

# *Single Factor Experiments*

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- Topic:
  - Comparison of more than 2 groups
  - One-Way Analysis of Variance
  - F test
- Learning Aims:
  - Understand model parametrization
  - Carry out an anova
- Reason: Multiple t tests won't do!

# Potatoe scab

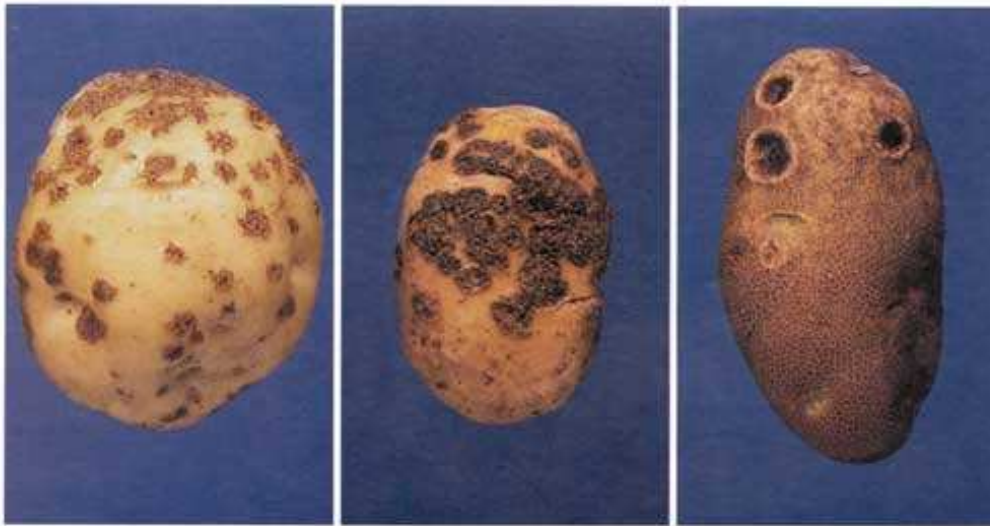


Figure 1.

Figure 2.

Figure 3.

- widespread disease
- causes economic loss
- known factors: variety, soil condition

# *Experiment with different treatments*

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- Compare 7 treatments for effectiveness in reducing scab
- Field with 32 plots, 100 potatoes are randomly sampled from each plot
- For each potatoe the percentage of the surface area affected was recorded. Response variable is the average of the 100 percentages.

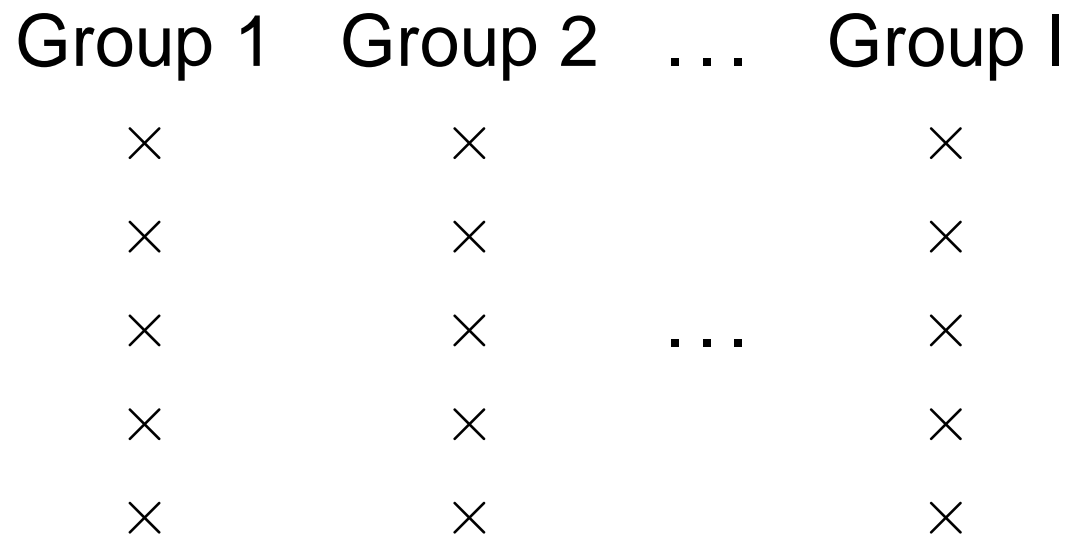
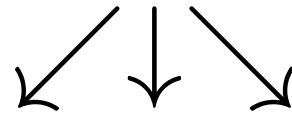
## *Field plan and data*

|    |    |    |    |    |    |    |    |
|----|----|----|----|----|----|----|----|
| 2  | 1  | 6  | 4  | 6  | 7  | 5  | 3  |
| 9  | 12 | 18 | 10 | 24 | 17 | 30 | 16 |
| 1  | 5  | 4  | 3  | 5  | 1  | 1  | 6  |
| 10 | 7  | 4  | 10 | 21 | 24 | 29 | 12 |
| 2  | 7  | 3  | 1  | 3  | 7  | 2  | 4  |
| 9  | 7  | 18 | 30 | 18 | 16 | 16 | 4  |
| 5  | 1  | 7  | 6  | 1  | 4  | 1  | 2  |
| 9  | 18 | 17 | 19 | 32 | 5  | 26 | 4  |

# 1-Factor Design

Plots, subjects

Randomisation



# *Complete Randomisation*

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- a) number the plots 1, ..., 32.
- b) construct a vector with 8 replicates of 1 and 4 replicates of 2 to 7.
- c) choose a random permutation and apply it to the vector in b).

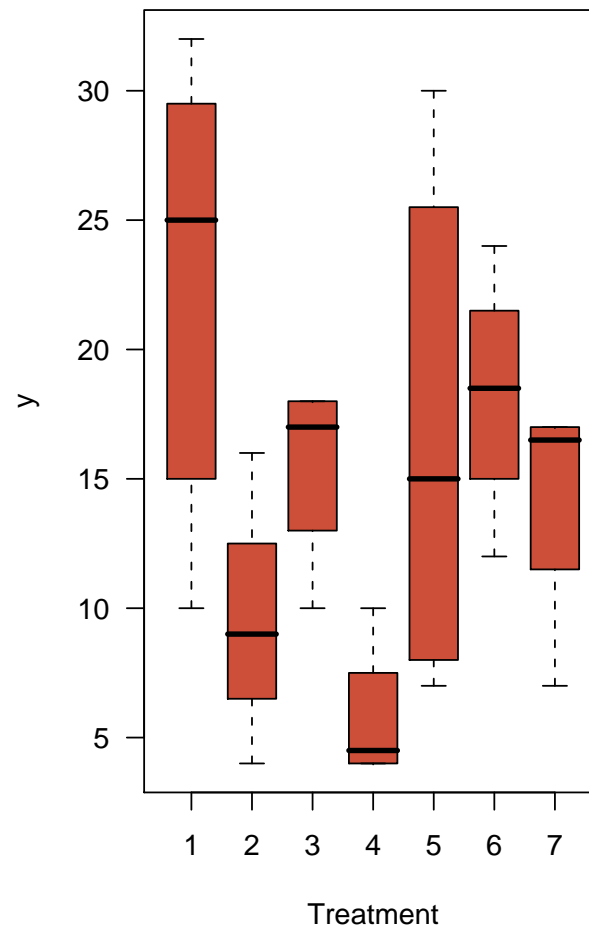
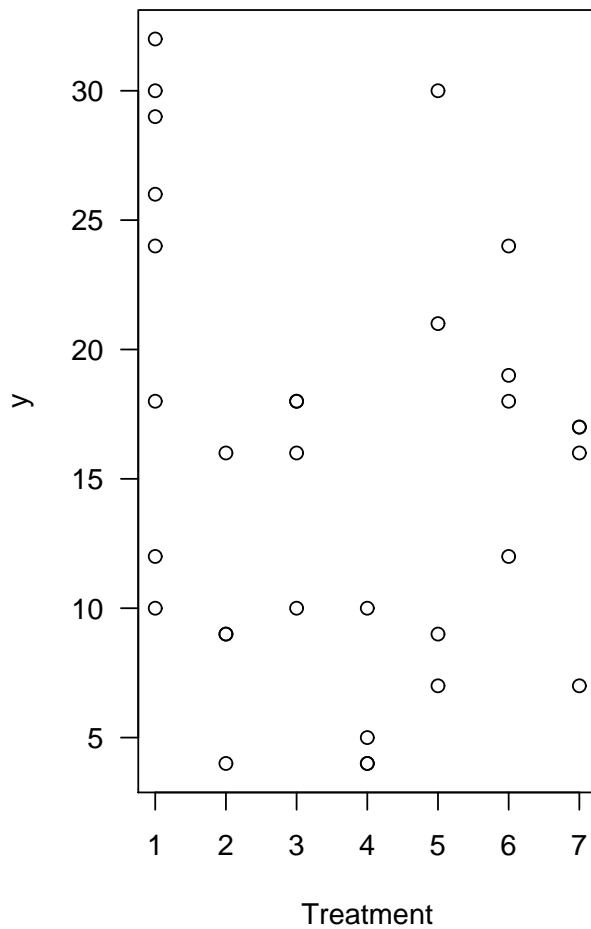
in R:

```
> treatment=factor(c(rep(1,8),rep(2:7,rep(4,6))))  
> sample(treatment)
```

# *Exploratory data analysis*

| Group | y  |    |    |    |    | $\bar{y}$ |    |    |        |
|-------|----|----|----|----|----|-----------|----|----|--------|
| 1     | 12 | 10 | 24 | 29 | 30 | 18        | 32 | 26 | 22.625 |
| 2     | 9  | 9  | 16 | 4  |    |           |    |    | 9.5    |
| 3     | 16 | 10 | 18 | 18 |    |           |    |    | 15.5   |
| 4     | 10 | 4  | 4  | 5  |    |           |    |    | 5.75   |
| 5     | 30 | 7  | 21 | 9  |    |           |    |    | 16.75  |
| 6     | 18 | 24 | 12 | 19 |    |           |    |    | 18.25  |
| 7     | 17 | 7  | 16 | 17 |    |           |    |    | 14.25  |

# Graphical display





# Two sample *t* tests

|         |   |         |   |                       |
|---------|---|---------|---|-----------------------|
| Group 1 | – | Group 2 | : | $H_0 : \mu_1 = \mu_2$ |
| Group 1 | – | Group 3 | : | $H_0 : \mu_1 = \mu_3$ |
| Group 1 | – | Group 4 | : | $H_0 : \mu_1 = \mu_4$ |
| Group 1 | – | Group 5 | : | $H_0 : \mu_1 = \mu_5$ |
| Group 1 | – | Group 6 | : | $H_0 : \mu_1 = \mu_6$ |
| Group 1 | – | Group 7 | : | $H_0 : \mu_1 = \mu_7$ |

...

$\alpha = 5\%$ ,  $P(\text{Test not significant} | H_0) = 95\%$

7 groups, 21 independent tests:

$P(\text{none of the tests sign.} | H_0) = 0.95^{21} = 0.34$

$P(\text{at least one test sign.} | H_0) = 0.66$  (more realistic: 0.42)

$$1 - (1 - \alpha)^n$$

# ***Bonferroni correction***

Choose  $\alpha_T$  such that

$$1 - (1 - \alpha_T)^n = \alpha_E = 5\%$$

( $\alpha_T = \alpha$  „testwise“,  $\alpha_E = \alpha$  „experimentwise“)

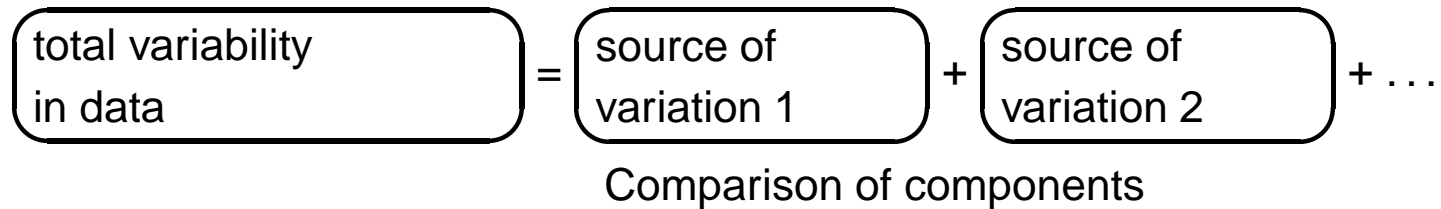
Since  $1 - (1 - \frac{\alpha}{n})^n \approx \alpha$ , the significance level for a single test has to be divided by the number of tests.

Overcorrection, not very efficient.

# Analysis of variance

- Comparison of more than 2 groups
- for more complex designs
- global F test

## Idea:



total = treatment + experimental error

total = variability of plots with different treatments + variability of plots with the same treatment

$\sigma^2 + \text{treatment effect}$

$\sigma^2$

# Definitions

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- **Factor**: categorical, explanatory variable  
**Level**: value of a factor  
Ex 1: Factor= soil treatment, 7 levels 1 – 7.  
⇒ One-way analysis of variance  
Ex 2: 3 varieties with 4 quantities of fertilizer  
⇒ Two-way analysis of variance
- **Treatment**: combination of factors
- **Plot, experimental unit**: smallest unit to which a treatment can be applied  
Ex: feeding (chicken, chicken-houses), dental medicine (families, people, teeth)

# One-way analysis of variance

Model:

response = treatment + error (Plot)

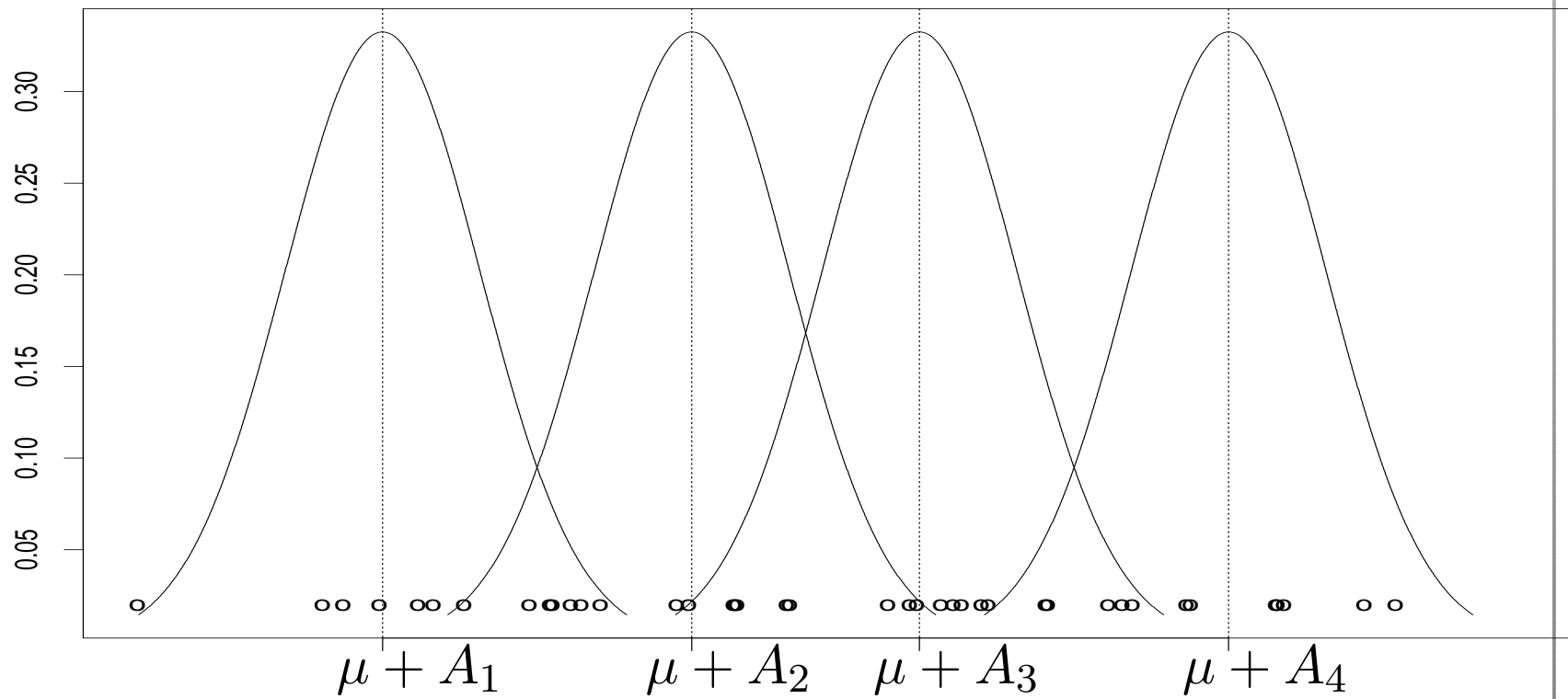
$$(1) \quad Y_{ij} = \mu + A_i + \epsilon_{ij}$$
$$i=1, \dots, I; j=1, \dots, J_i$$

$\mu$  = overall mean

$A_i$  =  $i$ th treatment effect

$\epsilon_{ij}$  = random error,  $\mathcal{N}(0, \sigma^2)$  iid.

# Illustration of model (1)



# Necessary constraint

Group mean

| $\mu + A_i$ | $\mu + A_i$ | $\mu + A_i$ |
|-------------|-------------|-------------|
| 4           | 4+0         | 3+1         |
| 5           | 4+1         | 3+2         |
| 3           | 4-1         | 3+0         |

usual constraint:  $\sum J_i A_i = 0$ ,  
 $\sum A_i = 0$  if  $J_i = J$  for all  $i$   
( $A_i$  deviation from overall mean)

or  $A_1 = 0$ , resp.  $A_I = 0$

# ***Decomposition of the deviation of a response from the overall mean***

$$y_{ij} - y_{..} = \underbrace{y_{i.} - y_{..}}_{\text{deviation of the group mean}} + \underbrace{y_{ij} - y_{i.}}_{\text{deviation from the group mean}}$$

$$y_{i.} = \frac{1}{J_i} \sum_j y_{ij} \text{ mean of group } i,$$

$$y_{..} = \frac{1}{N} \sum_i \sum_j y_{ij} \text{ overall mean, } N = \sum J_i.$$



# Analysis of variance identity

$$\underbrace{\sum_i \sum_j (y_{ij} - y_{..})^2}_{\text{total variability}} = \underbrace{\sum_i \sum_j (y_{i.} - y_{..})^2}_{\text{variability between groups}} + \underbrace{\sum_i \sum_j (y_{ij} - y_{i.})^2}_{\text{variability within groups}}$$

total sum of squares = treatment sum of squares + residual sum of squares

$$SS_{tot} = SS_{treat} + SS_{res}$$

# Mean squares

Total mean square:  $MS_{tot} = SS_{tot}/(N - 1)$

Residual mean square:  $MS_{res} = SS_{res}/(N - I)$

$$\frac{SS_{res}}{N - I} = \frac{\sum_i (J_i - 1) S_i^2}{\sum_i (J_i - 1)}, \quad S_i^2 = \frac{\sum_j (y_{ij} - y_{i.})^2}{J_i - 1}$$

$$MS_{res} = \hat{\sigma}^2 = \widehat{Var}(Y_{ij}), \quad E(MS_{res}) = \sigma^2$$

Treatment mean square:  $MS_{treat} = SS_{treat}/(I - 1)$

$$E(MS_{treat}) = \sigma^2 + \sum J_i A_i^2 / (I - 1)$$

$$df_{tot} = df_{treat} + df_{res}, \quad N - 1 = I - 1 + N - I$$

## ***R: anova table***

```
> mod1=aov(y~treatment,data=scab)
```

```
> summary(mod1)
```

|           | Df | Sum Sq  | Mean Sq | F value | Pr(>F) |   |
|-----------|----|---------|---------|---------|--------|---|
| treatment | 6  | 972.34  | 162.06  | 3.608   | 0.0103 | * |
| Residuals | 25 | 1122.88 | 44.92   |         |        |   |

$$F = \frac{MS_{treat}}{MS_{res}} \approx 1 \implies \text{no treatment differences}$$

$$F = \frac{MS_{treat}}{MS_{res}} \gg 1 \implies \text{treatment differences exist}$$

# ***F* test**

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$H_0$  : all  $A_i = 0$

$H_A$  : at least one  $A_i \neq 0$

Since  $\epsilon_{ij} \sim \mathcal{N}(0, \sigma^2)$ ,  $F = \frac{MS_{treat}}{MS_{res}}$  has under  $H_0$  an  $F$  distribution with  $I - 1$  and  $N - I$  degrees of freedom.

one-sided test:

reject  $H_0$  if  $F > F_{95\%, I-1, N-I}$

# Chisquare and $t$ distribution

- Let  $Z_1, \dots, Z_n \sim \mathcal{N}(0, 1)$ , *iid*. Then

$$X = Z_1^2 + Z_2^2 + \dots + Z_n^2$$

has a  $\chi^2$  distribution with  $n$  df,  $X \sim \chi_n^2$

- Let  $Z \sim \mathcal{N}(0, 1)$  and  $X \sim \chi_n^2$  be independent random variables. The distribution of

$$T = \frac{Z}{\sqrt{X/n}}$$

is called the  $t$  distribution with  $n$  df,  $T \sim t_n$

# *F distribution*

- Let  $X_1 \sim \chi_n^2$  and  $X_2 \sim \chi_m^2$  be independent random variables. The distribution of

$$F = \frac{X_1/n}{X_2/m}$$

is called the  $F$  distribution with  $n$  and  $m$  df,

$$F \sim F_{n,m}$$

Properties:

$$F_{1,m} = t_m^2$$
$$E(F_{n,m}) = \frac{m}{m-2}$$