Example p. 21 / Beispiel S. 21

FK

automatic

Working directory

```
> setwd("D:/kronthafranz/Documents/01Lehre/06Quantitative Forschungsmethoden dt en")
```

Load data

> load("D:/kronthafranz/Documents/01Lehre/06Quantitative Forschungsmethoden dt en/05ANOVA/production.RData")

Define factor

```
> production <- within(production, {
+  f_method <- as.factor(method)
+ })</pre>
```

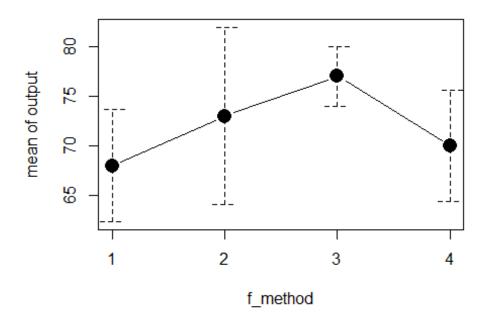
Descriptive statistics

```
> library(abind, pos=15)
> library(e1071, pos=16)
> numSummary(production[,"output", drop=FALSE], groups=production$f_method,
   statistics=c("mean", "sd", "quantiles"), quantiles=c(0,.25,.5,.75,1))
            sd 0%
                               75% 100% output:n
 mean
                    25% 50%
1 68 5.366563 61 65.50 67.5 69.50
                                     77
                                               6
2 73 8.532292 62 66.00 75.0 78.75
                                     83
                                               6
3 77 2.898275 73 75.25 77.0 78.75
                                     81
                                               6
4 70 5.329165 62 67.25 70.0 74.25
```

Plot means

```
> with(production, plotMeans(output, f_method, error.bars="conf.int",
level=0.95,
+ connect=TRUE))
```

Plot of Means



Check assumptions

Independence of observations

Matter of design of the experiment

Equality of variance

```
> with(production, tapply(output, f_method, var, na.rm=TRUE))

1  2  3  4
28.8 72.8 8.4 28.4
> with(production, tapply(output, f_method, var, na.rm=TRUE))

1  2  3  4
28.8 72.8 8.4 28.4
> leveneTest(output ~ f_method, data=production, center="mean")

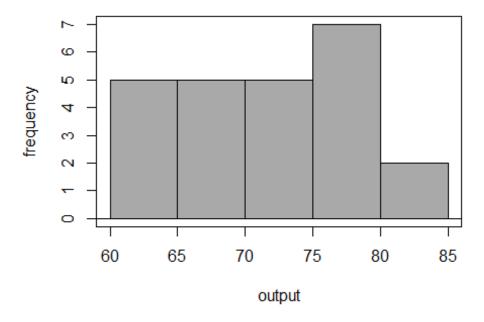
Levene's Test for Homogeneity of Variance (center = "mean")

Df F value Pr(>F)
group 3 2.6873 0.074 .

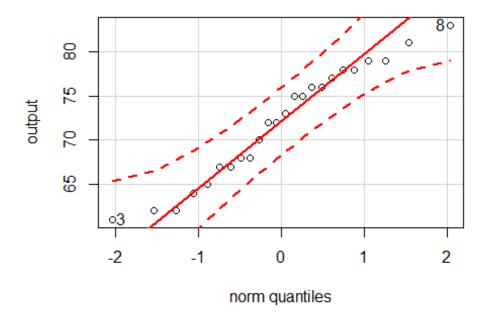
20
---
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

Normal distribution

```
> with(production, Hist(output, scale="frequency", breaks="Sturges",
col="darkgray"))
```



```
> with(production, qqPlot(output, dist="norm", id.method="y", id.n=2,
+ labels=rownames(production)))
```



```
3 8
1 24

> normalityTest(~output, test="shapiro.test", data=production)

Shapiro-Wilk normality test

data: output
W = 0.95365, p-value = 0.3245
```

ANOVA

```
> AnovaModel.1 <- aov(output ~ f_method, data=production)</pre>
> summary(AnovaModel.1)
            Df Sum Sq Mean Sq F value Pr(>F)
f_method
             3
                  276
                         92.0
                                2.659 0.0761 .
Residuals
            20
                  692
                         34.6
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
> with(production, numSummary(output, groups=f_method, statistics=c("mean",
"sd")))
  mean
             sd data:n
    68 5.366563
   73 8.532292
```

```
77 2.898275
                    6
   70 5.329165
                    6
> TukeyHSD(AnovaModel.1)
 Tukey multiple comparisons of means
    95% family-wise confidence level
Fit: aov(formula = output ~ f_method, data = production)
$f_method
   diff
                lwr
                        upr
                                p adj
2-1
      5 -4.5054003 14.5054 0.4717578
3-1
      9 -0.5054003 18.5054 0.0674888
4-1
    2 -7.5054003 11.5054 0.9342653
3-2 4 -5.5054003 13.5054 0.6472435
4-2
     -3 -12.5054003 6.5054 0.8134157
     -7 -16.5054003 2.5054 0.1998434
```

Interpretation:

H0 is rejected (significance level of 10%)

There is a difference in the productivity of production methods

Tukey Post Hoc shows that there is a difference between method 1 and method 3