Chapter 4

Descriptive measures of association for bivariate distributions

Exercise 14) Answer:

```
> duration = faithful$eruptions # the eruption durations
> waiting = faithful$waiting # the waiting period
> cov(duration, waiting) # apply the cov function
[1] 13.978
```

The covariance of the eruption duration and waiting time is 13.978. It indicates a positive linear relationship between the two variables.

(Exercise 15) \underline{Answer} :

```
> CA < -c(70,65,90,95,110,115,120,140,155,150)
> YA<-c(80,100,120,140,160,180,200,220,240,260)
> CB<-c(75,50,95,120,90,115,110,140,175,140)
> YB<-c(80,100,120,140,160,180,200,220,240,260)
> data<-data.frame(CA,YA,CB,YB)</pre>
       • > meanCA<-mean(CA)
         > meanYA<-mean(YA)</pre>
         > cat("meanCA=",meanCA," meanYA=",meanYA," sdCA=",sd(CA)," sdYA=",sd(YA),"\n")
         meanCA= 111 meanYA= 170 sdCA= 31.42893 sdYA= 60.55301
       • > answer1<-data.frame(CA,YA,col1=(CA-meanCA)\land2,col2=(YA-meanYA)\land2,
         + col3=(CA-meanCA)*(YA-meanYA))
         > answer1
                CA
                     YΑ
                         col1 col2 col3
                70
                         1681
                                8100
                                      3690
                     80
                    100
                         2116
                               4900
                                      3220
           3
                                2500
                90
                    120
                          441
                                      1050
           4
                95
                   140
                          256
                                 900
                                       480
                    160
                            1
                                 100
              110
                                        10
           6
              115
                    180
                           16
                                 100
                                        40
           7
              120
                    200
                           81
                                 900
                                       270
              140
                    220
                          841
                                2500
                                      1450
```

If we use the previous datas

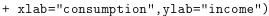
```
> n<-dim(answer1)</pre>
```

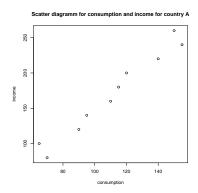
- > covA=sum(answer1\$col3)/(n-1)
- > sdCA=sqrt(sum(answer1\$col1)/(n-1))
- > sdYA=sqrt(sum(answer1\$col2)/(n-1))
- > corA=covA/(sdCA*sdYA)
- > cat("covA=",covA," corA=",corA,"\n")

If we use the R commands

```
> cov(CA, YA)
[1] 1866.667
> cor(CA, YA)
[1] 0.9808474
```

2. > plot(CA,YA,main="Scatter diagramm for consumption and income for country A",





- 3. > meanCB<-mean(CB)
 - > meanYB<-mean(YB)</pre>
 - > cat("meanCB=",meanYB=",meanYB=",sdCB=",sd(CB)," sdYB=",sd(YB),"\n") meanCB= 111 meanYB= 170 sdCB= 35.88562 sdYB= 60.55301
 - > answer2<-data.frame(CB,YB,col1=(CB-meanCB) \land 2,col2=(YB-meanYB) \land 2,
 - + col3=(CB-meanCB)*(YB-meanYB))
 - > answer2

	CB	YΒ	col1	col2	col3
1	75	80	1296	8100	3240
2	50	100	3721	4900	4270
3	95	120	256	2500	800
4	120	140	81	900	-270
5	90	160	441	100	210
6	115	180	16	100	40
7	110	200	1	900	-30
8	140	220	841	2500	1450
9	175	240	4096	4900	4480
Ω	140	260	841	8100	2610

By using the previous datas,

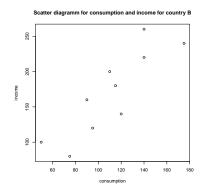
- > covB=sum(answer2\$col3)/(n-1)
- > sdCB=sqrt(sum(answer2\$col1)/(n-1))
- > sdYB=sqrt(sum(answer2\$col2)/(n-1))

```
> corB=covB/(sdCB*sdYB)
> cat("covB=",covB," corB=",corB,"\n")
covB= 1866.667 corB= 0.8590345
```

If we use the R commands

> cov(CB,YB)
[1] 1866.667
> cor(CB,YB)
[1] 0.8590345

> plot(CB, YB, main="Scatter diagramm for consumption and income for country B",
 + xlab="consumption", ylab="income")



- Since YA and YB identical, we have meanYA=meanYB and sdYA=sdYB. We remark next that meanCA=meanCB (even if CA≠CB) and sdCA<sdCB which means that the incomes of country A are less dispersed around their mean than the ones of country B.
- 4. We note that covA=covB. Since the covariance measures the degree to which two variables are linearly associated, the consumptions and incomes of country A are as much as (linearly) linked than the ones of country B.

The covariance is multiplied by 100^2 .

The covariance is multiplied by 100.

The covariance is multiplied by 7.85.

```
(d) > CAd<-CA+10
> YAd<-YA+10
> cov(CAd,YAd)
[1] 1866.667
```

The covariance remains unchanged.

```
(e) > CAe<-CA
> YAe<-YA+10
> cov(CAd, YAd)
[1] 1866.667
```

The covariance remains unchanged.

All these results are well known. Indeed, let a_1 , a_2 , b_1 and b_2 be scalars, and let X and Y be random variables. Then the covariance of $a_1X + b_1$ and $a_2Y + b_2$ can be given by

$$Cov(a_1X + b_1, a_2Y + b_2) = a_1a_2Cov(X, Y).$$

```
Exercise 16
```

```
> xi<-c(-20,-10,0,10,20)
> yi<-c(60,40,35,20,20)
> cor(xi,yi)
```

[1] -0.9534626

From the rule of thumb, we may conclude that X and Y are very strongly related.

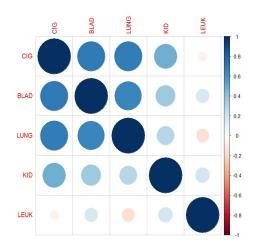
Exercise 17 We first download the datas and save them under csv format ("exo17.csv" in this answer). Next, when we have selected the correct directory, we may use the commands

```
> dat<-read.csv("exo17.csv",header=T,row.names=1)
> cor(dat)
```

CIG BLAD LUNG KID LEUK CIG 1.00000000 0.7036219 0.6974025 0.4873896 -0.06848123 BLAD 0.70362186 1.0000000 0.6585011 0.3588140 0.16215663 LUNG 0.69740250 0.6585011 1.0000000 0.2827431 -0.15158448 KID 0.48738962 0.3588140 0.2827431 1.0000000 0.18871294 -0.06848123 0.1621566 0.1887129 1.00000000 LEUK -0.1515845

The higher the coefficient is in absolute values, the stronger is the relation between two variables. We may note from the previous correlation matrix that bladder and lung cancers are strongly associated with smoking. Help to R, we may provide "graphical" correlations.

```
> library('corrplot') # package corrplot
> corrplot(cor(dat), method = "circle") # plot matrix
```



Exercise 18

```
> cor.test(judge1,judge2,method="pearson")
Pearson's product-moment correlation
```

```
data: judge1 and judge2
t = 2.9448, df = 8, p-value = 0.01857
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
0.1677685 0.9289897
sample estimates:
cor
0.7212121
```

Consequently, there exists a strong rank correlation between judge 1 and judge 2.

Exercise 19

Those tied in the x rankings are given a value of $\frac{2+3}{2} = 2.5$ and those tied in y are allocated $\frac{6+7+8}{3} = 7$. (In general, each tie is given the mean of the places that would have been occupied if a strict order had been produced.) The table, therefore, becomes

Ranks x	1	[2.5]	$\boxed{2.5}$	5	4	6	7	8
Ranks y	1	3	4	2	5	7	7	7

We may apply next the usual commands in R.

```
> ranks_x<-c(1,2.5,2.5,5,4,6,7,8)
> ranks_y<-c(1,3,4,2,5,7,7,7)
> cor.test(ranks_x,ranks_y,method="pearson")
Pearson's product-moment correlation
```

```
data: ranks_x and ranks_y
t = 3.5386, df = 6, p-value = 0.01224
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
0.2794869 0.9667584
```

```
sample estimates:
cor
0.8222252
```

We may conclude that there is positive correlation between rankings.

```
Exercise 20) We use the commands:
```

```
> mat=matrix(c(91,104,235,39,73,48,18,31,161),nrow=3)
> chisq.test(mat,correct=FALSE)

Pearson's Chi-squared test

data: mat
X-squared = 86.023, df = 4, p-value < 2.2e-16</pre>
```

Since the p-value is almost zero we reject the null hypothesis at $\alpha = 0.05$ that financial condition and education level are independent. In another words, financial condition and education level are not independent attributes and we may not reject the evidence of a relationship between the two variables.

Exercise 21 We use the commands:

```
> mat=matrix(c(18,8,7,17),nrow=2)
> chisq.test(mat,correct=FALSE)

Pearson's Chi-squared test
data: mat
X-squared = 8.0128, df = 1, p-value = 0.004645
```

Since the p-value is 0.004645 we reject the null hypothesis at $\alpha = 0.05$ level. Thus, age and cholesterol are not independent attributes.