● Intro: Small example and TV-data from B&O

Model

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Omputation - decomposition and the ANOVA table

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Post hoc analysis

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A complete example - from the book

Intro: Small example and TV-data from B&O

### TV set development at Bang & Olufsen

Sound and image quality is measured by th human perceptual instrument:



We developed a tool that is used by B&O to ANOVA (among other things) PanelCheck (Show Panelcheck programme with TV data)

### Course 02402 Introduction to Statistics Lecture 11:

### Twoway Analysis of Variance, ANOVA

### Per Bruun Brockhoff

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Intro: Small example and TV-data from B&O

### Agenda

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### Bang & Olufsen data in R:

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Intro: Small example and TV-data from B&O

### Twoway ANOVA - example

 Same data as for oneway, but now we know that the experiment was split in blocks

	Group A	Group B	Group C
Block 1	2.8	5.5	5.8
Block 2	3.6	6.3	8.3
Block 3	3.4	6.1	6.9
Block 4	2.3	5.7	6.1

- hence three *Groups* on four *blocks*
- or three treatments on four persons
- or three varieties on four fields (hence blocks)
- or similarly
- oneway vs. twoway ANOVA
- Completely randomized design vs. Randomized block design

### Bang & Olufsen data in R:

	TV3	TV2	TV1
Person 1	9.30	4.70	6.60
Person 2	10.20	7.00	8.80
Person 3	11.50	9.50	8.00
Person 4	11.90	6.60	8.20
Person 5	10.70	4.20	5.40
Person 6	10.90	9.10	7.10
Person 7	8.50	5.00	6.30
Person 8	12.60	8.90	10.70

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Intro: Small example and TV-data from B&O

### Twoway ANOVA - example

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	Group A	Group B	Group C	
Block 1	2.8	5.5	5.8	
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Block 4	2.3	5.7	6.1	

- Question: Is there a significant difference (in means) between the groups A, B and C?
- ANOVA can be used if the observations in each group are (approximately) normally distributed OR if  $n_i$ s are large enough (CLT)

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### The toy data in R

```
## Input data and plot
## Observations
y \leftarrow c(2.8, 3.6, 3.4, 2.3,
      5.5, 6.3, 6.1, 5.7,
      5.8, 8.3, 6.9, 6.1)
## treatments (Groups, varieties)
treatm <- factor(c(1, 1, 1, 1,
                  2, 2, 2, 2,
                  3, 3, 3, 3))
## blocks (persons, fields)
block <- factor(c(1, 2, 3, 4,
                 1, 2, 3, 4,
                 1, 2, 3, 4))
## for later formulas
(k <- length(unique(treatm)))</pre>
(1 <- length(unique(block)))</pre>
## Plots
par(mfrow=c(1,2))
## Plot histogramms by treatments
plot(treatm, y, xlab="Treatments", ylab="y")
## Plot histogrammer by blocks
plot(block, y, xlab="Blocks", ylab="y")
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```

Mode

### Twoway ANOVA, model

Express a model

$$Y_{ij} = \mu + \alpha_i + \beta_j + \varepsilon_{ij}, \quad \varepsilon_{ij} \sim N(0, \sigma^2)$$

where the deviations

$$arepsilon_{ij} \sim N(0, oldsymbol{\sigma}^2)$$
 and i.i.d.

- $\bullet$   $\mu$  is the overall mean
- $\alpha_i$  is the effect of treatment i
- $\beta_i$  is the level for Block i
- ullet there are k treatments and l blocks
- j indicates the observations in the groups, from 1 to  $n_i$  for treatment i

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Мо

### Estimates of parameters in the model

ullet We can compute the estimates of the parameters  $(\hat{\mu}$  and  $\hat{lpha}_i$ , and  $\hat{eta}_j)$ 

$$\hat{\mu} = \bar{y} = \frac{1}{k \cdot l} \sum_{i=1}^{k} \sum_{j=1}^{l} y_{ij}$$

$$\hat{\alpha}_i = \left(\frac{1}{l} \sum_{j=1}^l y_{ij}\right) - \hat{\mu}$$

$$\hat{\beta}_j = \left(\frac{1}{k} \sum_{i=1}^k y_{ij}\right) - \hat{\mu}$$

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Computation - decomposition and the ANOVA table

### Formulas for sums of squares

• Total sum of squares ("the total variance") (same as for oneway)

$$SST = \sum_{i=1}^{k} \sum_{j=1}^{l} (y_{ij} - \hat{\mu})^2$$

• treatment sum of squares ("Variance explained by the treatment part of the model")

$$SS(Tr) = l \cdot \sum_{i=1}^{k} \hat{\alpha}_i^2$$

## Twoway ANOVA, decomposition and the ANOVA table, Theorem 8.20

With the model

$$Y_{ij} = \mu + \alpha_i + \beta_j + \varepsilon_{ij}, \quad \varepsilon_{ij} \sim N(0, \sigma^2)$$

• the total variation in the data can be decomposed:

$$SST = SS(Tr) + SS(Bl) + SSE$$

- 'twoway' refers to the fact that there are two factors in the experiment(Two "ways" of the data table)
- The method is called <u>analysis</u> of <u>variance</u>, because the testing is carried out by comparing certain variances.

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Computation - decomposition and the ANOVA table

### Formulas for sums of squares

• Sum of squares for blocks (persons) ("Variance explained by the block part of the model")

$$SS(Bl) = k \cdot \sum_{i=1}^{l} \hat{\beta}_{j}^{2}$$

 The sum of squares for the residuals ("residual variance after model fit")

$$SSE = \sum_{i=1}^{k} \sum_{j=1}^{l} (y_{ij} - \hat{\alpha}_i - \hat{\beta}_j - \hat{\mu})^2$$

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Hypothesis test (F-test)

# Twoway ANOVA: hypothesis of no effect of persons (blocks), Theorem 8.22

• We want to compare (more than 2) means  $\mu + \beta_i$  in the model

$$Y_{ij} = \mu + \alpha_i + \beta_j + \varepsilon_{ij}, \quad \varepsilon_{ij} \sim N(0, \sigma^2)$$

• So we can express the hypothesis

 $H_{0.Bl}$ :  $\beta_i = 0$  for all i

 $H_{1,Bl}: \quad \beta_i \neq 0 \quad \text{for at least one } i$ 

• Under  $H_{0,Bl}$  the following is true:

$$F_{Bl} = \frac{SS(Bl)/(l-1)}{SSE/((k-1)(l-1))}$$

follows an F-distribution with l-1 and (k-1)(l-1) degrees of freedom

## Twoway ANOVA: hypothesis of no effect of treatment, Theorem 8.22

• We want to compare (more than 2) means  $\mu + \alpha_i$  in the model

$$Y_{ij} = \mu + \alpha_i + \beta_j + \varepsilon_{ij}, \quad \varepsilon_{ij} \sim N(0, \sigma^2)$$

So we can express the hypothesis:

 $H_{0,Tr}: \quad \alpha_i = 0 \quad \text{for all } i$ 

 $H_{1,Tr}: \quad \alpha_i \neq 0 \quad \text{for at least one } i$ 

• Under  $H_{0,Tr}$  the following is true:

$$F_{Tr} = \frac{SS(Tr)/(k-1)}{SSE/((k-1)(l-1))}$$

is F-distributed with k-1 and (k-1)(l-1) degrees of freedom

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Hypothesis test (F-test)

### F-distribution and treatments hypothesis

Hypothesis test (F-test)

### F-distribution and blocks hypothesis

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Post hoc analysis

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### Hypothesis test (F-test

### The twoway ANOVA table

Source of	Deg. of	Sums of	Mean sum of	Test-	<i>p</i> -
variation	freedom	squares	squares	statistic $F$	value
Treatment	k-1	SS(Tr)	$MS(Tr) = \frac{SS(Tr)}{k-1}$	$F_{\mathrm{Tr}} = \frac{MS(Tr)}{MSE}$	$P(F > F_{\mathrm{Tr}})$
Block	l-1	SS(Bl)	$MS(Bl) = \frac{SS(Bl)}{l-1}$	$F_{\rm Bl} = \frac{MS(Bl)}{MSE}$	$P(F > F_{\rm Bl})$
Residual	(k-1)(l-1)	SSE	$MSE = \frac{SSE}{(k-1)(l-1)}$		
Total	n-1	SST			

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Post hoc analys

### Post hoc confidence interval

- As for oneway (Use methods 8.9 and 8.10) substitute (n-k) degrees of freedom with (k-1)(l-1) (and use MSE from twoway).
- Can be done with either treatments or blocks
- A single pre-planned confidence interval for the difference between treatment *i* and *j* is found as:

$$\bar{y}_i - \bar{y}_j \pm t_{1-\alpha/2} \sqrt{\frac{SSE}{n-k} \left(\frac{1}{n_i} + \frac{1}{n_j}\right)} \tag{1}$$

where  $t_{1-\alpha/2}$  is based on the t-distribution with (k-1)(l-1) degrees of freedom.

• If all M = k(k-1)/2 combinations of pairwise confidence intervals are found use the formula M times but each time with  $\alpha_{\text{Bonferroni}} = \alpha/M$ .

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### Post hoc pairwise hypothesis test

• A single pre-planned level  $\alpha$  hypothesis tests:

$$H_0: \mu_i = \mu_j, \ H_1: \mu_i \neq \mu_j$$

is carried out as:

$$t_{\text{obs}} = \frac{\bar{y}_i - \bar{y}_j}{\sqrt{MSE\left(\frac{1}{n_i} + \frac{1}{n_j}\right)}} \tag{2}$$

and:

$$p$$
 – value =  $2P(t > |t_{obs}|)$ 

where the *t*-distribution with (k-1)(l-1) degrees of freedom is used.

• If all M = k(k-1)/2 combinations of pairwise confidence intervals are found use the formula M times but each time with  $\alpha_{\text{Bonferroni}} = \alpha/M$ .

Model control

### Variance homogeneity

Look at box-plot to check whether the variability seems different for the groups

```
## Check assumption of homogeneous variance
## Save the fit
fit <- lm(y ~ treatm + block)
## Box plot
par(mfrow=c(1,2))
plot(treatm, fit$residuals, y, xlab="Treatment")
## Box plot
plot(block, fit$residuals, xlab="Block")
```

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Model control

### Normal assumption

### Look at qq-normal plot

```
## Check the assumption of normality of residuals
## qq-normal plot of residuals
qqnorm(fit$residuals)
qqline(fit$residuals)
## Or with a Wally plot
require (MESS)
qqwrap <- function(x, y, ...) {qqnorm(y, main="",...);</pre>
## Can we see a deviating qq-norm plot?
wallyplot(fit$residuals, FUN = qqwrap)
```

A complete example - from the book

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Agenda

### Agenda

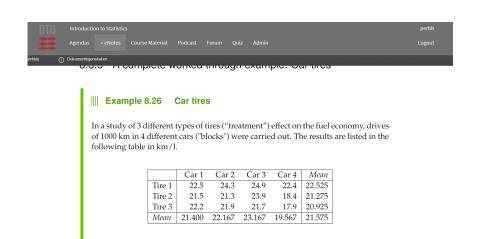
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A complete example - from the book

### A complete example - from the book

plotting:



Let us analyse these data with a two-way ANOVA model, but first some explorative