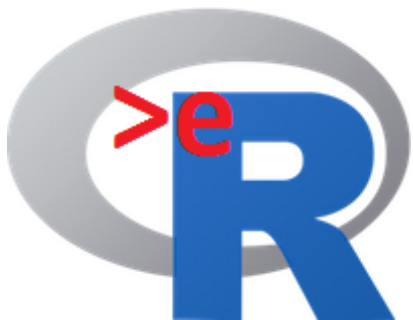




This course was developed as a part of the VLIR-UOS Cross-Cutting project s:

- Statistics: 2011-2016, 2017.
- Statistics: 2017.
- Statistics for development : 2018-2020.



The >eR-Biostat initiative
Making R based education materials in
statistics accessible for all

An introduction to R: Short Version (2017)

Part 4: statistical modeling 2

Developed by

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ER-BioStat



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Overview

1. Two-way ANOVA.
2. More about two-way ANOVA.
3. More about linear regression.

Statistical modeling : Two-way ANOVA

Model formulation

$$Y_{ijk} = \mu + \alpha_i + \beta_j + \alpha\beta_{ij} + \varepsilon_{ijk}$$

μ Overall mean

α_i Main effect of factor A

β_j Main effect of factor B

$\alpha\beta_{ij}$ Interaction effect

ε_{ijk} Random error

Example 1: Reading the data

```
> spwh3<-read.table('c:\\projects\\wseda\\spwh3.txt',  
  header=FALSE,na.strings="NA", dec=".")  
> names(spwh3)<-c("id","y","x1","gender")
```

Example 1: The data

```
> print(spwh3)
      id      y  x1 gender
1      1 10.111368   1     0
2      2  9.948930   1     0
3      3 10.322560   1     0
.      .      .     .     .
.      .      .     .     .
59    59 30.030490   3     1
60    60 29.541542   3     1
>
```

Both x1 and gender are numerical objects !!!!

For an ANOVA model the independent variables are suppose to be factors.

Example 2: The data

	y	f1	f2
1	10	A1	B1
2	11	A1	B1
3	12	A1	B1
4	9	A2	B1
5	7	A2	B1
6	6	A2	B1
7	11	A1	B2
8	13	A1	B2
9	14	A1	B2
10	7	A2	B2
11	5	A2	B2
12	8	A2	B2

Two factors: f1 and f2

Three observations per combination.

```
> f1<-c("A1","A1","A1","A2","A2","A2","A1","A1","A1","A2","A2","A2")
> f2<-c("B1","B1","B1","B1","B1","B1","B2","B2","B2","B2","B2","B2")
> y<-c(10,11,12,9,7,6,11,13,14,7,5,8)
> data.frame(y,f1,f2)
```


Which null hypotheses we test ?

$$H_0 : \alpha_1 = \alpha_2 \quad \text{No treatment effect of factor A}$$

$$H_0 : \beta_1 = \beta_2 \quad \text{No treatment effect of factor B}$$

$$\text{No interaction effects} \quad H_0 : \alpha\beta_{11} = \alpha\beta_{12} = \alpha\beta_{21} = \alpha\beta_{22}$$

Example 1: A model without interaction

```
> fit.1<-aov(y~as.factor(x1)+as.factor(gender))
```

```
> anova(fit.1)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
as.factor(x1)	2	1034.81	517.40	2244.8	< 2.2e-16 ***
as.factor(gender)	1	1509.98	1509.98	6551.3	< 2.2e-16 ***
Residuals	56	12.91	0.23		

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

Example 1: A model with interaction

```
fit.2<-aov(y~as.factor(x1)+as.factor(gender)
           +as.factor(x1)*as.factor(gender))
```

```
> anova(fit.2)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
as.factor(x1)	2	1034.81	517.40	2171.959	<2e-16	***
as.factor(gender)	1	1509.98	1509.98	6338.599	<2e-16	***
as.factor(x1):as.factor(gender)	2	0.04	0.02	0.091	0.9131	
Residuals	54	12.86	0.24			

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

```
>
```

Example 1: Testing model 1 versus model 2

```
> anova(fit.1,fit.2)
```

Analysis of Variance Table

Model 1: $y \sim \text{as.factor}(x1) + \text{as.factor}(\text{gender})$

Model 2: $y \sim \text{as.factor}(x1) + \text{as.factor}(\text{gender}) + \text{as.factor}(x1) * \text{as.factor}(\text{gender})$

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	56	12.9073				
2	54	12.8639	2	0.0434	0.091	0.9131



F-test for the interaction

Example 2: A model without interaction

```
> fit.1<-aov(y~f1+f2)
> anova(fit.1)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
f1	1	70.083	70.083	31.4066	0.0003325 ***
f2	1	0.750	0.750	0.3361	0.5763122
Residuals	9	20.083	2.231		

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

Example 2: A model with interaction

```
> fit.2<-aov(y~f1+f2+f1*f2)
> anova(fit.2)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
f1	1	70.083	70.083	35.0417	0.0003539	***
f2	1	0.750	0.750	0.3750	0.5572922	
f1:f2	1	4.083	4.083	2.0417	0.1909016	
Residuals	8	16.000	2.000			

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

Example 2: Testing model 1 versus model 2

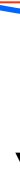
```
> anova(fit.1,fit.2)
```

Analysis of Variance Table

Model 1: $y \sim f1 + f2$

Model 2: $y \sim f1 + f2 + f1 * f2$

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	9	20.083				
2	8	16.000	1	4.0833	2.0417	0.1909



F-test for the interaction

Example 2: means by factor level

```
> tapply(y,f1,mean)
```

A1	A2
11.83333	12.00000

Factor 1

```
> tapply(y,f2,mean)
```

B1	B2
9.166667	14.666667

Factor 2

```
> ind<-list(f1,f2)
```

```
> ind
```

```
[[1]]
```

```
[1] "A1" "A1" "A1" "A2" "A2" "A2" "A1" "A1" "A1" "A2" "A2" "A2"
```

```
[[2]]
```

```
[1] "B1" "B1" "B1" "B1" "B1" "B1" "B1" "B2" "B2" "B2" "B2" "B2"
```

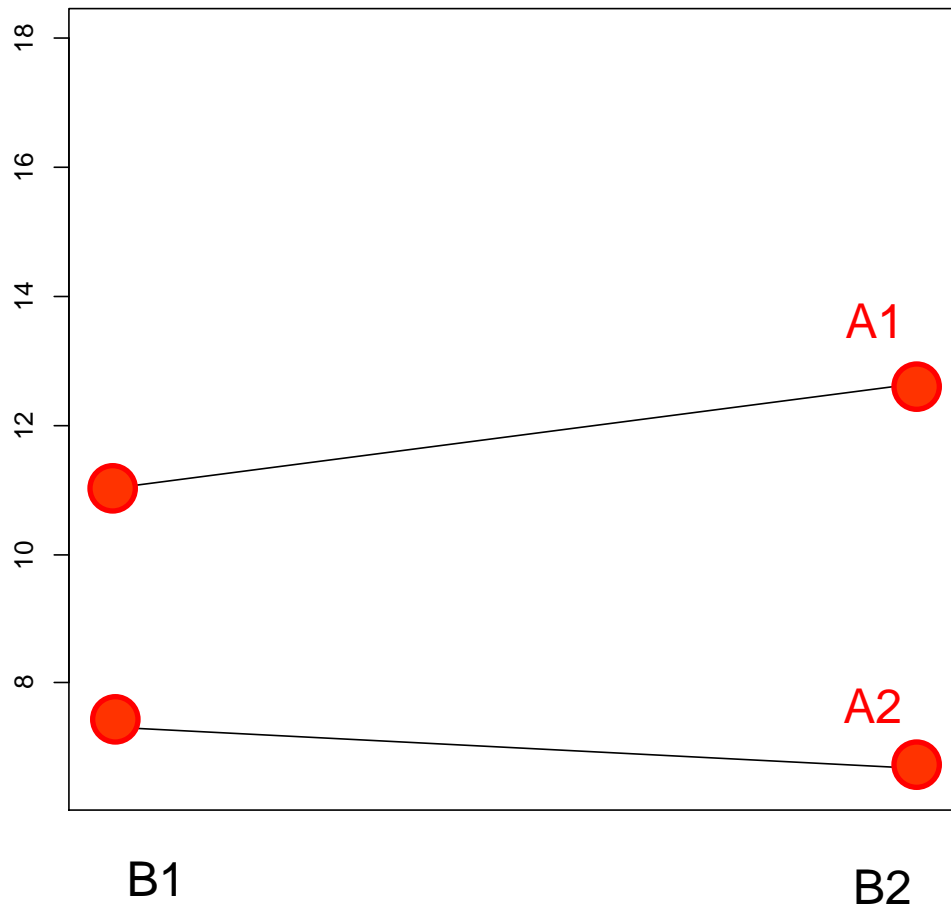
```
> m<-tapply(y,ind,mean)
```

```
> m
```

	B1	B2
A1	11.000000	12.666667
A2	7.333333	16.666667

Cell means

Interaction plot: Example 2



Cell means

	B1	B2
A1	11.000000	12.666667
A2	7.333333	6.666667

Example 3: The data

y	f1	f2
1	10	A1 B1
2	11	A1 B1
3	12	A1 B1
4	9	A2 B1
5	7	A2 B1
6	6	A2 B1
7	11	A1 B2
8	13	A1 B2
9	14	A1 B2
10	17	A2 B2
11	15	A2 B2
12	18	A2 B2

Two factors: f1 and f2

Three observations per combination.

```
> f1<-c("A1","A1","A1","A2","A2","A2","A1","A1","A1","A2","A2","A2")
> f2<-c("B1","B1","B1","B1","B1","B1","B2","B2","B2","B2","B2","B2")
> y<-c(10,11,12,9,7,6,11,13,14,17,15,18)
> data.frame(y,f1,f2)
```

Example 3: A model with interaction

```
> fit.2<-aov(y~f1+f2+f1*f2)
> anova(fit.2)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
f1	1	0.083	0.083	0.0417	0.8433536
f2	1	90.750	90.750	45.3750	0.0001471 ***
f1:f2	1	44.083	44.083	22.0417	0.0015517 **
Residuals	8	16.000	2.000		

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

Example 3: means by factor level

```
> tapply(y,f1,mean)
```

A1	A2
11.83333	12.00000

Factor 1

```
> tapply(y,f2,mean)
```

B1	B2
9.166667	14.666667

Factor 2

```
> ind<-list(f1,f2)
```

```
> ind
```

```
[[1]]
```

```
[1] "A1" "A1" "A1" "A2" "A2" "A2" "A1" "A1" "A1" "A2" "A2" "A2"
```

```
[[2]]
```

```
[1] "B1" "B1" "B1" "B1" "B1" "B1" "B1" "B2" "B2" "B2" "B2" "B2"
```

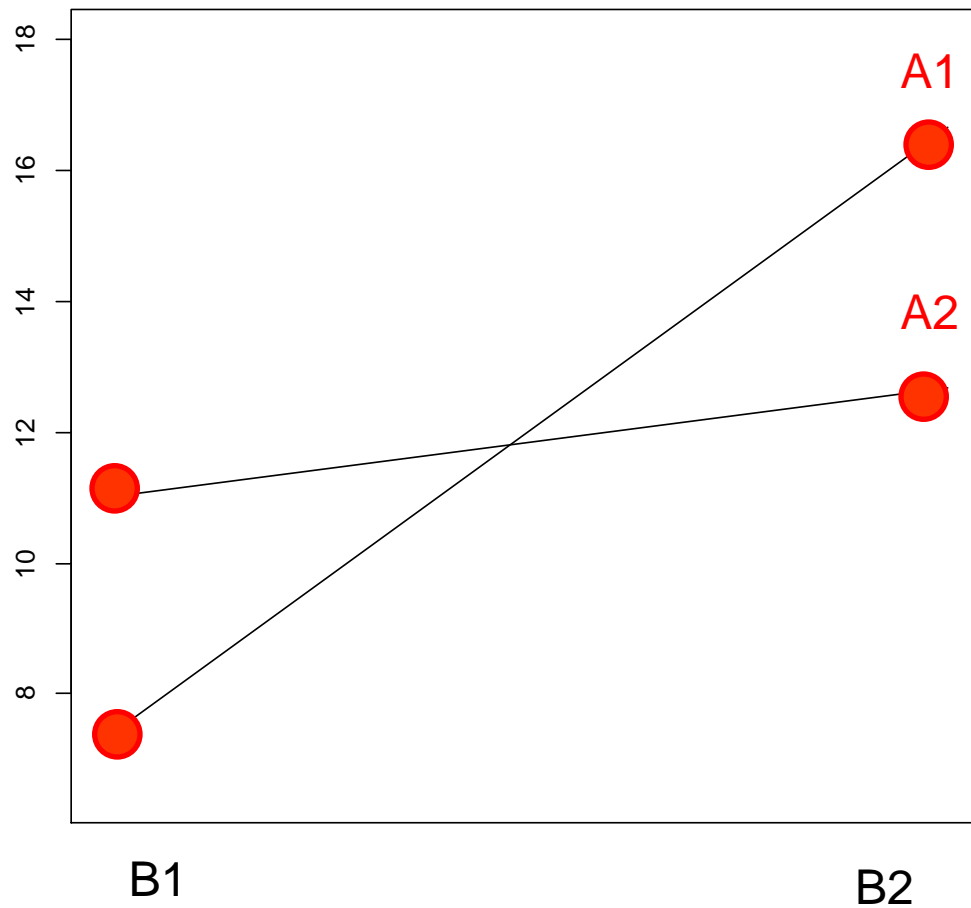
```
> m<-tapply(y,ind,mean)
```

```
> m
```

	B1	B2
A1	11.000000	12.666667
A2	7.333333	16.666667

Cell means

Interaction plot: Example 3



Cell means

	B1	B2
A1	11.000000	12.66667
A2	7.333333	16.66667

Statistical modeling : More about two-way ANOVA

Reading the data

```
> spwh3<-read.table('c:\\projects\\wseda\\spwh3.txt',  
header=FALSE,na.strings="NA", dec=".")  
> names(spwh3)<-c("id","y","x1","gender")  
> attach(spwh3)
```

Two-way ANOVA model

```
> fit.2<-aov(y~as.factor(x1)+as.factor(gender)+as.factor(x1)*as.factor(gender))  
> anova(fit.2)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
as.factor(x1)	2	1034.81	517.40	2171.959	<2e-16	***
as.factor(gender)	1	1509.98	1509.98	6338.599	<2e-16	***
as.factor(x1):as.factor(gender)	2	0.04	0.02	0.091	0.9131	
Residuals	54	12.86	0.24			

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

Stepwise procedure

```
> slm1 <- step(fit.2)
```

```
Start:  AIC=-80.4
```

```
y ~ as.factor(x1) + as.factor(gender) + as.factor(x1) * as.factor(gender)
```

	Df	Sum of Sq	RSS	AIC
- as.factor(x1):as.factor(gender)	2	0.043	12.907	-84.193
<none>			12.864	-80.395

```
Step:  AIC=-84.19
```

```
y ~ as.factor(x1) + as.factor(gender)
```

	Df	Sum of Sq	RSS	AIC
<none>			12.91	-84.19
- as.factor(x1)	2	1034.81	1047.72	175.60
- as.factor(gender)	1	1509.98	1522.89	200.04

Stepwise procedure

```
> summary(slm1)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
as.factor(x1)	2	1034.81	517.40	2244.8	< 2.2e-16	***
as.factor(gender)	1	1509.98	1509.98	6551.3	< 2.2e-16	***
Residuals	56	12.91	0.23			

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

Statistical modeling : More about Linear regression

Reading the data

```
> spwh2<-read.table('c:\\projects\\wseda\\spwh2.txt',
header=FALSE,
+                               ,na.strings="NA", dec=".")
> dim(spwh2)
[1] 100    5
>
> names(spwh2)<-c("id","y","x1","x2","x3")
> attach(spwh2)
```

The following object(s) are masked from spwh2 (position 3) :

```
id x1 x2 x3 y
```

Fitting two models

```
> fit.1<-lm(y~x1+x2)
> anova(fit.1)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
x1	1	164.2	164.2	27.152	1.059e-06 ***
x2	1	7409.7	7409.7	1224.980	< 2.2e-16 ***
Residuals	97	586.7	6.0		

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

```
> fit.2<-lm(y~x1+x2+x3)
> anova(fit.2)
```

Analysis of Variance Table

Response: y

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
x1	1	164.2	164.2	758.98	< 2.2e-16 ***
x2	1	7409.7	7409.7	34241.81	< 2.2e-16 ***
x3	1	566.0	566.0	2615.44	< 2.2e-16 ***
Residuals	96	20.8	0.2		

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

Testing model 1 versus model 2

```
> anova(fit.1,fit.2)
```

```
Analysis of Variance Table
```

```
Model 1: y ~ x1 + x2
```

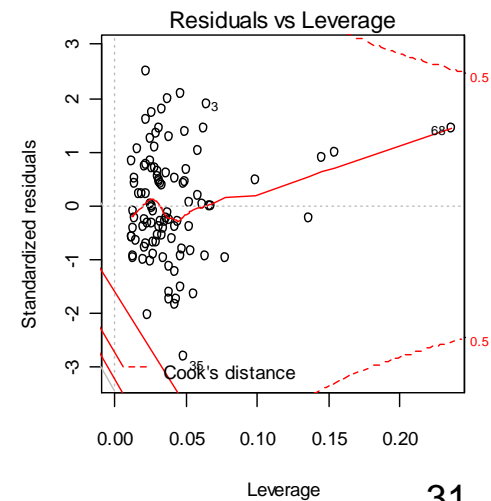
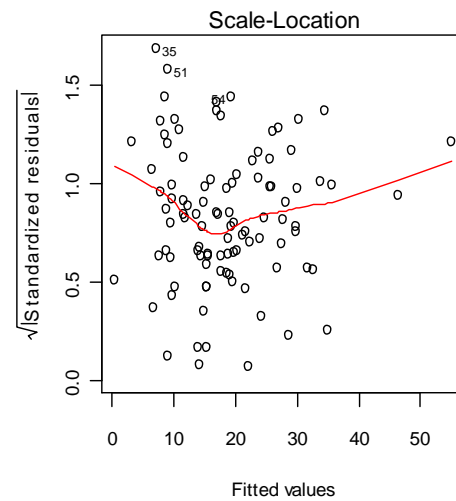
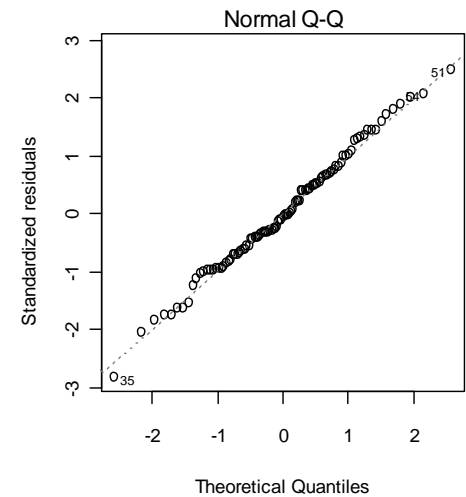
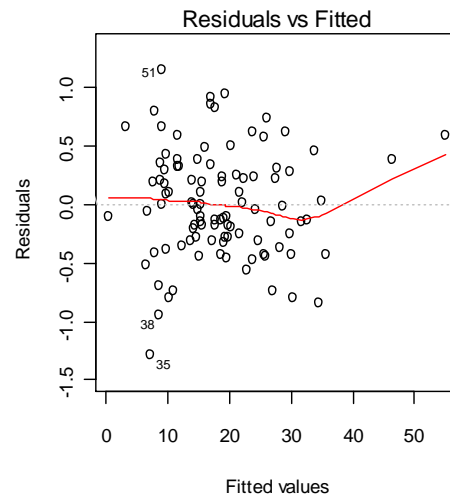
```
Model 2: y ~ x1 + x2 + x3
```

	Res.Df	RSS	Df	Sum of Sq	F	Pr(>F)
1	97	586.74				
2	96	20.77	1	565.97	2615.4	< 2.2e-16 ***

```
---
```

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

```
> par(mfrow=c(2,2))
> plot(fit.2)
```



Single terms deletions

```
> drop1(fit.2, test="F")  
Single term deletions
```

Model:

```
y ~ x1 + x2 + x3
```

	Df	Sum of Sq	RSS	AIC	F value	Pr(F)
<none>			20.8	-149.1		
x1	1	76.6	97.4	3.4	354.21	< 2.2e-16 ***
x2	1	7865.3	7886.1	442.8	36347.01	< 2.2e-16 ***
x3	1	566.0	586.7	182.9	2615.44	< 2.2e-16 ***

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

AIC and likelihood

```
> AIC(fit.2)
[1] 136.6403
> logLik(fit.2)
'log Lik.' -63.32017 (df=5)
```