

Biostatistics (MATH11230)

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Below I show a possible way of how to organise the stratified data in an array and pass this information to the `epi.2by2` function from the `epiR` package. This function, among other things, returns the Mantel-Haenszel adjusted OR and RR estimates and corresponding CIs, the stratum-specific estimates and corresponding CIs, and the results of the homogeneity test (to check if there is effect modification), There is a warning, that I do not know where it is coming from, but the results the function is returning are correct.

```
require(epiR)
```

```
stratified_data <- array(c(36, 50, 14, 50,
                          24,10,126,90),
                        dim = c(2,2,2),
                        list(area = c("Rural", "Urban"),
                             antibodies = c("Yes", "No"),
                             gender = c("Males", "Females"))
                        )
```

```
stratified_data
```

```
## , , gender = Males
##
##      antibodies
## area   Yes No
## Rural  36 14
## Urban  50 50
##
## , , gender = Females
##
##      antibodies
## area   Yes  No
## Rural  24 126
## Urban  10  90
```

```
res <- epi.2by2(stratified_data, units = 1, method = "cohort.count")
res
```

```
##           Outcome +      Outcome -      Total      Inc risk *
## Exposed +           60          140        200      0.30 (0.24 to 0.37)
## Exposed -           60          140        200      0.30 (0.24 to 0.37)
## Total             120          280        400      0.30 (0.26 to 0.35)
##
##
## Point estimates and 95% CIs:
## -----
## Inc risk ratio (crude)                1.00 (0.74, 1.35)
## Inc risk ratio (M-H)                 1.48 (1.13, 1.94)
```

```

## Inc risk ratio (crude:M-H) 0.67
## Inc odds ratio (crude) 1.00 (0.65, 1.53)
## Inc odds ratio (M-H) 2.13 (1.24, 3.63)
## Inc odds ratio (crude:M-H) 0.47
## Attrib risk in the exposed (crude) * 0.00 (-0.09, 0.09)
## Attrib risk in the exposed (M-H) * 0.12 (0.01, 0.22)
## Attrib risk (crude:M-H) 0.00
## -----
## M-H test of homogeneity of IRRs: chi2(1) = 0.078 Pr>chi2 = 0.780
## M-H test of homogeneity of ORs: chi2(1) = 0.580 Pr>chi2 = 0.446
## Test that M-H adjusted OR = 1: chi2(1) = 7.819 Pr>chi2 = 0.003
## Wald confidence limits
## M-H: Mantel-Haenszel; CI: confidence interval
## * Outcomes per population unit
names(res)

## [1] "method" "n.strata" "digits" "conf.level"
## [5] "interp" "units" "tab" "massoc.summary"
## [9] "massoc.interp" "massoc.detail"
res$massoc.detail$OR.strata.wald

## est lower upper
## 1 2.571429 1.2376221 5.342701
## 2 1.714286 0.7813461 3.761170

```