

HW9 KEY

28 points total, 2 points per problem part unless otherwise noted.

Q1 Toxic Tomatoes

```
#A
#New
5/50
```

```
## [1] 0.1
```

```
#Old
9/50
```

```
## [1] 0.18
```

```
#B
prop.test(c(5, 9), c(50, 50))
```

```
##
## 2-sample test for equality of proportions with continuity
## correction
##
## data: c(5, 9) out of c(50, 50)
## X-squared = 0.74751, df = 1, p-value = 0.3873
## alternative hypothesis: two.sided
## 95 percent confidence interval:
## -0.23510963 0.07510963
## sample estimates:
## prop 1 prop 2
## 0.10 0.18
```

```
#C
Sludge <- matrix(c(5, 45, 9, 41), byrow = TRUE, nrow = 2)
colnames(Sludge) <- c("Toxic", "NonToxic")
rownames(Sludge) <- c("New", "Old")
Sludge
```

```
##      Toxic NonToxic
## New      5      45
## Old      9      41
```

```
chisq.test(Sludge)
```

```
##
## Pearson's Chi-squared test with Yates' continuity correction
##
## data: Sludge
## X-squared = 0.74751, df = 1, p-value = 0.3873
```

```
#D
chisq.test(Sludge)$expected
```

```
##      Toxic NonToxic
```

```
## New      7      43
## Old      7      43
```

```
#E
```

```
fisher.test(Sludge)
```

```
##
## Fisher's Exact Test for Count Data
##
## data: Sludge
## p-value = 0.3881
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
##  0.1235385 1.8596694
## sample estimates:
## odds ratio
##  0.5095856
```

F. Since all expected counts > 5, chi-square test is fine.

```
#G
```

```
prop.test(c(5, 9), c(50, 50), alternative = "less")
```

```
##
## 2-sample test for equality of proportions with continuity
## correction
##
## data: c(5, 9) out of c(50, 50)
## X-squared = 0.74751, df = 1, p-value = 0.1936
## alternative hypothesis: less
## 95 percent confidence interval:
## -1.00000000 0.05338758
## sample estimates:
## prop 1 prop 2
##  0.10  0.18
```

Q2 Anesthesia

```
#A
```

```
#Drug1
```

```
(12+10)/45
```

```
## [1] 0.4888889
```

```
#Drug2
```

```
(12+9)/45
```

```
## [1] 0.4666667
```

```
#B (4 pts)
```

```
Anes <- matrix(c(12, 10, 9, 14), byrow=TRUE, nrow=2)
```

```
Anes
```

```
##      [,1] [,2]
## [1,]  12  10
## [2,]   9  14
```

```
mcnemar.test(Anes)
```

```
##  
## McNemar's Chi-squared test with continuity correction  
##  
## data: Anes  
## McNemar's chi-squared = 0, df = 1, p-value = 1
```

Q3 Case Control Study

```
library(epitools)  
Birds <- matrix(c(328, 141, 101, 98), byrow = TRUE, nrow = 2)  
colnames(Birds) <- c("Control", "Cancer")  
rownames(Birds) <- c("NoBird", "YesBird")  
oddsratio(Birds, method="wald")
```

```
## $data  
##      Control Cancer Total  
## NoBird      328     141   469  
## YesBird      101      98   199  
## Total        429     239   668  
##  
## $measure  
##              NA  
## odds ratio with 95% C.I. estimate lower upper  
##              NoBird 1.000000      NA      NA  
##              YesBird 2.257145 1.60518 3.173915  
##  
## $p.value  
##              NA  
## two-sided midp.exact fisher.exact chi.square  
## NoBird      NA      NA      NA  
## YesBird 3.052348e-06 3.938413e-06 2.243712e-06  
##  
## $correction  
## [1] FALSE  
##  
## attr(,"method")  
## [1] "Unconditional MLE & normal approximation (Wald) CI"
```

3A. OR = 2.26, the YesBird group has higher odds of lung cancer.

3B. (4 pts) 95%CI = (1.605, 3.174)

Since the CI does NOT include one, we can conclude that there is a relationship between bird ownership and lung cancer.

3C. Chi-square p-value < 0.001.

Reject H₀; conclude there is a relationship between bird ownership and lung cancer.