

# epidemiology measures

SURV686

Example from class:

Table: Data for Example 1 and 2

Smoker	Stroke		Total
	Yes	No	
Yes	171	3,264	3,435
No	117	4,320	4,437
Total	288	7,584	7,872

Please calculate incidence among smokers, non-smokers, and the relative risk and odds ratio for smokers compared to non-smokers.

We enter the data as separate vectors to facilitate calculations.

```
library(tidyverse)
library(epiR)
a <- 171
b <- 3264
c <- 117
d <- 4320

# total
t <- sum(a, b, c, d)
```

Calculate incidence among smokers and non-smokers using;

$$I_{Exposed} = \frac{A}{A+B}$$

$$I_{Un-exposed} = \frac{C}{C+D}$$

```
ir_exp <- a / (a+b) ; ir_exp
```

```
[1] 0.04978166
```

```
ir_unexp <- c / (c+d) ; ir_unexp
```

```
[1] 0.02636917
```

- About 5% of smokers had a stroke, almost twice as many as the non-smokers.

**Calculate relative risk:**

$$RR = \frac{I_{Exposed}}{I_{Un-exposed}}$$

\$\$

\$\$

```
RR <- round(ir_exp / ir_unexp, 4)
```

- Smokers are 1.88 times more at risk of having a stroke compared to non-smokers.

**Calculate Odds Ratio**

$$\hat{OR} = \frac{AD}{BC}$$

```
OR <- (a*d) / (b*c)
```

- Our odds ratio is close to our relative risk estimate.

**Calculate Variances for OR and RR, to estimate 95% CI.**

$$Var(\ln \hat{R}) = \frac{b}{a(a+b)} + \frac{d}{c(c+d)}$$

$$Var(\ln \hat{O}) = \frac{1}{a} + \frac{1}{b} + \frac{1}{c} + \frac{1}{d}$$

```
var_RR <- (b / (a*(a+b))) + (d / (c*(c+d))) ; var_RR
```

```
[1] 0.01387846
```

```
var_OR <- (1/a) + (1/b) + (1/c) + (1/d) ; var_OR
```

```
[1] 0.01493282
```

```
lower <- round(exp(log(RR) - 1.96 * sqrt(var_RR)) , 4)
upper <- round(exp(log(RR) + 1.96 * sqrt(var_RR)) , 4)

print(str_c("Printing 95% CI for relative risk . . . ", RR,
            " [" , lower, " , " , upper, "]" ))
```

```
[1] "Printing 95% CI for relative risk . . . 1.8879 [1.4986, 2.3783]"
```

```
lower <- round(exp(log(OR) - 1.96 * sqrt(var_OR)) , 4)
upper <- round(exp(log(OR) + 1.96 * sqrt(var_OR)) , 4)

print(str_c("Printing 95% CI for odds ratio. . . ", RR,
            " [" , lower, " , " , upper, "]" ))
```

```
[1] "Printing 95% CI for odds ratio. . . 1.8879 [1.5224, 2.4579]"
```

- The intervals do not contain 1, so they are statistically significant and reject the Null hypothesis that the odds are equal across smokers and non-smokers.

**Prospective study:**

```

a <- 23
b <- 125
c <- 13
d <- 150
t <- sum(a, b, c, d)
# check attrition
attrition <- sum(a/t, b/t, c/t, d/t) ; attrition

```

```
[1] 1
```

**Calculate relative risk, and odds ratio from formula from before.**

$$\hat{R} = \frac{a(c+d)}{c(a+b)}$$

```
R_hat <- (a*(c+d)) / (c*(a+b)) ; R_hat
```

```
[1] 1.948545
```

```
OR <- (a*d)/(b*c) ; OR
```

```
[1] 2.123077
```

- The odds ratio is slightly larger than the relative risk.

Calculate the attributed risk percent in the exposed group.

$$ARP_{Exposed} = \frac{\hat{R} - 1}{\hat{R}}$$

```
ARP_exposed <- (R_hat - 1) / R_hat ; ARP_exposed # fraction
```

```
[1] 0.4867965
```

- The amount of disease incidence is 49% which can be attributed to an exposure in a prospective study.
- This is also referred to ARP, attributable risk percent when converting to a percent.

The attributable risk (ratio) of the exposed is calculated as incidence of the exposed - incidence of the non-exposed.

```
ir_exp <- a / (a+b)
ir_unexp <- c / (c+d)
AR_exposed <- ir_exp - ir_unexp ; AR_exposed
```

```
[1] 0.0756508
```

To calculate 95% CI we would need to calculate the standard error as:

$$SE_{AR} = \sqrt{\frac{a+c}{N}(1 - \frac{a+c}{N})(\frac{1}{a+b} + \frac{1}{c+d})}$$

```
SE_AR <- sqrt(
  ((a+c)/t)*(1-((a+c)/t)) * ( (1/(a+b)) + (1/(c+d)) )
)

(AR_exposed - 1.97*SE_AR) * 100 # lower
```

```
[1] 0.4089381
```

```
(AR_exposed + 1.97*SE_AR) * 100 # Upper
```

```
[1] 14.72122
```

When we use the `epiR::epi.2by2()` we can see these results under “Attrib prev in the exposed.”

### Calculate the attributable risk in the population

$$\hat{A}_{pop} = \frac{ad - bc}{(a+c)(c+d)}$$

```
# reduction in incidence in the population that would
# occur in the absence of the risk factor
A_hat_pop <- (a*d - b*c) / ((a+c)*(c+d)) ; A_hat_pop # fraction
```

```
[1] 0.3110089
```

- This is the reduction in incidence of 31% if the whole population were unexposed, comparing with actual exposure.
- This is known as PARP, population attributable risk when converting to percent.

**Calculate the variance of the attributable risk in the population.**

$$V(\ln(1 - \hat{A}_{pop}) = \frac{b + \hat{A}_{pop}(a + d)}{tc}$$

```
V <- (b + A_hat_pop* (a+d)) / (t*c) ; V
```

```
[1] 0.04422571
```

**In a retrospective study we use the OR instead of or relative risk, and calculate the attributable risk in the exposed group and population.**

```
R_hat <- (a*d)/(b*c) ; R_hat
```

```
[1] 2.123077
```

```
A_exposed <- (R_hat - 1) / R_hat ; A_exposed # fraction
```

```
[1] 0.5289855
```

**For the population we change the denominator.**

$$\hat{A}_{pop} = \frac{ad - bc}{d(a + c)}$$

```
A_hat_pop <- (a*d - b*c) / ( d * (a+c)) ; A_hat_pop # fraction
```

```
[1] 0.337963
```

**The variance is also computed differently**

$$V(\ln(1 - \hat{A}_{pop}) = \frac{a}{c(a + c)} + \frac{b}{d(b + d)}$$

```
V <- (a / (c * (a + c))) +  
      (b / (d*(b+d))); V
```

```
[1] 0.0521756
```

**We can now calculate 95% CI.**

$$LCL = 1 - \exp(\ln(1 - \hat{A}_{pop}) + 1.96\sqrt{V(\ln(1 - \hat{A}_{pop}))})$$

$$UCL = 1 - \exp(\ln(1 - \hat{A}_{pop}) - 1.96\sqrt{V(\ln(1 - \hat{A}_{pop}))})$$

```
LCL <- 1 - exp( A_hat_pop + 1.96 * sqrt(V) ) ; LCL
```

```
[1] -1.193867
```

```
UCL <- 1 - exp( A_hat_pop - 1.96 * sqrt(V) ) ; UCL
```

```
[1] 0.1039328
```

**We can also use the r package epiR**

```
# create table for later  
bp <- matrix(c(a, b, c, d), ncol = 2, byrow = TRUE)  
epi.2by2(bp, method="cross.sectional")
```

	Outcome +	Outcome -	Total	Prev risk *
Exposed +	23	125	148	15.54 (10.11 to 22.40)
Exposed -	13	150	163	7.98 (4.31 to 13.25)
Total	36	275	311	11.58 (8.24 to 15.66)

Point estimates and 95% CIs:

---

Prev risk ratio	1.95 (1.02, 3.71)
Prev odds ratio	2.12 (1.03, 4.36)
Attrib prev in the exposed *	7.57 (0.40, 14.73)
Attrib fraction in the exposed (%)	48.68 (2.41, 73.01)
Attrib prev in the population *	3.60 (-1.87, 9.07)
Attrib fraction in the population (%)	31.10 (-4.04, 54.37)

---

Uncorrected chi2 test that OR = 1: chi2(1) = 4.337 Pr>chi2 = 0.037

Fisher exact test that OR = 1: Pr>chi2 = 0.050

Wald confidence limits

CI: confidence interval

\* Outcomes per 100 population units