

In this **R** program, we run a χ^2 test for homogeneity. The data is from the article by Rosenman and Hertzberg *et al.*, “Chronic Beryllium Disease and Sensitization at a Beryllium Processing Facility,” in *Environmental Health Perspectives*, 2005; as quoted by Navidi, *Statistics for Engineers and Scientists*, 2nd ed., Mc Graw Hill, New York, 2008. The authors discuss the effects of exposure to beryllium in a cohort of workers. Workers were categorized by their duration of exposure (in years) and by their disease status (chronic beryllium disease, sensitization to beryllium, or no disease). Can we conclude that the proportion of workers in the various disease categories differ among exposure levels? The results were as follows

Disease Status	Duration of Exposure		
	< 1Yr.	1 ≤ Yr. < 5	5 ≤ Yrs.
Diseased	10	8	23
Sensitized	9	19	11
Normal	70	136	206

We run our chi-squared test of Homogeneity. The null hypothesis is that the proportion of durations is the same for each level of disease. The test statistic

$$\chi^2 = \sum_{i=1}^3 \sum_{j=1}^3 \frac{(n_{ij} - \hat{e}_{ij})^2}{\hat{e}_{ij}} = 10.8286$$

where the estimated expected frequency in each cell is given by

$$\hat{e}_{ij} = \frac{n_{i.} \cdot n_{.j}}{n_{..}}$$

Since the $\alpha = .05$ critical value is $\chi_{\alpha, (I-1)(J-1)}^2 = \chi_{0.05, 4}^2 = 9.487729$, we reject the null hypothesis: there is significant evidence to indicate that workers having different disease levels received different levels of exposure. The p -value was 0.02856.

R Session:

```
R version 2.13.1 (2011-07-08)
Copyright (C) 2011 The R Foundation for Statistical Computing
ISBN 3-900051-07-0
Platform: i386-apple-darwin9.8.0/i386 (32-bit)
```

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Natural language support but running in an English locale
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Type 'demo()' for some demos, 'help()' for on-line help, or
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Type 'q()' to quit R.

[R.app GUI 1.41 (5874) i386-apple-darwin9.8.0]

[History restored from /Users/andrejstreibergs/.Rapp.history]

```
> ##### ENTER THE CONTINGENCY TABLE AS MATRIX #####

> exposure.disease=matrix(c(10,8,23,9,19,11,70,136,206),nrow=3,byrow=T)
> rownames(exposure.disease)=c("Diseased","Sensitized","Normal")
> colnames(exposure.disease)=c("<1 Yr","1 Yrs < 5","5 Yrs")
> names(dimnames(exposure.disease))=c("Disease Status","Duration of Exposure")
> exposure.disease
              Duration of Exposure
Disease Status <1 Yr 1 Yrs < 5 5 Yrs
      Diseased      10      8      23
      Sensitized     9      19     11
      Normal       70     136     206
>
> ##### AS A DATA FRAME OF COUNTS #####
> as.data.frame(as.table(exposure.disease))
  Disease.Status Duration.of.Exposure Freq
1      Diseased              <1 Yr    10
2      Sensitized             <1 Yr     9
3       Normal              <1 Yr    70
4      Diseased             1 Yrs < 5     8
5      Sensitized            1 Yrs < 5    19
6       Normal             1 Yrs < 5   136
7      Diseased             5 Yrs     23
8      Sensitized            5 Yrs     11
9       Normal             5 Yrs    206
>
> ##### RUN THE CHI SQUARED TEST OF HOMOGENEITY #####
> X2<-chisq.test(exposure.disease)
> X2
```

Pearson's Chi-squared test

```
data: exposure.disease
X-squared = 10.8286, df = 4, p-value = 0.02856
```

```

> ##### CHI-SQUARED TEST "BY HAND" #####

> ##### ROW SUMS, COLUMN SUMS, GRAND TOTAL #####
> rowsum = margin.table(exposure.disease,1); rowsum
Disease Status
  Diseased Sensitized      Normal
        41         39         412
> colsum = margin.table(exposure.disease,2); colsum
Duration of Exposure
  <1 Yr 1 Yrs < 5      5 Yrs
        89         163         240
> n=sum(exposure.disease); n
[1] 492

> ##### TABLE OF EXPECTED FREQUENCIES #####

> outer(rowsum, colsum)/n
      Duration of Exposure
Disease Status  <1 Yr 1 Yrs < 5      5 Yrs
  Diseased      7.416667    13.58333    20.00000
  Sensitized    7.054878    12.92073    19.02439
  Normal       74.528455   136.49593   200.97561
> X2$expected
      Duration of Exposure
Disease Status  <1 Yr 1 Yrs < 5      5 Yrs
  Diseased      7.416667    13.58333    20.00000
  Sensitized    7.054878    12.92073    19.02439
  Normal       74.528455   136.49593   200.97561

> X2$observed
      Duration of Exposure
Disease Status <1 Yr 1 Yrs < 5      5 Yrs
  Diseased      10          8          23
  Sensitized     9          19         11
  Normal        70         136         206

> ##### CONTRIBUTION TO CHI-SQ FROM EACH CELL ###

> E <- X2$expected; O <- X2$observed;
> (E-O)^2/E
      Duration of Exposure
Disease Status  <1 Yr 1 Yrs < 5      5 Yrs
  Diseased      0.8998127  2.294989775  0.4500000
  Sensitized    0.5362955  2.860325856  3.3846467
  Normal        0.2751554  0.001801896  0.1256098

```

```
> ##### CHI-SQ SUM #####
> ch2=sum((O-E)^2/E); ch2
[1] 10.82864

> ##### CRITICAL VALUE FOR ALPHA=.05, (I-1)(J-1)=4 #####
> qchisq(.05,4,lower.tail=F)
[1] 9.487729
> ##### SO REJECT H0 #####

> ##### P-VALUE #####
> pchisq(ch2,4,lower.tail=F)
[1] 0.02855892

>
```