

Multi-Factor Experiments

- Twoway anova
- Interactions
- More than two factors

Hypertension: Effect of biofeedback

Biofeedback + Medication	Biofeedback	Medication	Control
158	188	186	185
163	183	191	190
173	198	196	195
178	178	181	200
168	193	176	180

Main effects

Treatment means:

		Medication		Total
		no	yes	
Biofeedback	no	190	186	188
	yes	188	168	178
Total	189	177	183	

main effect of biofeedback: $188 - 178 = 10 \text{ mmHg}$

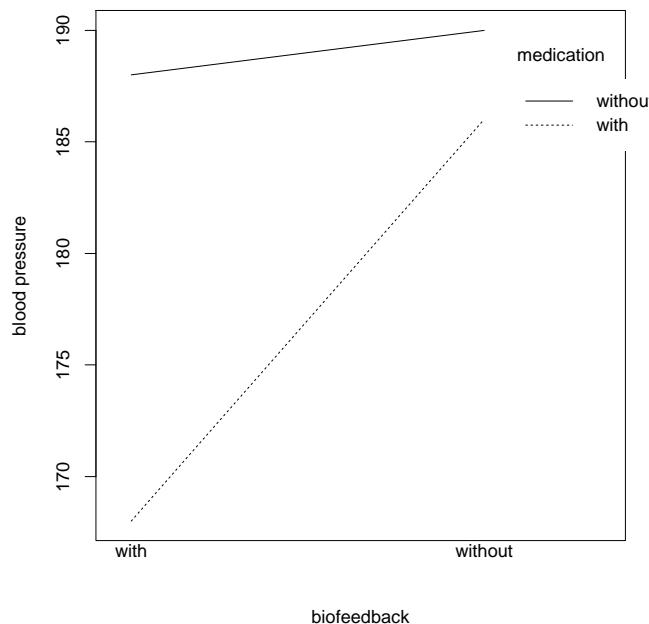
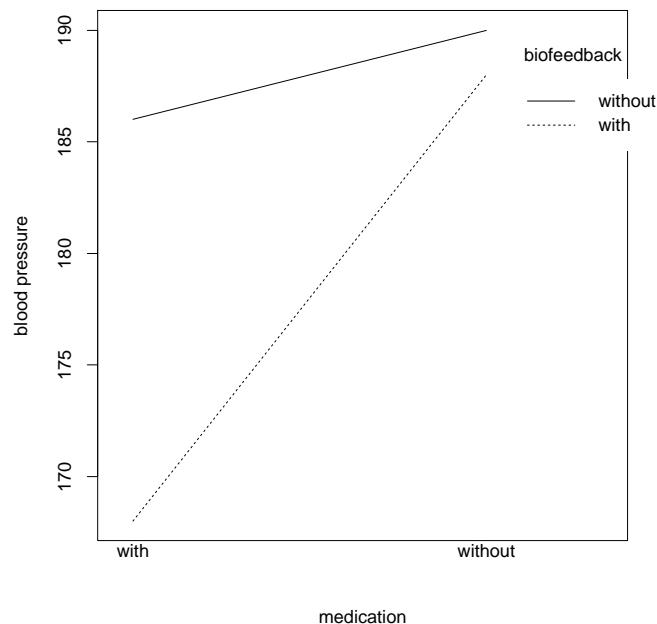
Interaction plot

Effect of biofeedback with medi: 18 mmHg

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Effect of biofeedback without medi: 2 mmHg

→ **Interaction**



Model for two factors

$$Y_{ijk} = \mu + A_i + B_j + (AB)_{ij} + \epsilon_{ijk}$$

$$i = 1, \dots, I; j = 1, \dots, J; k = 1, \dots, n.$$

$$\sum A_i = 0, \sum B_j = 0, \sum_i (AB)_{ij} = \sum_j (AB)_{ij} = 0.$$

A_i : i th effect of factor A

B_j : j th effect of factor B

$\mu + A_i + B_j$: overall mean + effect of factor A on level i
+ effect of factor B on level j

$(AB)_{ij}$: deviation from additive model

Parameter estimation

$\hat{\mu} = y_{...}$, $\hat{A}_i = y_{i..} - y_{...}$ and $\hat{B}_j = y_{.j.} - y_{...}$

$\widehat{AB}_{ij} = y_{ij.} - (\hat{\mu} + \hat{A}_i + \hat{B}_j) = y_{ij.} - y_{i..} - y_{.j.} + y_{...}$

Medication A

Biofeedback B	no	yes	Total
no	190	186	$y_{.1.} = 188$
yes	188	168	$y_{.2.} = 178$
Total	$y_{1..} = 189$	$y_{2..} = 177$	$y_{...} = 183$

$\hat{\mu} = 183$, $\hat{A}_1 = -\hat{A}_2 = 6$, $\hat{B}_1 = -\hat{B}_2 = 5$

Predicted values of the additive model

Treatment means: $(\hat{\mu} + \hat{A}_i + \hat{B}_j)$

		Medication	
		no	yes
Biofeedback	no	194	182
	yes	184	172

$$\widehat{AB}_{11} = \widehat{AB}_{22} = -\widehat{AB}_{12} = -\widehat{AB}_{21} = 4.$$

Decomposition of Variability

$$SS_{tot} = SS_A + SS_B + SS_{AB} + SS_{res}$$

$$SS_{tot} = \sum \sum \sum (y_{ijk} - y...)^2$$

$$SS_A = \sum \sum \sum (y_{i..} - y...)^2$$

$$SS_B = \sum \sum \sum (y_{.j.} - y...)^2$$

$$SS_{AB} = \sum \sum \sum (y_{ij.} - y_{i..} - y_{.j.} + y...)^2$$

SS_{res} = «Difference»

degrees of freedom: main effect with I levels: $I - 1$ df,
interaction between two factors with I and J levels:
 $(I - 1)(J - 1)$ df.

Anova table

Source	SS	df	MS	F	P value
medi	720	1	720	11.52	0.004
bio	500	1	500	8.00	0.012
medi:bio	320	1	320	5.12	0.038
residual	1000	16	62.5		
total	2540	19			

Treatment effects

	effect size	C.I.
medi without bio:	$y_{21.} - y_{11.} = 186 - 190 = -4$	(-18, 10)
medi with bio:	$y_{22.} - y_{12.} = 168 - 188 = -20$	(-34, -6)
bio without medi:	$y_{12.} - y_{11.} = 188 - 190 = -2$	(-16, 12)
bio with medi:	$y_{22.} - y_{21.} = 168 - 186 = -18$	(-32, -4)

(standard error: $\sqrt{2 \cdot MS_{res}/5} = 5$)

Efficiency of factorials

		factorial	
		medication	
biofeedback		no	yes
no		6	6
yes		6	6

Two separate studies

medication	
no	yes
6	6

and

biofeedback	
no	yes
6	6

or:

control	medi	bio
8	8	8

More than two factors

	bio/medi	bio	medi	control
without diet	158	188	186	185
	163	183	191	190
	173	198	196	195
	178	178	181	200
	168	193	176	180
with diet	162	162	164	205
	158	184	190	199
	153	183	169	171
	182	156	165	161
	190	180	177	179

Treatment means

without diet(C)				with diet (C)			
medication (A)				medication (A)			
bio (B)	no	yes	total	bio (B)	no	yes	total
no	190	186	188	no	183	173	178
yes	188	168	178	yes	173	169	171
total	189	177	183	total	178	171	174.5

Main effects and interactions

main effects: medication (A): $183.5 - 174 = 9.5$

biofeedback (B): $183 - 174.5 = 8.5$

diet(C): $183 - 174.5 = 8.5$

2-way interactions AB, AC, BC:

		medi (A)		
		Bio (B)	no	yes
Bio (B)	no	186.5	179.5	183
	yes	180.5	168.5	174.5
total		183.5	174	178.75

3-way interaction ABC

Model and Anova table

$$Y_{ijkl} = \mu + A_i + B_j + C_k + (AB)_{ij} + (AC)_{ik} + (BC)_{jk} + (ABC)_{ijk} + \epsilon_{ijkl}$$

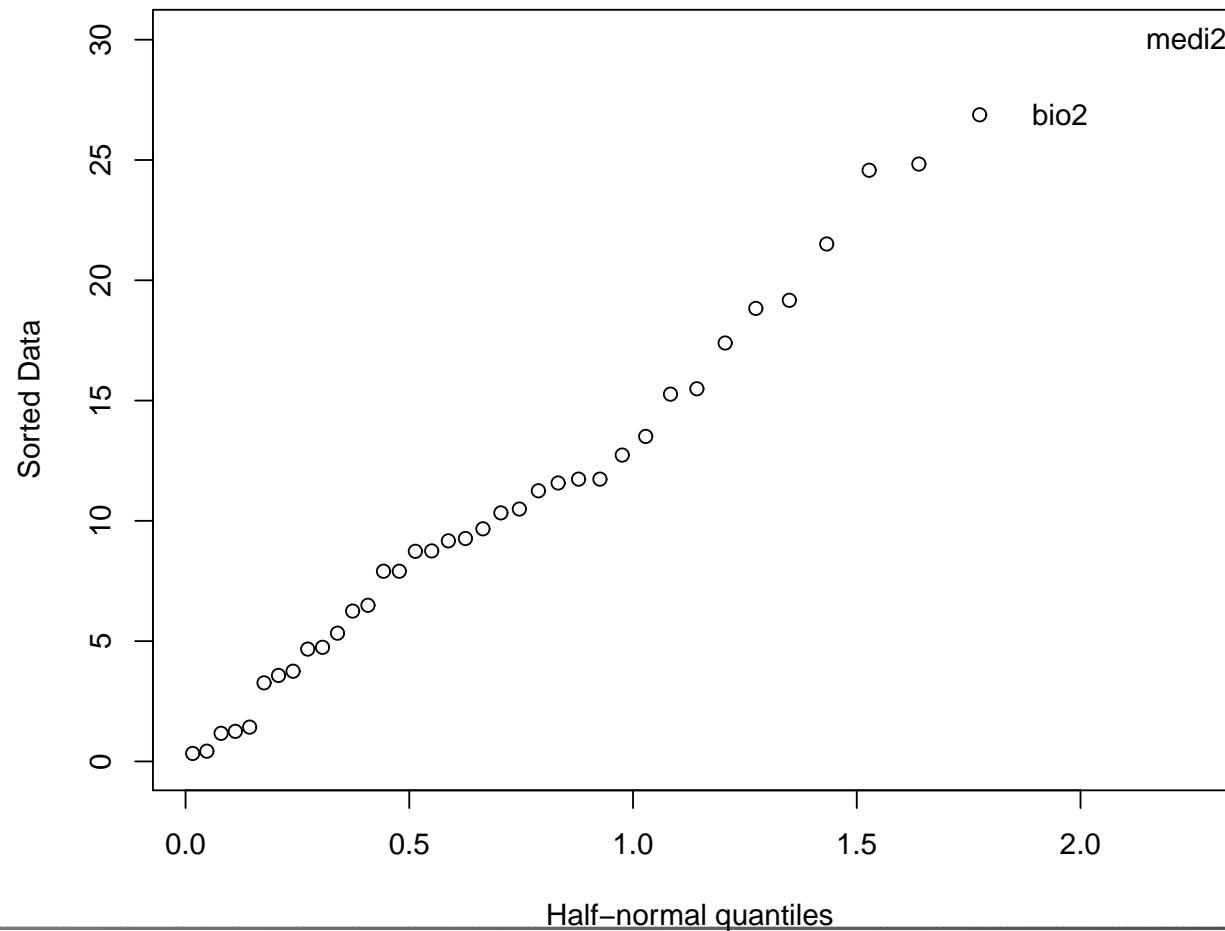
$$\sum A_i = 0, \dots, \sum_k (ABC)_{ijk} = 0$$

Source	df	MS=SS/df	F
A	$I - 1$	MS_A / MS_{res}	
B	$J - 1$	MS_B / MS_{res}	
C	$K - 1$	MS_C / MS_{res}	
AB	$(I - 1)(J - 1)$	MS_{AB} / MS_{res}	
AC	$(I - 1)(K - 1)$	MS_{AC} / MS_{res}	
BC	$(J - 1)(K - 1)$	MS_{BC} / MS_{res}	
ABC	$(I - 1)(J - 1)(K - 1)$	MS_{ABC} / MS_{res}	
Residual	«Difference»	$MS_{res} = \hat{\sigma}^2$	
Total	$IJKn - 1$		

Anova table

Source	SS	df	MS	F	P value
medi	902.5	1	902.5	6.33	0.017
bio	722.5	1	722.5	5.06	0.031
diet	722.5	1	722.5	5.06	0.031
medi:bio	62.5	1	62.5	0.44	0.51
medi:diet	62.5	1	62.5	0.44	0.51
bio:diet	22.5	1	22.5	0.16	0.69
medi:bio:diet	302.5	1	302.5	2.12	0.15
Residual	4566.0	32	142.7		
Total	7363.5	39			

Half normal plot



Unbalanced Factorials

uncorrelated estimators:

$$SS_{tot} = SS_A + SS_B + SS_{AB} + \underbrace{SS_{res}}_{SS_C + \dots + SS_{res'}}$$

correlated estimators:

$$SS_{tot} = SS'_A + SS'_B + SS'_{AB} + SS_C + \dots + SS_{res'}$$

ss Typ I: SS_A ignores all other SS

ss Typ II: SS_A takes into account all other main effects,
ignores all interactions

ss Typ III: SS_A takes into account all other effects

Calculation of SS's

by model comparison

For SS Typ I:

model 1: $Y_{ijk} = \mu + \epsilon_{ijk}$

$$SS_{e1} = SS_T$$

model 2: $Y_{ijk} = \mu + A_i + \epsilon_{ijk}$

$$SS_{e2}$$

model 3: $Y_{ijk} = \mu + A_i + B_j + \epsilon_{ijk}$

$$SS_{e3}$$

model 4: $Y_{ijk} = \mu + A_i + B_j + AB_{ij} + \epsilon_{ijk}$

$$SS_{e4} = SS_{res}$$

Rat genotype

- Litters of rats are separated from their natural mother and given to another female to raise.
- 2 factors: mother's genotype (A, B, I, J) and litter's genotype (A, B, I, J)
- response: average weight gain of the litter.

Full model

```
> summary(aov(y~mother*genotype,data=gen))
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
mother	3	771.61	257.202	4.7419	0.005869	*
genotype	3	63.63	21.211	0.3911	0.760004	
mother:genotype	9	824.07	91.564	1.6881	0.120053	
Residuals	45	2440.82	54.240			

```
> summary(aov(y~genotype*mother,data=gen))
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
genotype	3	60.16	20.052	0.3697	0.775221	
mother	3	775.08	258.360	4.7632	0.005736	*
genotype:mother	9	824.07	91.564	1.6881	0.120053	
Residuals	45	2440.82	54.240			

SS Typ II in

```
> drop1(mod1,test="F")  
Model: y ~ mother * genotype  
              Df  Sum Sq    RSS    AIC   F value Pr(F)  
<none>                 2440.8 257.04  
mother:geno  9   824.07 3264.9 256.79  1.6881 0.1201
```

```
> drop1(mod1,.~.,test="F")  
Model: y ~ mother * genotype  
              Df  Sum Sq    RSS    AIC   F value Pr(F)  
<none>                 2440.8 257.04  
mother      3   582.25 3023.1 264.09  3.5782 0.02099 *  
geno        3   591.69 3032.5 264.28  3.6362 0.01968 *  
mother:geno  9   824.07 3264.9 256.79  1.6881 0.12005
```

Offer for a 6-year old car

- Planned experiment to see whether the offered cash for the same medium-priced car depends on gender or age (young, middle, elderly) of the seller.
- 6 factor combinations with 6 replications each.
- Response variable y is offer made by a car dealer (in \$ 100)
- Covariate: overall sales volume of the dealer

Analysis of Covariance

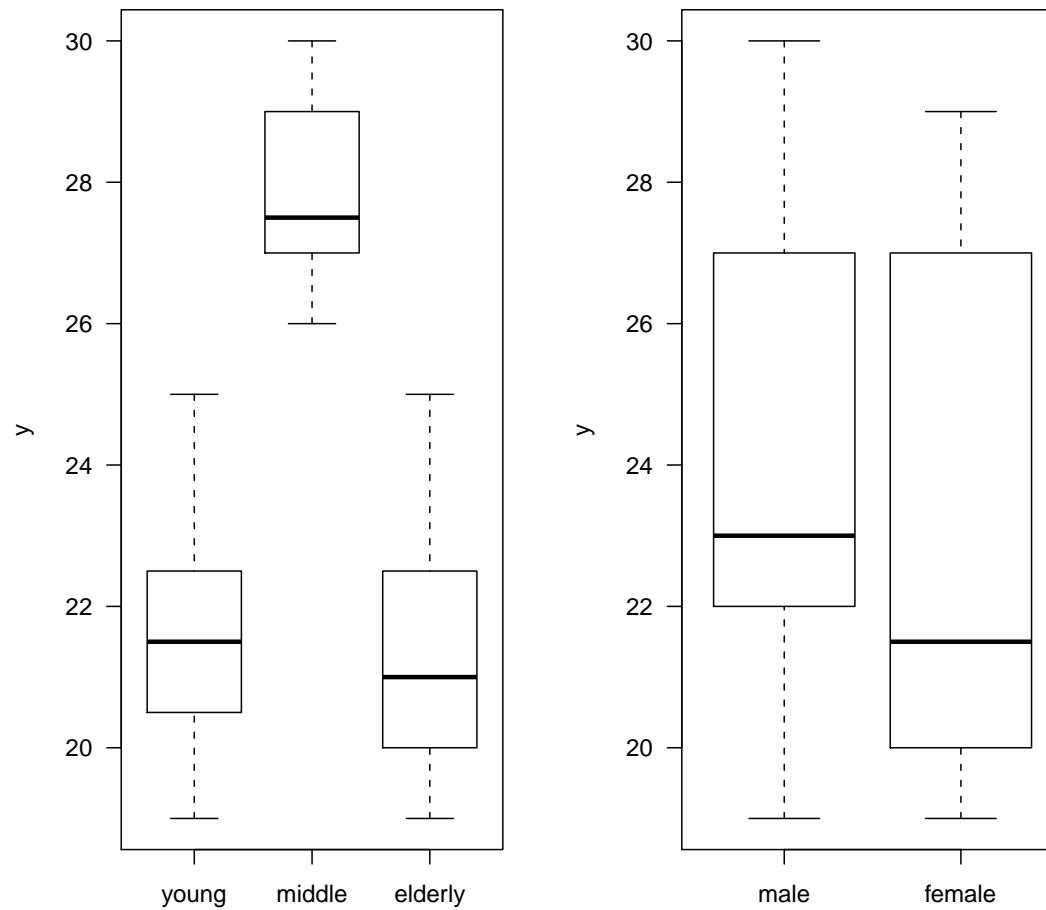
- Covariates can reduce MS_{res} , thereby increasing power for testing.
- Baseline or pretest values are often used as covariates. A covariate can adjust for differences in characteristics of subjects in the treatment groups.
- It should be related only to the response variable and not to the treatment variables (factors).
- We assume that the covariate will be linearly related to the response and that the relationship will be the same for all levels of the factor (no interaction between covariate and factors).

Model for two-way ANCOVA

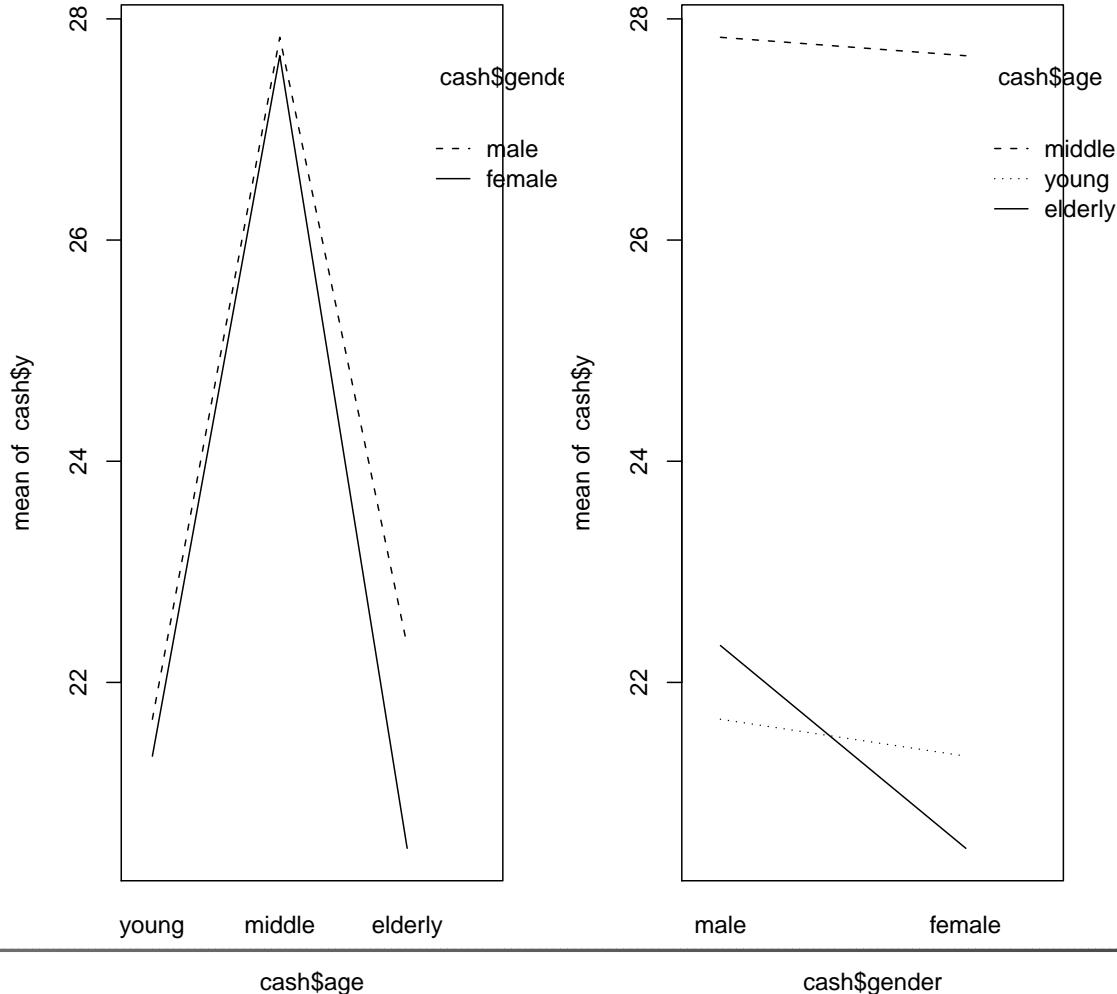
$$Y_{ijk} = \mu + \theta x_{ijk} + A_i + B_j + (AB)_{ij} + \epsilon_{ijk}$$

$$\sum A_i = \sum B_j = \sum (AB)_{ij} = 0, \quad \epsilon_{ijk} \sim \mathcal{N}(0, \sigma^2)$$

Effect of Age and Gender



Interaction effect of Age and Gender

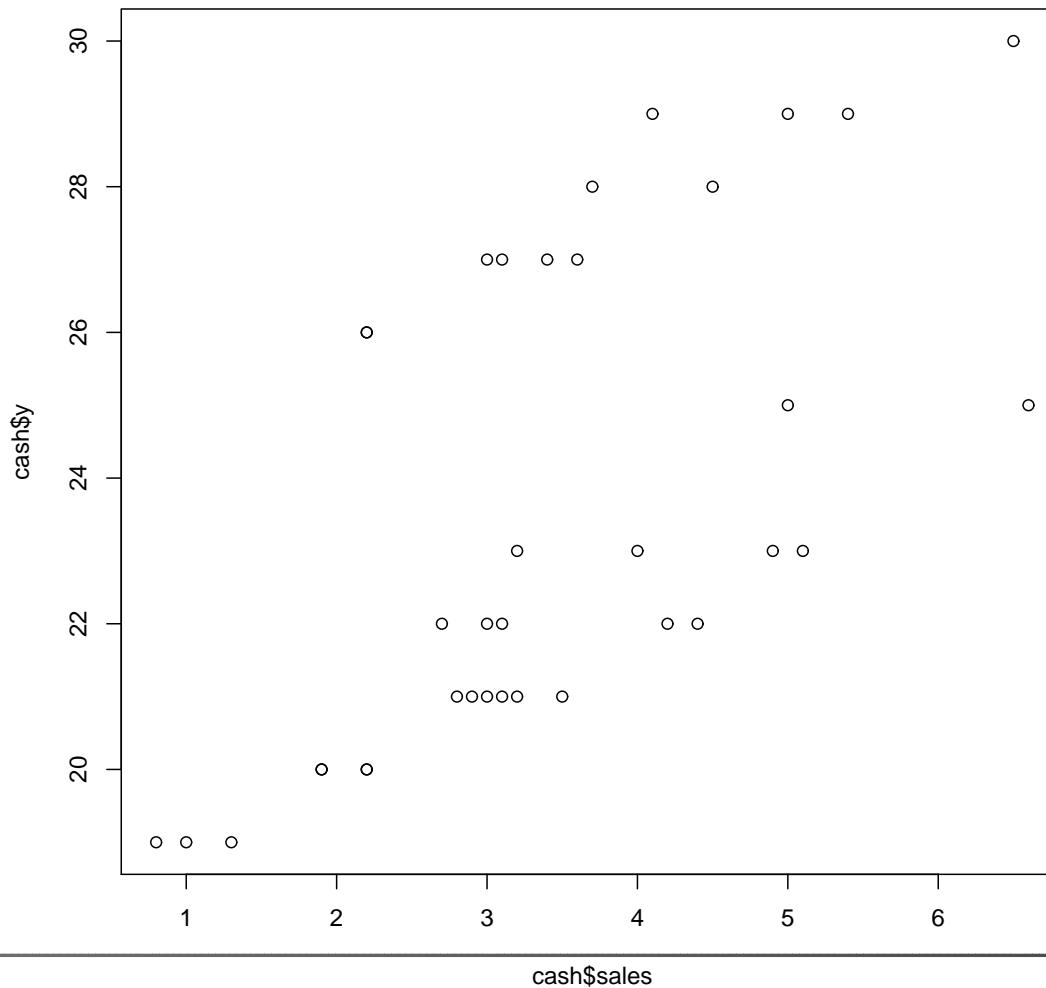


Two-way Anova

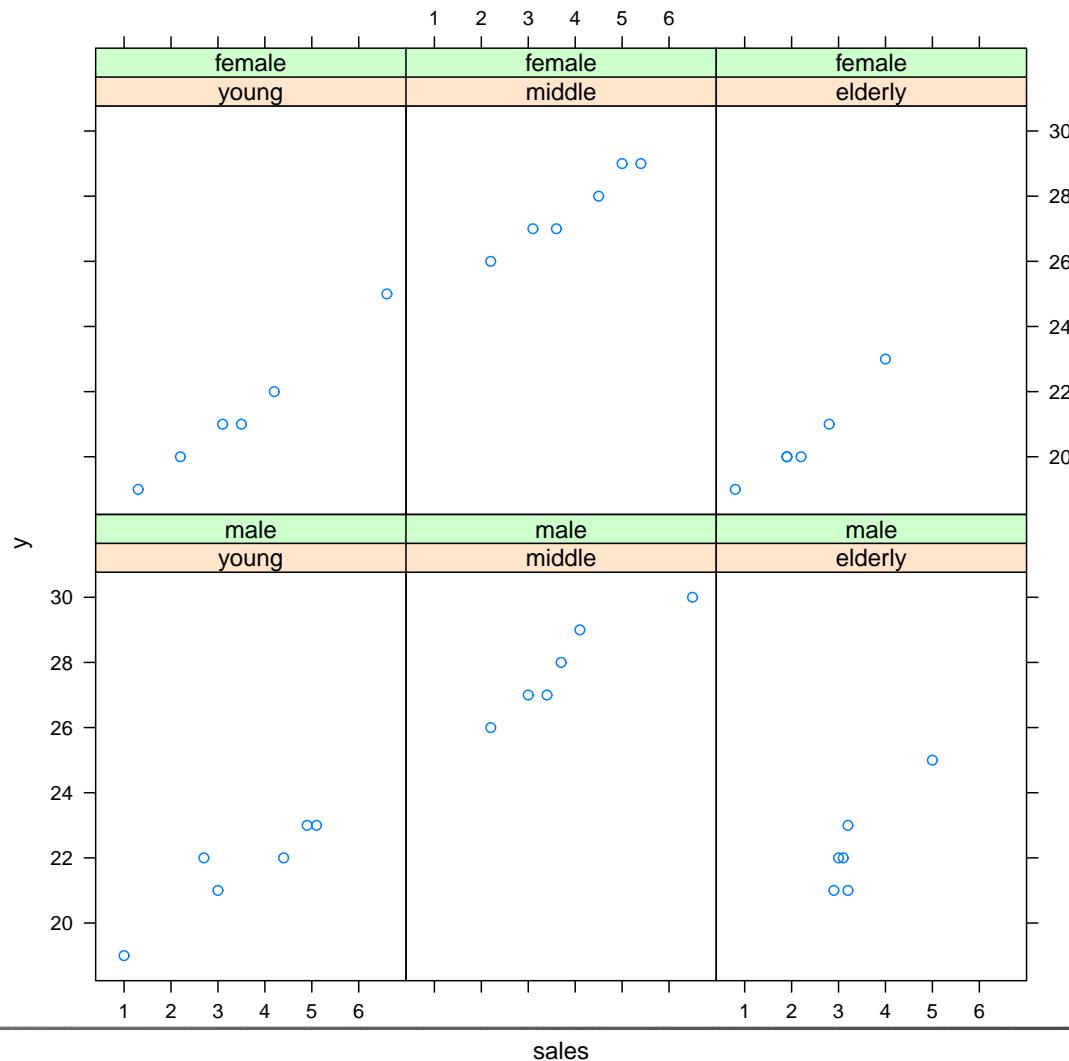
```
> mod1=aov(y~age*gender,data=cash)
> summary(mod1)

             Df  Sum Sq Mean Sq Fvalue    Pr(>F)
age           2 316.72 158.36 66.29 9.79e-12*
gender        1   5.44   5.44   2.28  0.1416
age:gender    2   5.06   2.53   1.06  0.3597
Residuals   30  71.67   2.39
```

Sales and Cash Offer



Sales and Cash Offer by Group



Two-way Ancova

```
> mod2=aov(y~sales+age*gender,data=cash)
> summary(mod2)

             Df  Sum Sq Mean Sq Fvalue    Pr(>F)
sales          1 157.37 157.37 550.22 < 2e-16 ***
age            2 231.52 115.76 404.75 < 2e-16 ***
gender         1   1.51   1.51    5.30  0.02874 *
age:gender     2   0.19   0.10    0.34  0.71422
Residuals     29   8.29   0.29
```