## Solution to Exercise 1: Introduction to R software: basics

## Key points:

a. $R$ is both a language and an environment of computing
b. Anything and everything can be assigned to an object
c. Object names are case-sensitive
d. A vector is a one-dimensional object of like elements
e. A matrix is a two-dimensional table (another object) of like elements
f. Vectors can be bound to matrices

## Task

In analogy to the calculation of the odds ratio, write an $R$ script e_ex01_rr.r that calculates the relative risk from two proportions (thus not ratio of incidence rates, but ratio of two prevalence proportions) for a study used in Altman's textbook. It has the isolation of Helicobacter pylori as the outcome and the history of an ulcer in the mother as the exposure. We use the following set-up of notations:

| Exposure | Outcome |  |  |
| :--- | :---: | :---: | :---: |
|  | Ill | Healthy | Total |
| Yes | A | B | $\mathrm{A}+\mathrm{B}$ |
| No | C | D | $\mathrm{C}+\mathrm{D}$ |
| Total | $\mathrm{A}+\mathrm{B}$ | $\mathrm{B}+\mathrm{D}$ | $\mathrm{A}+\mathrm{B}+\mathrm{C}+\mathrm{D}$ |

The data provided by Altman in table 7.2 (page 59) are as follows:

| Mother with a <br> history of ulcer | H pylori isolated |  |  |
| :--- | :---: | :---: | :---: |
|  | Yes | No | Total |
| Yes | 6 | 16 | 22 |
| No | 112 | 729 | 841 |
| Total | 118 | 745 | 863 |

The relative risk is calculated by:
$R=(A /(A+B)) /(C /(C+D))$

## The standard error of the $\log _{e} R$ is:

$$
\text { se }=\operatorname{SQRT}(1 / A-1 /(A+B)+1 / C-(1 /(C+D))
$$

## The 95\% CI is thus:

$95 \% C I=\exp \left(\log _{\mathrm{e}} \mathrm{R} \pm 1.96^{*}\right.$ se $)$

## Solution:

The solution for the e_ex01_rr.r is a straight forward modification of the e_ex01_or.r:
\# Complete 2 by 2 table from
\# A, B, C, D and calculate relative risk with Wolf CI
A <- 54
B <- 89
C <- 60
D <- 245

```
AB<- c(A, B, A+B)
CD <- C(C, D, C+D)
ABCD <- rbind(AB, CD)
rtot <- abcd[1,]+abcd[2,]
tab <- rbind(ABCD, rtot)
colnames(tab) <- cbind("Ill", "Healthy", "Total")
rownames(tab) <- rbind("Exp+", "Exp-", "Total")
names(dimnames(tab)) <- c("Exposure", "Status")
se <- sqrt(1/A-1/(A+B)+1/C-1/(C+D))
rr<- (tab[1,1]/tab[1,3])/(tab[2,1]/tab[2,3]); rr
rr.ci.lower <- exp(log(rr)-1.96*se)
rr.ci.upper <- exp(log(rr)+1.96*se)
print(tab)
cat("\nRR:", round(rr, digits=3), "\n95% CI:", round(rr.ci.lower,
digits=3), round(rr.ci.upper, digits=3), "\n")
```

The result is:

|  | I11 | Healthy | Tota1 |
| :--- | ---: | ---: | ---: |
| Exp+ | 6 | 16 | 22 |
| Exp- | 112 | 729 | 841 |
| Total | 118 | 745 | 863 |

RR: 2.048
95\% CI: 1.0134 .14

