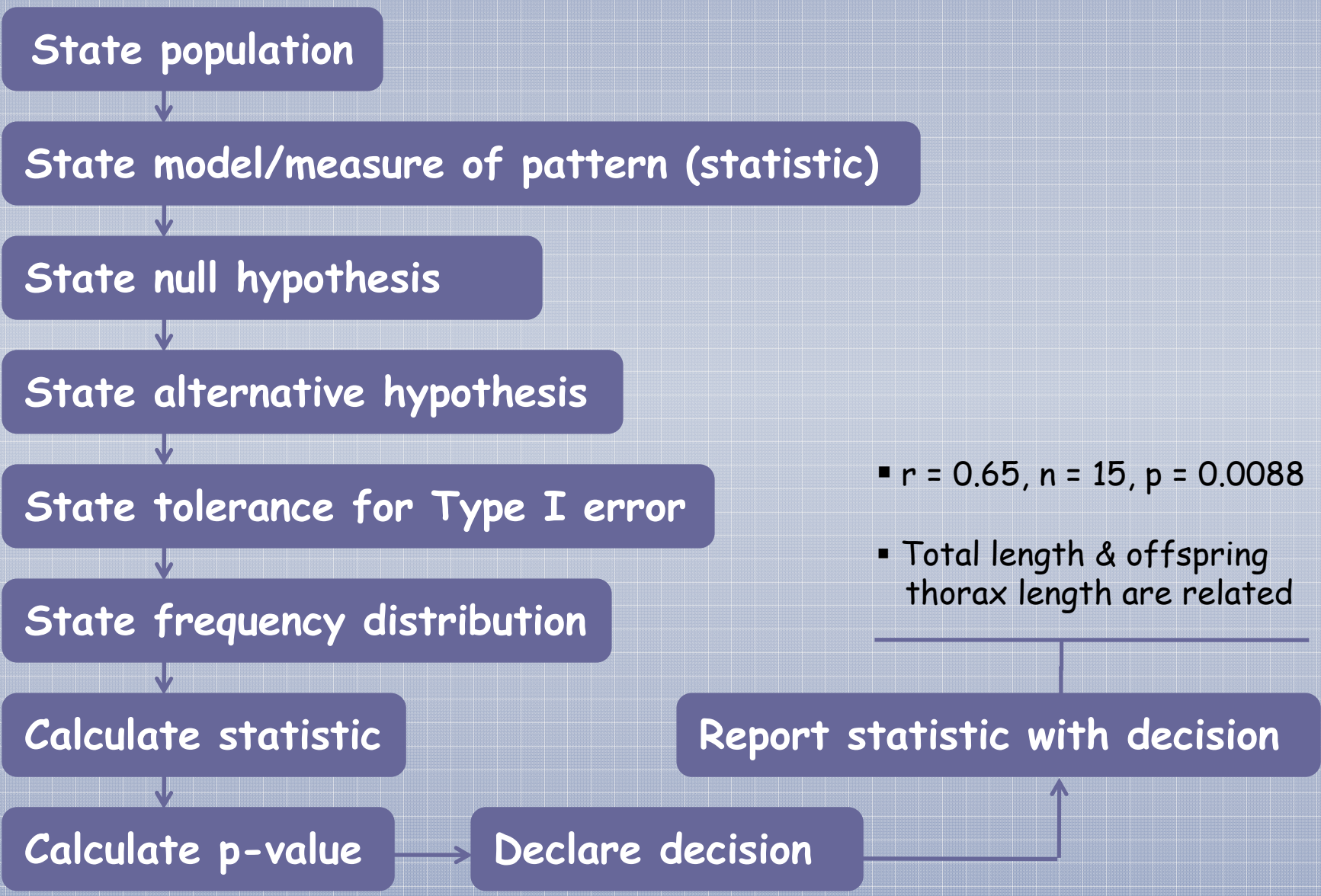
Calculate statistic

Calculate p-value

Declare decision

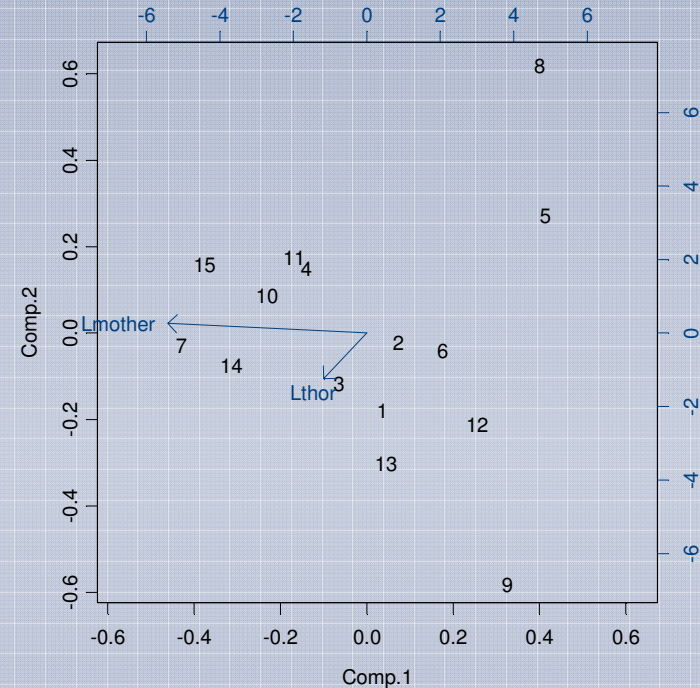
Report statistic with decision

- $r = 0.65, n = 15, p = 0.0088$
- Total length & offspring thorax length are related



Pearson's Correlation - Assumptions

- Assumptions
- Normal & independent errors
- Homogeneous around straight line

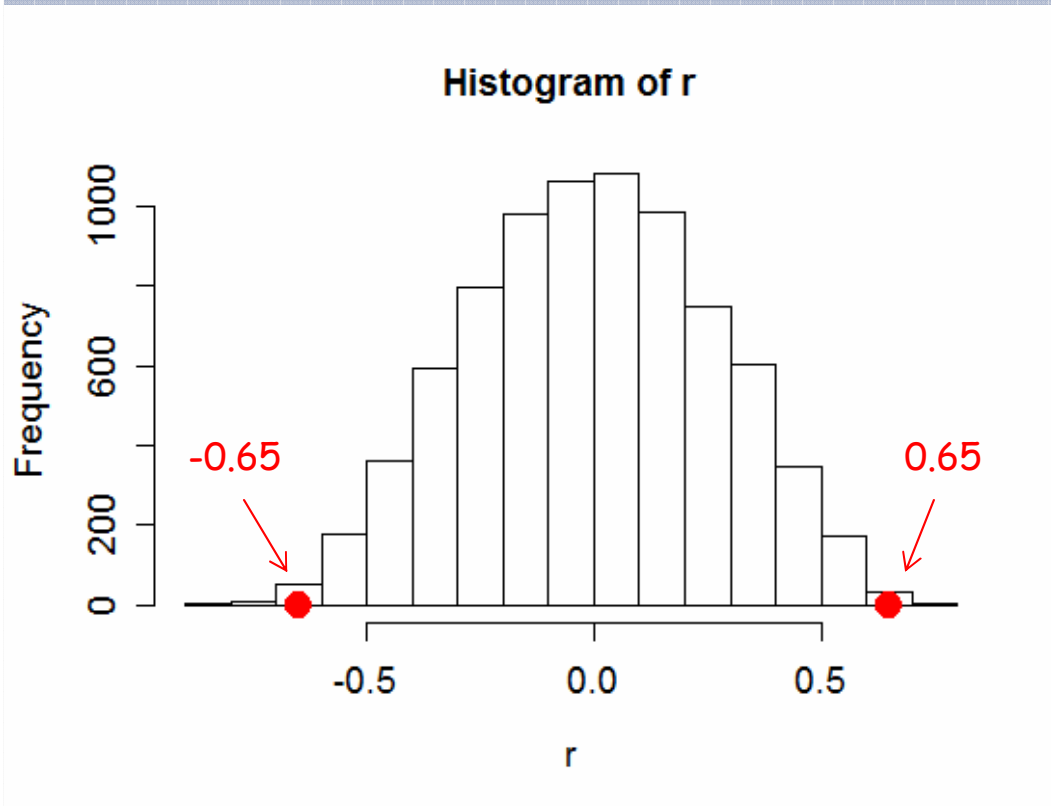


- What if assumptions for Pearson test not met?
- Here are the observations relative to the correlation line (comp 1)
- Not homogeneous, due to outliers (observations 8 & 9)

Pearson's Correlation - Randomization test

- Significance test with no distributional assumptions
- Hold one variable, permute the other one many times
- A new r from each new permutation
- Construct empirical frequency distribution
- Compare the empirical distribution with the observed r

Pearson's Correlation - Randomization test



- 8000 times
- $p_1 = p(r > 0.65) = 0.001875$
- $p_2 = p(r < -0.65) = 0.003875$
- $p = p_1 + p_2 = 0.00575 < \alpha = 0.05$
- Reject Null, accept alternative
- Consistent with testing result from theoretical t-distribution, for this data

Pearson's Correlation coefficient - Confidence Limit

- 95% confidence limit (tolerance of Type I error @ 5%)
- t-distribution (df = n - 2) (NO)

a). $H_0: \rho = 0$ was rejected

b). Distribution of r is negatively skewed

c). Fisher's transformation

- $z = \frac{1}{2} \ln\left(\frac{1+r}{1-r}\right); \frac{z-\eta}{1/\sqrt{n-3}} \sim N(0, 1) \text{ or } t_{[\infty]}$

$$\eta = \frac{1}{2} \ln\left(\frac{1+\rho_1}{1-\rho_1}\right)$$

Pearson's Correlation coefficient - Confidence Limit

C. I. for η :

- $$\begin{cases} z_l = z - z_{(1-\alpha/2)} \cdot \sqrt{1/(n-3)} \\ z_u = z + z_{(1-\alpha/2)} \cdot \sqrt{1/(n-3)} \end{cases}, z_{(1-\alpha/2)}, \text{ critical value from } N(0, 1) \text{ at } p = 1-\alpha/2$$

C. I. for ρ :

- $$\begin{cases} r_l = \tanh(z_l) = \frac{\exp(2z_l) - 1}{\exp(2z_l) + 1} \\ r_u = \tanh(z_u) = \frac{\exp(2z_u) - 1}{\exp(2z_u) + 1} \end{cases}$$

For our example:

95 percent confidence interval:

$$r_l = 0.207$$

$$r_u = 0.872$$

Nonparametric: Spearman's Rho

- Measure of monotone association used when the distribution of the data make Pearson's correlation coefficient undesirable or misleading
- Spearman's correlation coefficient (Rho) is defined as the Pearson's correlation coefficient between the ranked variables

- $$Rho = \frac{\sum_i (y_{1i} - \bar{y}_1)(y_{2i} - \bar{y}_2)}{\sqrt{\sum_i (y_{1i} - \bar{y}_1)^2 \sum_i (y_{2i} - \bar{y}_2)^2}}, \text{ where } y_{1i}, y_{2i} \text{ are ranks of } Y_{1i}, Y_{2i}$$

- $$\text{If no ties, } Rho = 1 - \frac{6 \sum d_i^2}{n(n^2 - 1)}, \text{ where } d_i = y_{1i} - y_{2i}$$

- Randomization test for significance (option)

Nonparametric: Kendall's Tau

- Concordant pairs (Y_{1i}, Y_{2i}) and (Y_{1j}, Y_{2j}) :

If $Y_{1i} > Y_{1j}$ and $Y_{2i} > Y_{2j}$ or if $Y_{1i} < Y_{1j}$ and $Y_{2i} < Y_{2j}$

(if the ranks for both elements agree)

- Discordant pairs (Y_{1i}, Y_{2i}) and (Y_{1j}, Y_{2j}) :

If $Y_{1i} > Y_{1j}$ and $Y_{2i} < Y_{2j}$ or if $Y_{1i} < Y_{1j}$ and $Y_{2i} > Y_{2j}$

(if the ranks for both elements disagree)

- Neither concordant or discordant

If $Y_{1i} = Y_{1j}$ or $Y_{2i} = Y_{2j}$

Nonparametric: Kendall's Tau

- Kendall's Tau =

$$\left\{ \begin{array}{l} \frac{n_c - n_d}{\frac{1}{2}n(n-1)} \quad \text{(no ties)} \\ \frac{n_c - n_d}{n_c + n_d} \quad \text{(in the case of ties)} \end{array} \right.$$

, where n_c = number of concordant pairs
 n_d = number of discordant pairs

Properties:

Gamma coefficient or Goodman correlation coefficient

- The denominator is the total number of pairs, $-1 \leq \text{tau} \leq 1$
- $\text{tau} = 1$, for perfect ranking agreement
- $\text{tau} = -1$, for perfect ranking disagreement
- $\text{tau} \approx 0$, if two variables are independent
- For large samples, the sampling distribution of tau is approximately normal

Nonparametric

For more information on nonparametric test of correlation
e.g., significance test, etc.

References:

- Conover, W.J. (1999) "Practical nonparametric statistics", 3rd ed. Wiley & Sons
- Kendall, M. (1948) "Rank Correlation Methods", Charles Griffin & Company Limited
- Caruso, J. C. & N. Cliff. (1997) "Empirical Size, Coverage, and Power of Confidence Intervals for Spearman's Rho", Ed. and Psy. Meas., 57 pp. 637-654
- Corder, G.W. & D.I. Foreman. (2009) "Nonparametric Statistics for Non-Statisticians: A Step-by-Step Approach", Wiley

Data

Total length of aphid stem mothers (Y_1)
Vs.

Mean thorax length of their parthenogenetic offspring (Y_2)

#	Y_1	Y_2
1	8.7	5.95
2	8.5	5.65
3	9.4	6.00
4	10.0	5.70
5	6.3	4.70
6	7.8	5.53
7	11.9	6.40
8	6.5	4.18
9	6.6	6.15
10	10.6	5.93
11	10.2	5.70
12	7.2	5.68
13	8.6	6.13
14	11.1	6.30
15	11.6	6.03

Total length of mothers vs. Mean thorax length of offspring

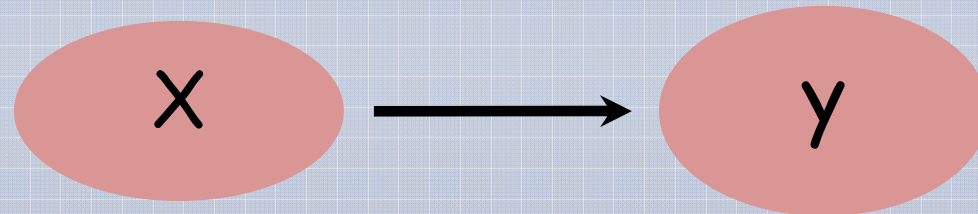
#	RAW		RANK	
	Y_1	Y_2	y_1	y_2
1	8.7	5.95	8	9
2	8.5	5.65	6	4
3	9.4	6.00	9	10
4	10.0	5.70	10	6.5
5	6.3	4.70	1	2
6	7.8	5.53	5	3
7	11.9	6.40	15	15
8	6.5	4.18	2	1
9	6.6	6.15	3	13
10	10.6	5.93	12	8
11	10.2	5.70	11	6.5
12	7.2	5.68	4	5
13	8.6	6.13	7	12
14	11.1	6.30	13	14
15	11.6	6.03	14	11

Group Activity

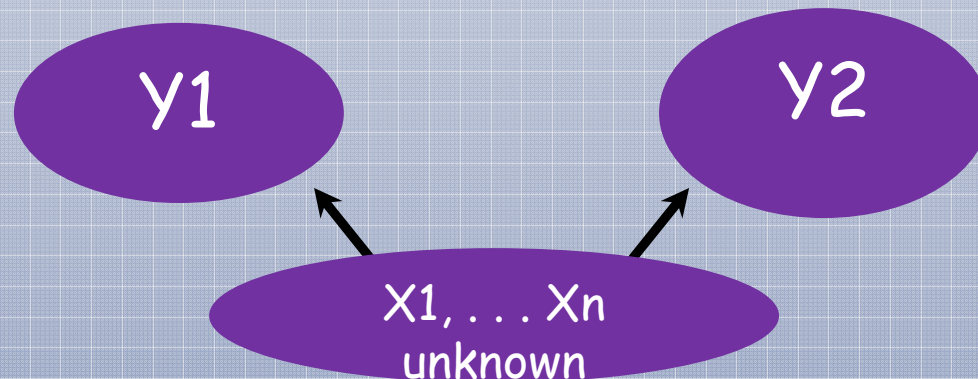
Activity Instructions

- Question: **REGRESSION** or **CORRELATION**?
- Justification guideline:

Regression:



Correlation:



Activity Instructions

- Form small groups or 2-3 people.
- Each group is assigned a number
- Group members work together on each example for 5 minutes, come up with an answer & your justifications
- A number will be randomly generated from the group #'s
- The corresponding group will have to present their answer & justifications
- Go for the next example . . .

Activity Instructions

There is NO RIGHT/WRONG

ANSWER (for these examples),

as long as your justifications are

LOGICAL

Example 1

Height and ratings of physical attractiveness vary across individuals. Would you analyze this as regression or correlation?

Subject	Height	Phy
1	69	7
2	61	8
3	68	6
4	66	5
5	66	8
.	..	.
48	71	10

Example 2

Airborne particles such as dust and smoke are an important part of air pollution. Measurements of airborne particles made every six days in the center of a small city and at a rural location 10 miles southwest of the city

(Moore & McCabe, 1999. Introduction to the Practice of Statistics).

Would you analyze this relation as regression or correlation?

Example 3

A study conducted in the Egyptian village of Kalama examined the relation between birth weights of 40 infants and family monthly income

(El-Kholy et al. 1986, Journal of the Egyptian Public Health Association, 61: 349).

Would you analyze this relation as regression or correlation?