Statistics 305/605: Introduction to Biostatistical Methods for Health Sciences

R Demo for Chapter 15, part 3: McNemar's Test

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2018-10-22

McNemar's Test for the Smoking Data

▶ We have r = 97 and s = 76. The test statistic with continuity correction is

$$\frac{(|97-76|-1)^2}{97+76} = 2.31$$

and the corresponding p-value is 0.128:

```
X2 <- (abs(97-76)-1)^2/(97+76)
X2

## [1] 2.312139

pval<-pchisq(X2,df=1,lower.tail=FALSE)
pval

## [1] 0.1283673</pre>
```

McNemar's test on T1D Data

Read in the data and cross tabulate:

```
uu<-url("http://people.stat.sfu.ca/~jgraham/Teaching/S305_17/Data/tdt.csv")
tdtex <- read.csv(uu)
head(tdtex,n=8)</pre>
```

```
tt <- table(tdtex)
tt</pre>
```

```
## untransmitted
## transmitted 1 X
## 1 0 78
## X 46 0
```

▶ Now apply McNemar's test *without* the continuity correction to the cross-tabulated data:

```
mcnemar.test(tt,correct=FALSE)
```

```
##
## McNemar's Chi-squared test
##
## data: tt
## McNemar's chi-squared = 8.2581, df = 1, p-value = 0.004057
```

▶ Apply McNemar's test *with* the (default) continuity correction:

```
mcnemar.test(tt)
```

```
##
## McNemar's Chi-squared test with continuity correction
##
## data: tt
## McNemar's chi-squared = 7.75, df = 1, p-value = 0.005371
```