R Handout - Chi-Square Tests

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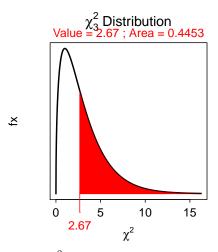
First Commands

> library(NCStats)

Chi-Square Distribution Calculations

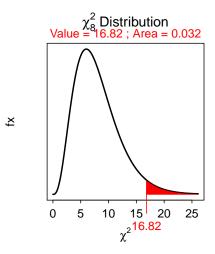
An example of computing the p-value $\chi^2=2.67$ and df=3.

```
> ( distrib(2.67,distrib="chisq",df=3,lower.tail=FALSE) )
[1] 0.4453
```



An example of computing the p-value if $\chi^2=16.82$ and df=8.

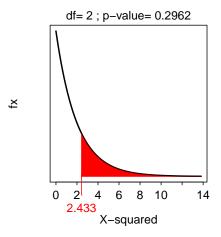
```
> ( distrib(16.82,distrib="chisq",df=8,lower.tail=FALSE) )
[1] 0.03204
```



Goodness-of-Fit Test

Four-o'clocks (*Mirabilis jalapa*) are plants native to tropical America. Individual four-o'clocks can have red, white, or pink flowers. Flower color in this species is thought to controlled by a single gene locus with two alleles experssing incomplete dominance, so that heterozygotes are pink-flower, while homozygotes for one allele are white-flowered and homozygotes for the other allele are red-flowered. According to Mendelian genetic principles, self-pollination of pink-flowered plants should produce progeny that have red, pink, and white flowers in a 1:2:1 ratio. A horticulturist self-pollinates several pink-flowered plants and produces 240 progeny with 55 that are red-flowered, 132 that are pink-flowered, and 53 that are white-flowered. Use the results to determine, at the 5% level, if the theoretical 1:2:1 ratio is upheld with these data¹.

```
> obs <- c(red=55,pink=132,white=53)</pre>
> exp.p <- c(red=1/4, pink=2/4, white=1/4)
> chi1 <- chisq.test(obs,p=exp.p,rescale=TRUE,correct=FALSE)</pre>
> chi1$expected
  red pink white
        120
               60
   60
> chi1$observed
  red pink white
        132
   55
               53
> chi1
Chi-squared test for given probabilities with obs
X-squared = 2.433, df = 2, p-value = 0.2962
> plot(chi1)
```



¹This is example 11.1 from Glover, T. and K. Mitchell. 2008. An Introduction to Biostatistics, 2nd edition. Waveland Press.

Chi-Square Test

From Summarized Observed Table

Do the dominant food items in lake trout and chinook salmon differ at the 5% level? The summarized observed table was given in class.

```
> freq <- c(32,10,8,18,18,4)
> ( obs <- matrix(freq,nrow=2,byrow=TRUE) )</pre>
    [,1] [,2] [,3]
[1,] 32 10 8
[2,]
     18
          18
> rownames(obs) <- c("lake trout", "chinook salmon")</pre>
> colnames(obs) <- c("lake herring", "smelt", "mysis")</pre>
> obs
              lake herring smelt mysis
lake trout
                        32 10 8
                      18
chinook salmon
                           18
> chi1 <- chisq.test(obs,correct=FALSE)</pre>
> chi1$expected
              lake herring smelt mysis
lake trout
               27.78 15.56 6.667
                     22.22 12.44 5.333
chinook salmon
> chi1$expected >= 5
             lake herring smelt mysis
lake trout
               TRUE TRUE TRUE
chinook salmon
                    TRUE TRUE TRUE
> chi1
Pearson's Chi-squared test with obs
X-squared = 6.508, df = 2, p-value = 0.03861
> chi1$residuals
              lake herring smelt mysis
lake trout
                  0.8011 -1.409 0.5164
chinook salmon
                   -0.8957 1.575 -0.5774
> percTable(obs,margin=1,digits=1)
              lake herring smelt mysis Sum
               64 20 16 100
lake trout
chinook salmon
                      45 45
                                  10 100
```

From Raw Data

On the GSS, respondents were asked to state their opinion on how true the following statement was "All radioactivity is made by humans." Respondents were also categorized by their highest educational degree. Use the results in the SciTest1.txt data file to determine, at the 5% level, if the response to the question differs among levels of education

```
> setwd("C:/aaaWork/Class Materials/MTH107/Lecture/HOs")
> ST1 <- read.table("SciTest1.txt",head=TRUE)
> str(ST1)
'data.frame': 2549 obs. of 2 variables:
$ degree : Factor w/ 5 levels "bach", "grad",..: 5 5 5 5 5 5 5 5 5 5 5 ...
$ scitest: Factor w/ 4 levels "def.not","def.true",..: 2 2 2 2 2 2 2 2 2 ...
> levels(ST1$degree)
[1] "bach" "grad" "hs"
                           "jc"
                                   "lt.hs"
> ST1$fdegree <- factor(ST1$degree,levels=c("lt.hs","hs","jc","bach","grad"))</pre>
> levels(ST1$scitest)
[1] "def.not"
              "def.true" "prob.not" "prob.true"
> ST1$fscitest <- factor(ST1$scitest,levels=c("def.not","prob.not","prob.true","def.true"))
> ( freq.tbl <- xtabs(~fdegree+fscitest,data=ST1) )</pre>
      fscitest
fdegree def.not prob.not prob.true def.true
 lt.hs
           52
                    112
                              155
                                        70
 hs
           366
                    451
                              437
                                       114
            60
                                         9
                     44
                               36
 jс
 bach
           214
                    135
                               78
                                        12
           123
                     57
                               18
                                         6
 grad
> ST1.chi <- chisq.test(freq.tbl,correct=FALSE)
> ST1.chi$expected
      fscitest
fdegree def.not prob.not prob.true def.true
 lt.hs 124.38 121.93 110.49
                                    32.20
        437.40 428.81
 hs
                         388.56
                                   113.24
         47.64 46.70
                           42.32
                                     12.33
 jс
 bach
       140.36 137.61 124.69
                                   36.34
         65.23 63.95
                           57.94
                                   16.89
 grad
> ST1.chi$expected >= 5
      fscitest
fdegree def.not prob.not prob.true def.true
 lt.hs
          TRUE
                  TRUE
                             TRUE
                                      TRUE
 hs
          TRUE
                   TRUE
                             TRUE
                                      TRUE
 jс
          TRUE
                   TRUE
                             TRUE
                                      TRUE
 bach
          TRUE
                   TRUE
                             TRUE
                                      TRUE
          TRUE
                   TRUE
                             TRUE
                                      TRUE
 grad
> all(chi1$expected >= 5)
[1] TRUE
> ST1.chi
```

Pearson's Chi-squared test with freq.tbl X-squared = 288.2, df = 12, p-value < 2.2e-16

> ST1.chi\$residuals

fscitest

> percTable(freq.tbl,margin=1,digits=1)

fscitest

| fdegree | <pre>def.not</pre> | <pre>prob.not</pre> | <pre>prob.true</pre> | <pre>def.true</pre> | Sum |
|---------|--------------------|---------------------|----------------------|---------------------|-------|
| lt.hs | 13.4 | 28.8 | 39.8 | 18.0 | 100.0 |
| hs | 26.8 | 33.0 | 31.9 | 8.3 | 100.0 |
| jс | 40.3 | 29.5 | 24.2 | 6.0 | 100.0 |
| bach | 48.7 | 30.8 | 17.8 | 2.7 | 100.0 |
| grad | 60.3 | 27.9 | 8.8 | 2.9 | 99.9 |