

Practical: Effect Estimates Binary Data

Alice Kamau

18 Feb 2016

We will use various R packages to compute these measures such as epicalc, Epi, epiR, and epitools to obtain effect estimates for binary data

Part 1: Load the dataset

We will use the lung capacity dataset to compute effect estimates for binary data (Risk Ratio and Odds Ratio)

Load the *LungCapData.csv* dataset into memory.

```
LungCapData<-read.csv("/Users/akamau/Desktop/Stats forum/Stat training/LungCapData.csv", header=TRUE)
```

Part 2: Explore the dataset

```
attach(LungCapData)
names(LungCapData)
class(Gender)
levels(Gender)
class(Smoke)
levels(Smoke)
```

Explore the relationship between various variables

Load the R packages 'ggplot2', 'lme4', 'descr'.

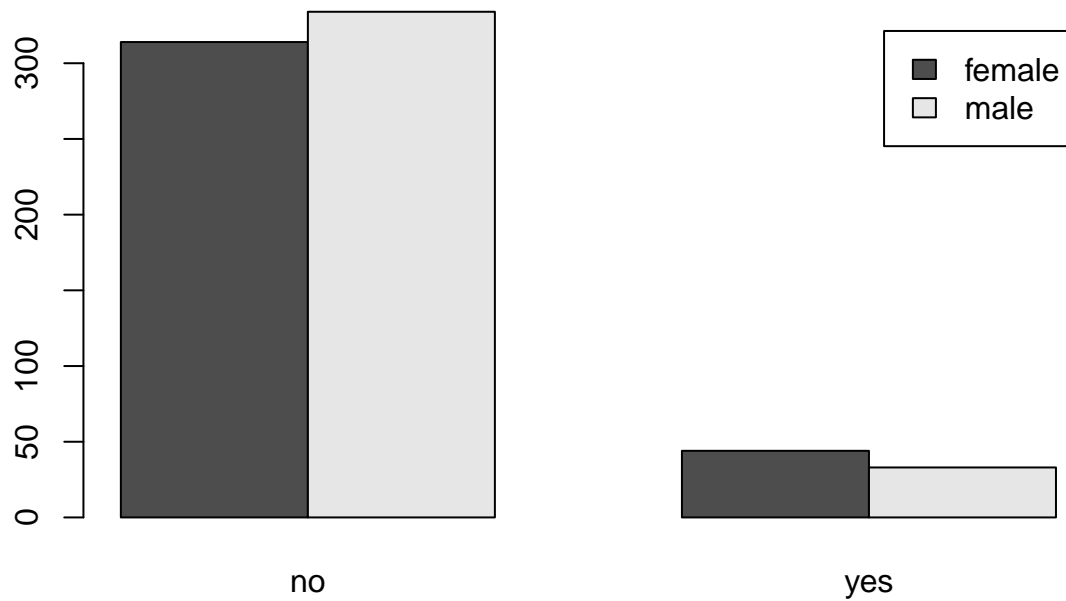
You can use the help function to determine what each package does e.g. ?ggplot

```
library(ggplot2)
library(lme4)
library(descr)
```

```
tab<-table(Gender, Smoke)
tab
```

```
##           Smoke
## Gender    no yes
##  female 314  44
##   male  334  33
```

```
barplot(tab, beside=T, legend=T)
```



```
chisq.test(tab, correct = F)
```

```
##
## Pearson's Chi-squared test
##
## data:  tab
## X-squared = 2.0773, df = 1, p-value = 0.1495
```

```
fisher.test(tab, conf.int=T, conf.level=0.95)
```

```
##
## Fisher's Exact Test for Count Data
##
## data:  tab
## p-value = 0.1845
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
##  0.4233701 1.1659426
## sample estimates:
## odds ratio
##  0.7054345
```

Part 3: Load the R package ‘epitools’, ‘epiR’, ‘abd’ to calculate the RR and OR

```
library(epitools)
library(epiR)
library(abd)
library(epiDisplay)
```

Part 4: Calculate the Risk Ratio and Odds Ratios

Recall the table 'tab'. We need to set it in the standard abcd format. The standard format dictates that the exposure be presented in rows while the outcome be presented in columns

```
tab2<-cbind(tab[,2],tab[,1])
colnames(tab2)<-c("yes","no")
tab2
```

```
##           yes  no
## female   44 314
## male     33 334
```

```
fisher.test(tab2, conf.int=T, conf.level=0.95)
```

```
##
## Fisher's Exact Test for Count Data
##
## data:  tab2
## p-value = 0.1845
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
##  0.8576752 2.3619999
## sample estimates:
## odds ratio
##  1.417566
```

Using epitools package

```
epitab(tab2, method="oddsratio")
```

```
## $tab
##           yes      p0  no      p1 oddsratio      lower      upper      p.value
## female   44 0.5714286 314 0.4845679  1.000000         NA         NA         NA
## male     33 0.4285714 334 0.5154321  1.418259 0.8803057 2.284955 0.1845426
##
## $measure
## [1] "wald"
##
## $conf.level
## [1] 0.95
##
## $pvalue
## [1] "fisher.exact"
```

Using epiR package

```
epi.2by2(tab2, method="cohort.count", conf.level=0.95)
```

```
##           Outcome +   Outcome -   Total   Inc risk *
## Exposed +           44         314     358        12.29
## Exposed -           33         334     367         8.99
## Total              77         648     725        10.62
##           Odds
## Exposed +         0.1401
## Exposed -         0.0988
## Total            0.1188
##
## Point estimates and 95 % CIs:
## -----
## Inc risk ratio (W)                1.37 (0.89, 2.10)
## Odds ratio (W)                   1.42 (0.88, 2.28)
## Attrib risk (W) *                 3.30 (-1.19, 7.79)
## Attrib risk in population (W) *   1.63 (-2.06, 5.32)
## Attrib fraction in exposed (%)    26.84 (-12.15, 52.28)
## Attrib fraction in population (%) 15.34 (-8.10, 33.69)
## -----
## X2 test statistic: 2.077 p-value: 0.15
## W: Wald confidence limits
## * Cases per 100 population units
```

Part 5: Interpretation

The odds of smoking among female is 1.42 times higher than the odds of smoking among male

Refer to the example on the relationship between lung cancer and smoking **

Using the example used in the slides to compute effect estimates using R **

```
smoking <- matrix(c(709, 154, 142, 308), nrow = 2)
rownames(smoking) <- c("smokers", "non-smokers")
colnames(smoking) <- c("lungcancer", "no-lung cancer")
```

```
smoking
```

```
##           lungcancer no-lung cancer
## smokers           709           142
## non-smokers        154           308
```

```
epi.2by2(smoking, method="cohort.count", conf.level=0.95)
```

```
## Outcome + Outcome - Total Inc risk *
## Exposed + 709 142 851 83.3
## Exposed - 154 308 462 33.3
## Total 863 450 1313 65.7
## Odds
## Exposed + 4.99
## Exposed - 0.50
## Total 1.92
##
## Point estimates and 95 % CIs:
## -----
## Inc risk ratio (W) 2.50 (2.19, 2.85)
## Odds ratio (W) 9.99 (7.67, 13.01)
## Attrib risk (W) * 49.98 (45.01, 54.96)
## Attrib risk in population (W) * 32.39 (27.39, 37.40)
## Attrib fraction in exposed (%) 59.99 (54.33, 64.95)
## Attrib fraction in population (%) 49.29 (43.36, 54.59)
## -----
## X2 test statistic: 332.057 p-value: < 0.001
## W: Wald confidence limits
## * Cases per 100 population units
```

```
epitab(smoking, method="oddsratio", rev="both")
```

```
## $tab
## no-lung cancer p0 lungcancer p1 oddsratio
## non-smokers 308 0.6844444 154 0.1784473 1.000000
## smokers 142 0.3155556 709 0.8215527 9.985915
## lower upper p.value
## non-smokers NA NA NA
## smokers 7.66614 13.00766 4.826448e-74
##
## $measure
## [1] "wald"
##
## $conf.level
## [1] 0.95
##
## $pvalue
## [1] "fisher.exact"
```

Part 6: Assignment

1. Evaluate the birthweight2 data
2. Look at the association between LBW and gestational weeks. Divide gestational week in two groups i.e. group1 < 35 weeks and group2 >= 35 weeks. Then calculate the odds ratio between LBW and gestational weeks.
3. Check the Odds ratio for the association between LBW and gender
4. Make Conclusion