

Part 6: Assignment

1. Evaluate the birthweight2 data

```
setwd("/Users/akamau/Work/OneDrive - Kemri Wellcome Trust/Stats forum/Stat training")
bw.data<-read.csv("Data/birthweight2.csv", header=TRUE)
```

```
names(bw.data)
class(bw.data$matage)
class(bw.data$ht)
class(bw.data$gestwks)
```

```
library(epitools)
library(epiR)
```

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.

```
mytable1 <- table(bw.data$lbw, bw.data$gestwks)
mytable1
```

```
##
##           25  26  28  29  30  31  32  33  34  35  36  37  38  39  40
## Normal 2500+  0   0   0   0   0   0   0   0   5   8  11  30  87 167 146
## Weight<2500  1   1   3   1   3   5   7   6   7   9   6  11  14   3   3
##
##           41  42
## Normal 2500+ 87  20
## Weight<2500  0   0
```

```
bw.data$gestwks1 <- NULL
bw.data$gestwks1[bw.data$gestwks < 35] <- 1
bw.data$gestwks1[bw.data$gestwks > 34] <- 2
bw.data$gestwks1 <- factor(bw.data$gestwks1 , labels = c("< 35 weeks" , ">= 35 weeks"))
```

```
mytable2 <- table(bw.data$gestwks1, bw.data$lbw)
mytable2
```

```
##
##           Normal 2500+ Weight<2500
## < 35 weeks           5           34
## >= 35 weeks          556          46
```

```
chisq.test(mytable1)
```

```
## Warning in chisq.test(mytable1): Chi-squared approximation may be incorrect
```

```
##
## Pearson's Chi-squared test
##
## data:  mytable1
## X-squared = 301.99, df = 16, p-value < 2.2e-16
```

```
chisq.test(mytable2)
```

```
## Warning in chisq.test(mytable2): Chi-squared approximation may be incorrect
```

```
##
```

```
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: mytable2
## X-squared = 204.92, df = 1, p-value < 2.2e-16
fisher.test(mytable2)

##
## Fisher's Exact Test for Count Data
##
## data: mytable2
## p-value < 2.2e-16
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.003574824 0.033832466
## sample estimates:
## odds ratio
## 0.01236775
mytable3<-cbind(mytable2[,2],mytable2[,1])
colnames(mytable3)<-c("Weight<2500", "Normal 2500+")
mytable3

##           Weight<2500 Normal 2500+
## < 35 weeks           34           5
## >= 35 weeks          46          556
mytable2

##           Normal 2500+ Weight<2500
## < 35 weeks           5           34
## >= 35 weeks          556          46
epitab(mytable3, method="oddsratio", rev= "both")

## Warning in chisq.test(xx, correct = correction): Chi-squared approximation
## may be incorrect

## $tab
##           Normal 2500+           p0 Weight<2500           p1 oddsratio           lower
## >= 35 weeks          556 0.991087344           46 0.575           1.0000           NA
## < 35 weeks           5 0.008912656           34 0.425           82.1913 30.67057
##           upper           p.value
## >= 35 weeks           NA           NA
## < 35 weeks 220.2571 4.58709e-29
##
## $measure
## [1] "wald"
##
## $conf.level
## [1] 0.95
##
## $pvalue
## [1] "fisher.exact"

Using epiR package
```

```
epi.2by2(mytable3, method="cohort.count", conf.level=0.95)
```

```
##           Outcome +      Outcome -      Total      Inc risk *
## Exposed +           34           5           39           87.18
## Exposed -           46          556          602           7.64
## Total              80          561          641          12.48
##
##           Odds
## Exposed +           6.8000
## Exposed -           0.0827
## Total              0.1426
##
## Point estimates and 95% CIs:
## -----
## Inc risk ratio              11.41 (8.43, 15.44)
## Odds ratio                  82.19 (30.67, 220.26)
## Attrib risk *              79.54 (68.83, 90.24)
## Attrib risk in population * 4.84 (1.52, 8.16)
## Attrib fraction in exposed (%) 91.24 (88.14, 93.52)
## Attrib fraction in population (%) 38.77 (27.10, 48.58)
## -----
## Test that odds ratio = 1: chi2(1) = 212.138 Pr>chi2 = < 0.001
## Wald confidence limits
## CI: confidence interval
## * Outcomes per 100 population units
```

3. Check the Odds ratio for the association between LBW and gender

```
mytable4 <- table( bw.data$sex, bw.data$lbw)
mytable4
```

```
##
##           Normal 2500+ Weight<2500
## Female           270           45
## Male            291           35
```

```
mytable5<-cbind(mytable4[,2],mytable4[,1])
colnames(mytable5)<-c("Weight<2500", "Normal 2500+")
mytable5
```

```
##           Weight<2500 Normal 2500+
## Female           45           270
## Male            35           291
```

```
epitab(mytable5, method="oddsratio", rev = "both")
```

```
## $tab
##           Normal 2500+      p0 Weight<2500      p1 oddsratio      lower
## Male           291 0.5187166           35 0.4375 1.000000      NA
## Female          270 0.4812834           45 0.5625 1.385714 0.8645583
##           upper  p.value
## Male           NA      NA
## Female 2.221023 0.1895456
##
## $measure
## [1] "wald"
##
```

```
## $conf.level
## [1] 0.95
##
## $pvalue
## [1] "fisher.exact"
```

Using epiR package

```
epi.2by2(mytable5, method="cohort.count", conf.level=0.95)
```

```
##           Outcome +      Outcome -      Total      Inc risk *
## Exposed +           45          270          315           14.3
## Exposed -           35          291          326           10.7
## Total              80          561          641           12.5
##
##           Odds
## Exposed +       0.167
## Exposed -       0.120
## Total          0.143
##
## Point estimates and 95% CIs:
## -----
## Inc risk ratio                1.33 (0.88, 2.01)
## Odds ratio                    1.39 (0.86, 2.22)
## Attrib risk *                 3.55 (-1.57, 8.67)
## Attrib risk in population *   1.74 (-2.48, 5.97)
## Attrib fraction in exposed (%) 24.85 (-13.66, 50.31)
## Attrib fraction in population (%) 13.98 (-8.59, 31.85)
## -----
## Test that odds ratio = 1: chi2(1) = 1.848 Pr>chi2 = 0.174
## Wald confidence limits
## CI: confidence interval
## * Outcomes per 100 population units
```