

## Ch. 5 Two-way ANOVA: Fixed effect model

### Equal sample sizes

### 1 Assumptions and models

- There are two factors, factors A and B, that are of interest.
- Factor A is studied at  $a$  levels, and factor B at  $b$  levels;  
All  $ab$  factor level combinations are included in the study.
- Format of (balanced) data set

		Factor B				
		level 1	...	level $j$	...	level $b$
Factor A	level 1	$Y_{111}, \dots, Y_{11n}$		$Y_{1j1}, \dots, Y_{1jn}$		$Y_{1b1}, \dots, Y_{1bn}$
	$\vdots$					
	level $i$	$Y_{i11}, \dots, Y_{i1n}$		$Y_{ij1}, \dots, Y_{ijn}$		$Y_{ib1}, \dots, Y_{ibn}$
	$\vdots$					
	level $a$	$Y_{a11}, \dots, Y_{a1n}$		$Y_{aj1}, \dots, Y_{ajn}$		$Y_{ab1}, \dots, Y_{abn}$

#### Mean parameters

		Factor B				
		level 1	...	level $j$	...	level $b$
Factor A	level 1	$\mu_{11}$		$\mu_{1j}$		$\mu_{1b} \quad \mu_{1\cdot}$
	$\vdots$					
	level $i$	$\mu_{i1}$		$\mu_{ij}$		$\mu_{ib} \quad \mu_{i\cdot}$
	$\vdots$					
	level $a$	$\mu_{a1}$		$\mu_{aj}$		$\mu_{ab} \quad \mu_{a\cdot}$
		$\mu_{\cdot 1}$		$\mu_{\cdot j}$		$\mu_{\cdot b}$

### 1.1 Cell means model (for balanced data)

$$Y_{ijk} = \mu_{ij} + \varepsilon_{ijk}, \quad i = 1, \dots, a, \quad j = 1, \dots, b, \quad k = 1, \dots, n.$$

$\mu_{ij}$ ,  $i = 1, \dots, a, j = 1, \dots, b$ , are mean parameters;

Error terms  $\varepsilon_{ijk} \stackrel{\text{i.i.d}}{\sim} \mathcal{N}(0, \sigma^2)$ ,  $i = 1, \dots, a, j = 1, \dots, b, k = 1, \dots, n$ .

- Feature of model

$$Y_{ijk} \stackrel{\text{independent}}{\sim} \mathcal{N}(\mu_{ij}, \sigma^2), \quad i = 1, \dots, a, \quad j = 1, \dots, b, \quad k = 1, \dots, n.$$

### 1.2 Factor Effects Model

$$Y_{ijk} = \mu_{..} + \alpha_i + \beta_j + (\alpha\beta)_{ij} + \varepsilon_{ijk}, \quad i = 1, \dots, a, \quad j = 1, \dots, b, \quad k = 1, \dots, n.$$

$\mu_{..}$  is a constant or overall mean  $(\mu_{..} = \frac{1}{ab} \sum_{i=1}^a \sum_{j=1}^b \mu_{ij})$ ;

$\alpha_i$  are constants subject to the restriction  $\sum_{i=1}^a \alpha_i = 0$   
 $(\alpha_i = \mu_{i.} - \mu_{..}, \quad \text{the main effect for factor A at the } i\text{th level})$ ;

$\beta_j$  are constants subject to the restriction  $\sum_{j=1}^b \beta_j = 0$   
 $(\beta_j = \mu_{.j} - \mu_{..}, \quad \text{the main effect for factor B at the } j\text{th level})$ ;

$(\alpha\beta)_{ij}$  are constants subject to the restrictions

$$\sum_{i=1}^a (\alpha\beta)_{ij} = 0, \quad j = 1, \dots, b, \quad \sum_{j=1}^b (\alpha\beta)_{ij} = 0, \quad i = 1, \dots, a$$

$(\alpha\beta)_{ij} = \mu_{ij} - \mu_{i.} - \mu_{.j} + \mu_{..}$ ,  
the interaction effect for factor A at the  $i$ th level and factor B at the  $j$ th level).

- Feature of model

$$Y_{ijk} \stackrel{\text{independent}}{\sim} \mathcal{N}(\mu_{..} + \alpha_i + \beta_j + (\alpha\beta)_{ij}, \sigma^2), \quad i = 1, \dots, a, \quad j = 1, \dots, b, \quad k = 1, \dots, n.$$

## 1.3 Notations

## Means

$$\begin{aligned}\bar{Y}_{...} &= \frac{1}{nab} \sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n Y_{ijk} && \text{(The overall mean)} \\ \bar{Y}_{i..} &= \frac{1}{nb} \sum_{j=1}^b \sum_{k=1}^n Y_{ijk} && \text{(The mean for the } i\text{th factor level of A)} \\ \bar{Y}_{.j.} &= \frac{1}{na} \sum_{i=1}^a \sum_{k=1}^n Y_{ijk} && \text{(The mean for the } j\text{th factor level of B)} \\ \bar{Y}_{ij.} &= \frac{1}{n} \sum_{k=1}^n Y_{ijk} && \text{(The mean for the } (i, j)\text{th treatment)}\end{aligned}$$

## Sums of Squares

$$\begin{aligned}SS_{TR} &= n \sum_{i=1}^a \sum_{j=1}^b (\bar{Y}_{ij.} - \bar{Y}_{...})^2 \\ SS_E &= \sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n (Y_{ijk} - \bar{Y}_{ij.})^2 \\ SS_T &= \sum_{i=1}^a \sum_{j=1}^b \sum_{k=1}^n (Y_{ijk} - \bar{Y}_{...})^2 \\ SS_A &= nb \sum_{i=1}^a (\bar{Y}_{i..} - \bar{Y}_{...})^2 && \text{(Factor A sum of squares)} \\ SS_B &= na \sum_{j=1}^b (\bar{Y}_{.j.} - \bar{Y}_{...})^2 && \text{(Factor B sum of squares)} \\ SS_{AB} &= n \sum_{i=1}^a \sum_{j=1}^b (\bar{Y}_{ij.} - \bar{Y}_{i..} - \bar{Y}_{.j.} + \bar{Y}_{...})^2 && \text{(AB interaction sum of squares)}\end{aligned}$$

## Relationships (partitioning)

$$\underbrace{Y_{ijk} - \bar{Y}_{...}}_{\text{Total deviation}} = \underbrace{\bar{Y}_{ij.} - \bar{Y}_{...}}_{\substack{\text{Deviation of} \\ \text{estimated treatment mean} \\ \text{around overall mean}}} + \underbrace{Y_{ijk} - \bar{Y}_{ij.}}_{\substack{\text{Deviation around} \\ \text{estimated treatment mean}}}$$

$$SS_T = SS_{TR} + SS_E, \quad SS_{TR} \perp SS_E$$

$$\underbrace{\bar{Y}_{ij.} - \bar{Y}_{...}}_{\substack{\text{Deviation of} \\ \text{estimated treatment mean} \\ \text{around overall mean}}} = \underbrace{(\bar{Y}_{i..} - \bar{Y}_{...})}_{\text{A main effect}} + \underbrace{(\bar{Y}_{.j.} - \bar{Y}_{...})}_{\text{B main effect}} + \underbrace{(\bar{Y}_{ij.} - \bar{Y}_{i..} - \bar{Y}_{.j.} + \bar{Y}_{...})}_{\text{AB interaction effect}}$$

$$SS_{TR} = SS_A + SS_B + SS_{AB}$$

## 2 Estimation for mean parameters and $\sigma^2$

### 2.1 Cell means model

The least squares and maximum likelihood estimators of  $\mu_{ij}$

$$\hat{\mu}_{ij} = \bar{Y}_{ij}.$$

Fitted value of  $Y_{ijk}$ :

$$\hat{\mu}_{ij} = \bar{Y}_{ij}.$$

Residual related to  $Y_{ijk}$ :

$$e_{ijk} = Y_{ijk} - \bar{Y}_{ij}.$$

Parameter	Estimator	Confidence interval
$\mu_{ij}$	$\hat{\mu}_{ij} = \bar{Y}_{ij} \sim \mathcal{N}(\mu_{ij}, \frac{\sigma^2}{n})$	$\bar{Y}_{ij} \pm qt(1 - \alpha/2; (n - 1)ab) \sqrt{\frac{MSE}{n}}$
$\mu_{i.}$	$\hat{\mu}_{i.} = \bar{Y}_{i.} \sim \mathcal{N}(\mu_{i.}, \frac{\sigma^2}{bn})$	$\bar{Y}_{i.} \pm qt(1 - \alpha/2; (n - 1)ab) \sqrt{\frac{MSE}{bn}}$
$\mu_{.j}$	$\hat{\mu}_{.j} = \bar{Y}_{.j} \sim \mathcal{N}(\mu_{.j}, \frac{\sigma^2}{an})$	$\bar{Y}_{.j} \pm qt(1 - \alpha/2; (n - 1)ab) \sqrt{\frac{MSE}{an}}$
$\sigma^2$	$\widehat{\sigma^2} = MSE, \frac{SS_E}{\sigma^2} \sim \chi^2_{(n-1)ab}$	

### 2.2 Factor effects model

Parameter	(LS or MLE) Estimator
$\mu_{..}$	$\hat{\mu}_{..} = \bar{Y}_{..}$
$\alpha_i = \mu_{i.} - \mu_{..}$	$\hat{\alpha}_i = \bar{Y}_{i.} - \bar{Y}_{..}$
$\beta_j = \mu_{.j} - \mu_{..}$	$\hat{\beta}_j = \bar{Y}_{.j} - \bar{Y}_{..}$
$(\alpha\beta)_{ij} = \mu_{ij} - \mu_{i.} - \mu_{.j} + \mu_{..}$	$\widehat{(\alpha\beta)}_{ij} = \bar{Y}_{ij} - \bar{Y}_{i.} - \bar{Y}_{.j} + \bar{Y}_{..}$

### 3 F tests

**Two-Way ANOVA table**

Source of Variation	df	Sum of Squares (SS)	Mean Square (MS)	$F^*$ -value	Test for
Factor A	$a - 1$	$SS_A$	$MS_A = \frac{SS_A}{a-1}$	$F^* = \frac{MS_A}{MS_E}$	$\mathcal{H}_0 : \mu_{.1} = \dots = \mu_{.a}$ (factor A main effects)
Factor B	$b - 1$	$SS_B$	$MS_B = \frac{SS_B}{b-1}$	$F^* = \frac{MS_B}{MS_E}$	$\mathcal{H}_0 : \mu_{.1} = \dots = \mu_{.b}$ (factor B main effects)
AB interactions	$(a - 1)(b - 1)$	$SS_{AB}$	$MS_{AB} = \frac{SS_{AB}}{(a-1)(b-1)}$	$F^* = \frac{MS_{AB}}{MS_E}$	$\mathcal{H}_0 : \text{all } (\alpha\beta)_{ij} = 0$ (interactions)
Error(Residuals)	$ab(n - 1)$	$SS_E$	$MS_E = \frac{SS_E}{ab(n-1)}$		
Total	$nab - 1$	$SS_T$			

*Castle Bakery Example*, p.817.

Factor A (display height) has 3 factor levels, and factor B (display width) has 2 factor levels.

```
> y <- read.table("CH19TA07.DAT")
  V1 V2 V3 V4
  1  47  1  1  1
  .....
  4  40  1  2  2
  5  62  2  1  1
  .....
  8  71  2  2  2
  9  41  3  1  1
  .....
 12 46  3  2  2
```

Set the data frame, as the one-way layout case.

```
> data <- y[,1]
> height <- y[,2]
> width <- y[,3]

> sales.df <- data.frame(data=data,height=factor(height),width=factor(width))
```

The first step is often to look at the data graphically.

```
> plot.factor(height,data)      # Factor A  (boxplot)
> plot(height,data)             (dot plot)

> plot.factor(width,data)      # Factor B
> plot(width,data)

> plot(sales.df)               # data frame
```

To obtain the ANOVA table, type `aov`, and then `summary`.

```
> anova <- aov(data~height+width+height*width,sales.df)
      # or      anova <- aov(data~height*width,sales.df)
      # compare aov(data~height,sales.df) aov(data~width,sales.df)
> summary(anova)
      # or      anova(lm(data~height*width,sales.df))
      Df  Sum Sq  Mean Sq  F value  Pr(>F)
height    2   1544.00   772.00   74.7097  5.754e-05
width     1    12.00    12.00    1.1613  0.3226
height:width 2    24.00    12.00    1.1613  0.3747
Residuals  6    62.00    10.33

> model.tables(anova,type="means")      # treatment means
```

Tables of means

Grand mean 51

height	1	2	3
	44	67	42

width	1	2
	50	52

height:width	Dim	1 = height,	2 = width
	1	2	
	1	45	43
	2	65	69
	3	40	44

## 4 Analysis of factor effects when factors do not interact

- Where are we?

Two possibilities:

- (i) Based on a F test for testing  $\mathcal{H}_0: \text{all } (\alpha\beta)_{ij} = 0$ , we accept  $\mathcal{H}_0$ . This means that factors do not interact.
- (ii) Factors A and B interact only in an unimportant fashion.

- What's next?

Examine whether the main effects for factors A and B are important.

For important A or B main effects, describe the nature of these effects in terms of the factor level means  $\mu_i$ . or  $\mu_j$ , respectively.

*Castle Bakery Example*, p.817.

```
> y <- read.table("CH19TA07.DAT")
> anova(lm(data~height*width,sales.df))
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)	
height	2	1544.00	772.00	74.7097	5.754e-05	# factor A
width	1	12.00	12.00	1.1613	0.3226	# factor B
height:width	2	24.00	12.00	1.1613	0.3747	# AB interaction
Residuals	6	62.00	10.33			

The p-value for testing  $\mathcal{H}_0: \text{all } (\alpha\beta)_{ij} = 0$  is 37.5%, which leads us to accept  $\mathcal{H}_0$ . This means that factors do not interact.

The p-value for testing  $\mathcal{H}_0: \text{all } \beta_j = 0$  is 32.3%, which leads us to accept  $\mathcal{H}_0$ . This means that factor B is not important.

How about factor A? Look at the p-value, which is 5.754e-05. This means that factor A is important. We need analyze the factor effects in terms of  $\mu_1, \mu_2$ , and  $\mu_3$ .

→ Multiple comparison procedure

○ Ch. 3 for analysis of one-way (fixed) factor level effects

Multiple comparison procedures

(cf. Ch. 3 for analysis of one-way (fixed) factor level effects)

- \* (1- $\alpha$ )100 % Tukey simultaneous confidence intervals for all pairwise comparisons  $\mu_i - \mu_{i'}$ :

$$\bar{Y}_i - \bar{Y}_{i'} \pm T * \sqrt{MS_E \left( \frac{1}{n_i} + \frac{1}{n_{i'}} \right)}, \quad i \neq i', i, i' \in \{1, \dots, r\}$$

where

$$T \triangleq \frac{1}{\sqrt{2}} q(1 - \alpha; r, n_T - r), \quad (\text{Table B.9})$$

- \* (1- $\alpha$ )100 % Scheffé simultaneous confidence intervals for the family of contrasts  $L$ :

$$\sum_{i=1}^r c_i \bar{Y}_i \pm S * \sqrt{MS_E \sum_{i=1}^r \frac{c_i^2}{n_i}},$$

where

$$S^2 \triangleq (r - 1) * qF(1 - \alpha; r - 1, n_T - r).$$

- \* (1- $\alpha$ )100 % Bonferroni simultaneous confidence intervals for the  $g$  linear combinations  $L$ :

$$\sum_{i=1}^r c_i \bar{Y}_i \pm B * \sqrt{MS_E \sum_{i=1}^r \frac{c_i^2}{n_i}},$$

where

$$B \triangleq qt\left(1 - \frac{\alpha}{2g}; n_T - r\right).$$



## 5 Analysis of factor effects when interactions are important

- Where are we?

Based on a F test for testing  $\mathcal{H}_0$ : all  $(\alpha\beta)_{ij} = 0$ , we accept  $\mathcal{H}_a$ . This means that factors A and B interact in an important fashion.

- What's next?

Analyze the two factor effects jointly in terms of the treatment means  $\mu_{ij}$

*Hay fever relief*, CH19PR14.DAT.

```
> y <- read.table("CH19PR14.DAT")
> data <- y[,1]
> ingredient1 <- y[,2]
> ingredient2 <- y[,3]

> fever.df <- data.frame(data=data, ingredient1=factor(ingredient1),
                        ingredient2=factor(ingredient2))

> anova(lm(data~ingredient1+ingredient2+ingredient1*ingredient2, fever.df))
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
ingredient1	2	220.020	110.010	1827.86	< 2.2e-16
ingredient2	2	123.660	61.830	1027.33	< 2.2e-16
ingredient1:ingredient2	4	29.425	7.356	122.23	< 2.2e-16
Residuals	27	1.625	0.060		

The p-value for testing  $\mathcal{H}_0$ : all  $(\alpha\beta)_{ij} = 0$  is  $< 2.2\text{e-}16$ , which leads us to accept  $\mathcal{H}_a$ . This means that factors do interact. We need analyze the factor effects in terms of  $\mu_{ij}$

→ Multiple comparison procedure

Multiple pairwise comparisons of treatment means**1° Tukey procedure**

The Tukey  $1 - \alpha$  multiple comparison confidence limits for all pairwise comparisons:

$$D = \mu_{ij} - \mu_{i'j'}$$

are

$$\bar{Y}_{ij.} - \bar{Y}_{i'j'}. \pm T \sqrt{\frac{2MS_E}{n}},$$

where

$$T = \frac{1}{\sqrt{2}} q[1 - \alpha; ab, (n - 1)ab] \quad (\text{Table B.9})$$

**2° Bonferroni procedure**

The Bonferroni  $1 - \alpha$  multiple comparison confidence limits for a group of  $g$  pairwise comparisons:

$$\bar{Y}_{ij.} - \bar{Y}_{i'j'}. \pm B \sqrt{\frac{2MS_E}{n}},$$

where

$$B = t[1 - \alpha/(2g); (n - 1)ab].$$

Multiple Contrasts of Treatment Means**3° Scheffe procedure**

The joint  $(1 - \alpha)$  confidence limits for contrasts of the form:

$$L = \sum \sum c_{ij} \mu_{ij}, \quad \text{where} \quad \sum \sum c_{ij} = 0,$$

are

$$\sum \sum c_{ij} \bar{Y}_{ij.} \pm S * \sqrt{\frac{MS_E}{n} \sum \sum c_{ij}^2},$$

where

$$S^2 = (ab - 1)F[1 - \alpha; ab - 1, (n - 1)ab].$$

**4° Bonferroni procedure (for small number of contrasts)**

$$\sum \sum c_{ij} \bar{Y}_{ij.} \pm B * \sqrt{\frac{MS_E}{n} \sum \sum c_{ij}^2},$$

where

$$B = t[1 - \alpha/(2g); (n - 1)ab].$$

Problem 20.7: *Hay fever relief*, CH19PR14.DAT.

Part c. The Scheffe multiple comparison.

```
> scheffe <- function(coef){
  s.value <- sqrt((3*3-1)*qf(1-.10,3*3-1,(4-1)*3*3));
  lowerbd <- sum(coef*cell.mean)-s.value*sqrt(MSE/n*sum(coef^2));
  upperbd <- sum(coef*cell.mean)+s.value*sqrt(MSE/n*sum(coef^2));
  scheffe <- c(lowerbd, upperbd);
  scheffe
}
```

where `cell.mean` is obtained from

```
> model.tables(aov(data~ingredient1+ingredient2+ingredient1*ingredient2, fever.df),
  type="means")
Grand mean    7.183333

ingredient1    1     2     3
              3.883 7.833 9.833

ingredient2    1     2     3
              4.633 7.933 8.983

ingredient1:ingredient2
              ingredient2
ingredient1  1     2     3
            1  2.475 4.600 4.575
            2  5.450 8.925 9.125
            3  5.975 10.275 13.250

> cell.mean <- matrix(c(2.475,5.450,5.975,4.600,8.925,10.275,4.575,9.125,13.250),
  3, 3)      # Enter a matrix by column.
            # The matrix can also be filled row-wise as
            # matrix(c(2.475, 4.600, ...), 3, 3, byrow=T)
```

Now CI for  $L_1 = (\mu_{12} + \mu_{13})/2 - \mu_{11}$  is

```
> scheffe(matrix(c(-1,.5,.5,0,0,0,0,0,0),3,3, byrow=T))
[1] 1.526296 2.698704
```

Part d. The Tukey multiple comparison procedure

```
> q.value <- 4.32 # In this case, a=3, b=3, n=4, alpha=.1
                  # From Table B.9, q.value=q(1-alpha; a*b, (n-1)*a*b)
                  #                               =q(.9, 3*3, 3*3*3)=4.32
> T <- q.value/sqrt(2)

> tukey <- function(x,y){
  tukey <- c(mean(x)-mean(y)-sqrt(2*MS_E/n)*T,
             mean(x)-mean(y)+sqrt(2*MS_E/n)*T);
  tukey
}
```

Now CI for  $\mu_{11} - \mu_{12}$  is

```
> tukey(x[,1][x[,2]==1&x[,3]==1], x[,1][x[,2]==1&x[,3]==2])
[1] -2.654090 -1.595910
```

Similarly, for  $\mu_{11} - \mu_{13}$

```
> tukey(x[,1][x[,2]==1&x[,3]==1], x[,1][x[,2]==1&x[,3]==3])
[1] -2.629090 -1.570910
```

There are  $\binom{3 \times 4}{2} = 66$  pairwise CIs. From the Tukey multiple comparison procedure, the longest mean relief is that with largest cell mean, i.e.,  $\mu_{33}$ .

## 6 One case per treatment

Question: What happens if  $n = 1$ ?

$$\text{df for Error(Residuals)} = 0; \quad SS_E = 0; \quad SS_T = SS_B = SS_A + SS_B + SS_{AB}.$$

No-interaction model:

$$Y_{ij} = \mu_{..} + \alpha_i + \beta_j + \varepsilon_{ij}, \quad i = 1, \dots, a, \quad j = 1, \dots, b,$$

where

$$\alpha_i = \mu_{i.} - \mu_{..}, \quad \beta_j = \mu_{.j} - \mu_{..}$$

**Two-Way ANOVA Table (n=1)**

Source of Variation	df	Sum of Squares	Mean Square (MS)	F*-value	Null hypothesis
Factor A	$a - 1$	$SS_A$	$MS_A = \frac{SS_A}{a-1}$	$F^* = \frac{MS_A}{MS_{AB}}$	$\mathcal{H}_0 : \alpha_1 = \dots = \alpha_a = 0$
Factor B	$b - 1$	$SS_B$	$MS_B = \frac{SS_B}{b-1}$	$F^* = \frac{MS_B}{MS_{AB}}$	$\mathcal{H}_0 : \beta_1 = \dots = \beta_b = 0$
Error	$(a - 1)(b - 1)$	$SS_{AB}$	$MS_{AB} = \frac{SS_{AB}}{(a-1)(b-1)}$	$\sigma^2$	
Total	$ab - 1$	$SS_T$			

### Sums of Squares

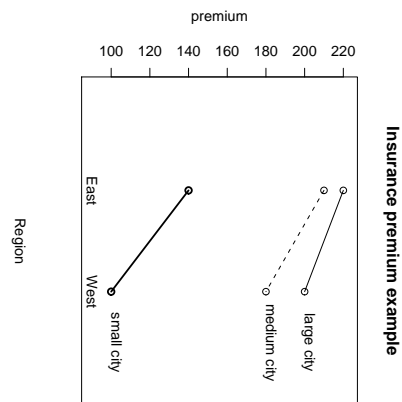
$$SS_T = \sum_{i=1}^a \sum_{j=1}^b (Y_{ij} - \bar{Y}_{..})^2$$

$$SS_A = b \sum_{i=1}^a (\bar{Y}_{i.} - \bar{Y}_{..})^2 \quad (\text{Factor A sum of squares})$$

$$SS_B = a \sum_{j=1}^b (\bar{Y}_{.j} - \bar{Y}_{..})^2 \quad (\text{Factor B sum of squares})$$

$$SS_{AB} = \sum_{i=1}^a \sum_{j=1}^b (Y_{ij} - \bar{Y}_{i.} - \bar{Y}_{.j} + \bar{Y}_{..})^2 \quad (\text{Error})$$

Insurance premium example, p.878



The **R** code for the figure is

```
> y <- read.table("CH21TA02.DAT")
> small <- y[,1][y[,2]==1]
> medium <- y[,1][y[,2]==2]
> large <- y[,1][y[,2]==3]
> region <- c(1,2)

> plot(region, small, type="o", lwd=2,xlim=c(0,3), ylim=c(90,222),
       xlab="Region", ylab="premium", xaxt="n", main="Insurance premium example")

> lines(region, medium,type="o", lty=2 )
> lines(region, large, type="o")

> text(2.5, 102, "small city")
```

```
> text(2.5, 182, "medium city")
> text(2.5, 202, "large city")

> text(1, 90, "East")
> text(2, 90, "West")
```

The ANOVA table is obtained as before.

```
> data <- y[,1]
> size <- y[,2]
> region <- y[,3]
> premium.df <- data.frame(data=data, size=factor(size), region=factor(region))

> anova <- aov(data~size+region, premium.df)
      # compare: aov(data~size+region+size*region, premium.df)
> summary(anova)
```

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
size	2	9300	4650	93	0.01064
region	1	1350	1350	27	0.03510
Residuals	2	100	50		