# Multivariate Statistics in Ecology and Quantitative Genetics Hotellings T<sup>2</sup>

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http://evol.bio.lmu.de/\_statgen

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- Multivariate normal distribution
- 4 The multivariate normal distribution in R

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- 5 Hotellings T<sup>2</sup>-test
- Testing for normality

We want to compare the vegetative growth of a mutated raspberry (*Rubus idaeus*) with the wildtype raspberry.



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We want to compare the vegetative growth of a mutated raspberry (*Rubus idaeus*) with the wildtype raspberry.

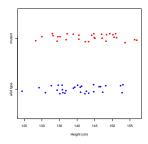


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We use different quantities as a measure for "growth":

- Height of bush (cm)
- Width of bush (cm)

#### Comparison of height:



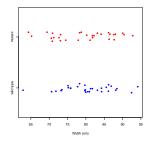
```
> raspberry <- read.table("raspberry.csv",sep=",",h=T)
> attach(raspberry)
> height.wt <- height[type=="wild type"]
> height.mu <- height[type=="mutant"]
> t.test(height.wt,height.mu)
```

```
Welch Two Sample t-test
```

```
data: height.wt and height.mu
t = -1.5441, df = 57.334, p-value = 0.1281
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
   -6.7627773 0.8735087
sample estimates:
mean of x mean of y
140.2768 143.2215
```

#### Motivating example

#### Comparison of width:



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```
> width.wt <- width [type=="wild type"]
> width.mu <- width [type=="mutant"]
> t.test(width.wt,width.mu)
```

```
Welch Two Sample t-test
```

data: width.wt and width.mu
t = 0.5717, df = 56.645, p-value = 0.5698
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -2.709472 4.874489
sample estimates:
mean of x mean of y
80.47013 79.38762

Both height and width are not significantly different for the the types of raspberry.

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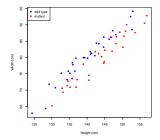
Is there no difference?



Both height and width are not significantly different for the the types of raspberry.

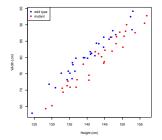
#### Is there no difference?

So far, we have not exploited the correlation between height and width. Let's look at the bivariate (two variables) data:



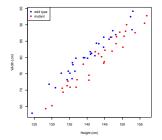
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We see a difference "by eye". Which test can detect that?





We see a difference "by eye". Which test can detect that? Answer: We need the multivariate analogon of the t-test. This multivariate version of the t-test is called

#### Hotelling's T<sup>2</sup>-test

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(Before we come to this test, we need to learn some theory)

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• The vector (height,width) is a two-dimensional random vector (with values in  $\mathbb{R}^2)$ 

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• An *d*-dimensional random vector is a vector of *d* random elements

- The vector (height,width) is a two-dimensional random vector (with values in  $\mathbb{R}^2$ )
- An *d*-dimensional random vector is a vector of *d* random elements
- The expectation of a random vector X = (X<sub>1</sub>, X<sub>2</sub>, ..., X<sub>d</sub>)<sup>T</sup> is the vector of the expectations:

$$\mathbb{E}X = \mathbb{E}\begin{pmatrix} X_1 \\ \vdots \\ X_d \end{pmatrix} = \begin{pmatrix} \mathbb{E}X_1 \\ \vdots \\ \mathbb{E}X_d \end{pmatrix}$$

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The expectation of a random matrix M = (M<sub>ij</sub>)<sub>i=1..n,j=1..d</sub> is the matrix of the expectations:

$$\mathbb{E}\begin{pmatrix} M_{11} & M_{12} & \cdots & M_{1d} \\ \vdots & \ddots & \vdots \\ M_{n1} & M_{n2} & \cdots & M_{nd} \end{pmatrix} = \begin{pmatrix} \mathbb{E}M_{11} & \mathbb{E}M_{12} & \cdots & \mathbb{E}M_{1d} \\ \vdots & \ddots & & \vdots \\ \mathbb{E}M_{n1} & \mathbb{E}M_{n2} & \cdots & \mathbb{E}M_{nd} \end{pmatrix}$$

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• reminder: The variance of a univariate random variable X is  $Var(X) = \mathbb{E}\left[(X - \mathbb{E}X)^2\right] = \mathbb{E}\left[X^2\right] - (\mathbb{E}X)^2$ .

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- reminder: The variance of a univariate random variable X is  $Var(X) = \mathbb{E}\left[(X \mathbb{E}X)^2\right] = \mathbb{E}\left[X^2\right] (\mathbb{E}X)^2$ .
- The analog in the multivariate case is the so called *covariance* matrix (or dispersion matrix or variance-covariance matrix). The covariance matrix Var(X) = Σ of X = (X<sub>1</sub>,...,X<sub>d</sub>)<sup>T</sup> is

$$\Sigma = \begin{pmatrix} \operatorname{Cov}(X_1, X_1) & \operatorname{Cov}(X_1, X_2) & \cdots & \operatorname{Cov}(X_1, X_d) \\ \operatorname{Cov}(X_2, X_1) & \operatorname{Cov}(X_2, X_2) & \cdots & \operatorname{Cov}(X_2, X_d) \\ \vdots & & \ddots & \vdots \\ \operatorname{Cov}(X_d, X_1) & \operatorname{Cov}(X_d, X_2) & \cdots & \operatorname{Cov}(X_d, X_d) \end{pmatrix} \\ = \mathbb{E} \left[ \begin{pmatrix} X_1 - \mathbb{E}X_1 \\ \vdots \\ X_d - \mathbb{E}X_d \end{pmatrix} \cdot \begin{pmatrix} X_1 - \mathbb{E}X_1, \cdots, X_d - \mathbb{E}X_d \end{pmatrix} \right] \\ = \mathbb{E} \left[ (X - \mathbb{E}X) \cdot (X - \mathbb{E}X)^T \right] \\ = \mathbb{E} \left[ X \cdot X^T \right] - \mathbb{E}X \cdot (\mathbb{E}X)^T$$

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Linearity of the expectation is analogos to the univarite case: Let X = (X<sub>1</sub>,..., X<sub>d</sub>) be a random vector and C = (C<sub>ij</sub>)<sub>i=1..n,j=1..d</sub> be a deterministic matrix. Then

$$\mathbb{E}(C \cdot X) = C \cdot \mathbb{E}(X)$$

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$$\mathbb{E}(C \cdot X) = C \cdot \mathbb{E}(X)$$

• If  $Y := X - \mathbb{E}(X)$ , then

$$Var(C \cdot X) = Var(C \cdot Y)$$
$$= \mathbb{E} [C \cdot Y \cdot (C \cdot Y)^{T}]$$
$$= \mathbb{E} [C \cdot Y \cdot Y^{T} \cdot C^{T}]$$
$$= C \cdot \mathbb{E} [Y \cdot Y^{T}] \cdot C^{T}$$
$$= C \cdot Var(Y) \cdot C^{T}$$
$$= C \cdot Var(X) \cdot C^{T}$$

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#### Motivating example

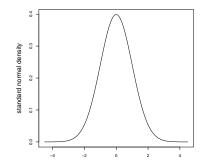
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Reminder: Univariate normal distribution N(μ, σ<sup>2</sup>) with mean μ ∈ ℝ and variance σ<sup>2</sup> ∈ (0, ∞) has the density

$$f(x) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(x-\mu)^2}{2\sigma^2}\right)$$



Remember:  $\mathbb{P}(\mu - \sigma < X < \mu + \sigma) = 0.68$  and  $\mathbb{P}(\mu - 1.96\sigma < X < \mu + 1.96\sigma) = 0.95$ 

• The density of the *d*-dimensional normal distribution with mean  $\mu \in \mathbb{R}^d$  and covariance matrix  $\Sigma \in \mathbb{R}^{d \times d}$  is analogous:

$$f(x) = \frac{1}{\sqrt{(2\pi)^d \det(\Sigma)}} \exp\left(-\frac{(x-\mu)^T \Sigma^{-1} (x-\mu)}{2}\right)$$

for  $x \in \mathbb{R}^d$  where det $(\Sigma)$  is the determinand of  $\Sigma$ , and  $\Sigma^{-1}$  is the inverse matrix.

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We write  $\mathcal{N}_{d}(\mu, \Sigma)$  for this distribution.

 The density of the *d*-dimensional normal distribution with mean μ ∈ ℝ<sup>d</sup> and covariance matrix Σ ∈ ℝ<sup>d×d</sup> is analogous:

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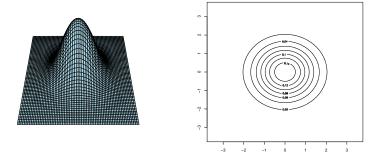
for  $x \in \mathbb{R}^d$  where det( $\Sigma$ ) is the determinand of  $\Sigma$ , and  $\Sigma^{-1}$  is the inverse matrix. We write  $\mathcal{N}_d(\mu, \Sigma)$  for this distribution.

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 The standard multivariate normal distribution has mean μ = 0 and the identity matrix Σ = 1 as covariance matrix.

Correlation 0.0:  

$$\Sigma = \begin{pmatrix} 1 & 0 \\ 0 & 1 \end{pmatrix}$$
,  $Var(X_1) = 1 = Var(X_2)$ ,  $Cov(X_1, X_2) = 0.0$ 

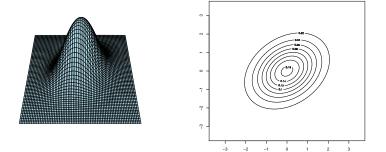


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Correlation 0.3:  

$$\Sigma = \begin{pmatrix} 1 & 0.3 \\ 0.3 & 1 \end{pmatrix}$$
,  $Var(X_1) = 1 = Var(X_2)$ ,  $Cov(X_1, X_2) = 0.3$ 



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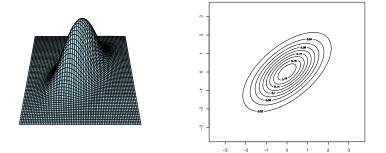
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Correlation 0.6:  

$$\Sigma = \begin{pmatrix} 1 & 0.6 \\ 0.6 & 1 \end{pmatrix}$$
,  $Var(X_1) = 1 = Var(X_2)$ ,  $Cov(X_1, X_2) = 0.6$ 



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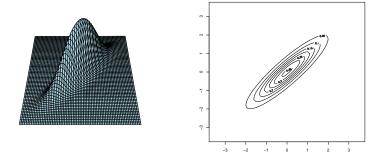
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Correlation 0.9:  

$$\Sigma = \begin{pmatrix} 1 & 0.9 \\ 0.9 & 1 \end{pmatrix}$$
,  $Var(X_1) = 1 = Var(X_2)$ ,  $Cov(X_1, X_2) = 0.9$ 



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Multivariate normal distribution

Properties: (Let the distribution of *X* be  $\mathcal{N}_d(\mu, \Sigma)$ )

Linear combinations are univariate normal distributed:
 ⟨c, X⟩ ~ N (⟨c, μ⟩, cΣc<sup>T</sup>)

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•  $X_i$  and  $X_j$  are independent  $\iff \text{Cov}(X_i, X_j) = 0$ 

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   ⟨c, X⟩ ~ N (⟨c, μ⟩, cΣc<sup>T</sup>)
- $X_i$  and  $X_j$  are independent  $\iff \text{Cov}(X_i, X_j) = 0$
- The standardized normal distribution is standard normal distributed

$$\Sigma^{-\frac{1}{2}} \cdot (\boldsymbol{X} - \mu) \sim \mathcal{N}_{\boldsymbol{d}}(\boldsymbol{0}, \mathbb{1})$$

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where  $M = \Sigma^{-\frac{1}{2}}$  is a matrix such that  $M^T \cdot M \cdot \Sigma = \mathbb{1}$ .

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- $X_i$  and  $X_j$  are independent  $\iff \text{Cov}(X_i, X_j) = 0$
- The standardized normal distribution is standard normal distributed

$$\Sigma^{-\frac{1}{2}} \cdot (X - \mu) \sim \mathcal{N}_d(0, \mathbb{1})$$

where  $M = \Sigma^{-\frac{1}{2}}$  is a matrix such that  $M^T \cdot M \cdot \Sigma = \mathbb{1}$ .

• The square of the standardized normal distribution is chi-squared distributed with *d* degrees of freedom:

$$(\boldsymbol{X} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1} (\boldsymbol{X} - \boldsymbol{\mu}) \sim \chi_d^2.$$

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- Linear combinations are univariate normal distributed:
   ⟨c, X⟩ ~ N (⟨c, μ⟩, cΣc<sup>T</sup>)
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• The square of the standardized normal distribution is chi-squared distributed with *d* degrees of freedom:

$$(\boldsymbol{X} - \boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1} (\boldsymbol{X} - \boldsymbol{\mu}) \sim \chi_d^2.$$

If Y<sub>1</sub>, Y<sub>2</sub>,..., Y<sub>d</sub> are independent and standard normal distributed, then (Y<sub>1</sub>,..., Y<sub>d</sub>) ~ N(0, 1).

• If  $M \in \mathbb{R}^{p \times d}$  is a non-random matrix, then  $M \cdot X \sim \mathcal{N}_p(M \cdot \mu, M \Sigma M^T)$ 

### Estimating $\mu$ and $\Sigma$

Let  $Y_1, \ldots, Y_n$  be a sample of independent observations from a  $\mathcal{N}_d(\mu, \Sigma)$  distribution. As usual, we write the variables as columns and the different observations as rows:

$$Y = \begin{pmatrix} Y_{11} & Y_{12} & \cdots & Y_{1d} \\ Y_{21} & Y_{22} & \cdots & Y_{2d} \\ \vdots & & \ddots & \vdots \\ Y_{n1} & Y_{n2} & \cdots & Y_{nd} \end{pmatrix}$$

• The sample mean  $\bar{Y}$  where  $\bar{Y}_j := \frac{1}{n} \sum_{i=1}^n Y_{ij}$  is an estimator for  $\mu$ .

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- The sample mean  $\overline{Y}$  where  $\overline{Y}_j := \frac{1}{n} \sum_{i=1}^n Y_{ij}$  is an estimator for  $\mu$ .
- The sample covariance matrix  $S := (S_{ij})_{i,j=1..d}$  where

$$S_{ij} = \frac{1}{n-1} \sum_{k=1}^{n} \left( Y_{ki} - \bar{Y}_i \right) \left( Y_{kj} - \bar{Y}_j \right)$$

is an (unbiased) estimator for  $\Sigma$ .

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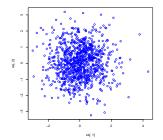
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- > library("mvtnorm") # multivariate t- and normal distrib.
- > Sigma0 <- matrix(c(1,0,0,1),ncol=2)</pre>
- > dmvnorm(c(0,0),mean=c(0,0),sigma=Sigma0)
- [1] 0.1591549
- > dmvnorm(c(0,0)) # same, the standard normal is the default
  [1] 0.1591549

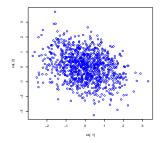
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- [1] 0.1591549
- > dmvnorm(c(0,0)) # same, the standard normal is the default
  [1] 0.1591549
- > xx <- rmvnorm(1000,mean=c(0,0),sigma=Sigma0)</pre>
- > plot(xx[,1],xx[,2])



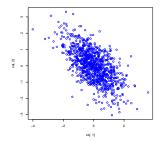
Correlation -0.3:

- > Sigma3 <- matrix(c(1,-0.3,-0.3,1),ncol=2)</pre>
- > xx <- rmvnorm(1000,mean=c(0,0),sigma=Sigma3)</pre>
- > plot(xx[,1],xx[,2])



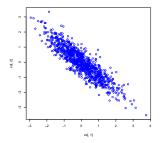
Correlation -0.6:

- > Sigma6 <- matrix(c(1,-0.6,-0.6,1),ncol=2)</pre>
- > xx <- rmvnorm(1000,mean=c(0,0),sigma=Sigma6)</pre>
- > plot(xx[,1],xx[,2])



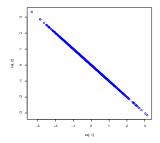
Correlation -0.9:

- > Sigma9 <- matrix(c(1,-0.9,-0.9,1),ncol=2)</pre>
- > xx <- rmvnorm(1000,mean=c(0,0),sigma=Sigma9)</pre>
- > plot(xx[,1],xx[,2])



Correlation -1:

- > Sigma10 <- matrix(c(1,-1,-1,1),ncol=2)</pre>
- > xx <- rmvnorm(1000,mean=c(0,0),sigma=Sigma10)</pre>
- > plot(xx[,1],xx[,2])



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Reminder for univarite case d = 1: If X = (X<sub>1</sub>,..., X<sub>n</sub>) is an independent sample from the N<sub>1</sub>(μ, σ<sup>2</sup>) distribution, then the t-statistics

$$t := \frac{\bar{X} - \mu}{sd(X)/\sqrt{n}}$$

is *t*-distributed with n - 1 degrees of freedom. This fact is used for the t-test.

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$$t := \frac{\bar{X} - \mu}{sd(X)/\sqrt{n}}$$

is *t*-distributed with n - 1 degrees of freedom. This fact is used for the t-test. In addition,  $t^2 \sim \mathcal{F}_{1,n-1}$  is Fisher-distributed with 1 numerator degree of freedom and n - 1 denominator degrees of freedom.

 Multivariate case: If X := (X<sub>1</sub>,..., X<sub>n</sub>, be a sample from the *N<sub>d</sub>* (μ, Σ) distribution with n > d, then the statistics

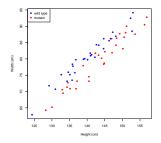
$$T^2 := n \left( \bar{X} - \mu \right)^T S^{-1} \left( \bar{X} - \mu \right)$$

is called *Hotellings*  $T^2$ . The matrix  $S^{-1}$  is the inverse of the sample covariance matrix. It has been shown that

$$T^2 \sim \frac{(n-1)d}{n-d} \mathcal{F}_{d,n-d}$$

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Back to our raspberry example:



We see a difference "by eye". Does Hotellings  $T^2$ -test detect this difference?

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- > raspberry <- read.table("raspberry.csv",sep=",",header=T)</pre>
- > raspberry.wt<-subset(raspberry,type=="wild type",select=-3)</pre>
- > raspberry.mu<-subset(raspberry,type=="mutant",select=-3)</pre>
- > library(rrcov) #one out of many libraries with Hotelling T2
- > T2.test(raspberry.wt,raspberry.mu)

```
Hotelling's two sample T2-test
```

Hotellings T<sup>2</sup>-test detects a significant difference

- # alternative syntax with a formula
- > T2.test(cbind(height,width)~type,data=raspberry)

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- # alternative syntax with a formula
- > T2.test(cbind(height,width)~type,data=raspberry)

Syntax:

## Default S3 method: T2.test(x, y = NULL, mu = 0, conf.level = 0.95, ...)

## S3 method for class 'formula': T2.test(formula, data, subset, na.action, ...)

If y = NULL, then perform a one sample T<sup>2</sup>-test with null hypothesis  $H_0$ : true mean is mu

 The T<sup>2</sup>-test (like the t-test) is prone to outliers. A single outlier can considerably decrease the power of the test. So check for outliers (possible mismeasurements)

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- The T<sup>2</sup>-test (more than the t-test) looses test power if the true distribution (of the population) deViates from the normal distribution (e.g., the Shapiro-Wilk test tests for normality)
- If one of the variables shows already a significant difference between the group, then also the T<sup>2</sup>-test yields a significant difference.

# Contents

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- 3 Multivariate normal distribution
- 4 The multivariate normal distribution in R

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- 5 Hotellings T<sup>2</sup>-test
- Testing for normality

#### Testing for normality

A multivariate Shapiro-Wilk test for normality is mshapiro.test() in the library mvnormtest:

# install.packages("mvnormtest") # install library if

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- > library(mvnormtest)
- > library(mvtnorm)
- > x <- rmvnorm(100,mean=c(0,0))
- > mshapiro.test(t(x))

```
Shapiro-Wilk normality test
```

```
data: Z
W = 0.9915, p-value = 0.7839
```