Chisq.test explorations

The question is: can chisq.test() be used to compare the frequency distributions of a table of data, and a subset of that table? More specifically, we have a data set that includes columns for "ideology" (Conservative, Moderate, Liberal, Refused, Don't Know), and for "voted" (Biden, Trump, Refused, Someone Else, Don't Know). We want to see if the distribution of "ideology" for the subset of rows for which "voted" = "Refused" (refused to say) is the same as the distribution of "ideology" for all rows, or if it tilts in some direction.

```
# Note: the data file `31118130.csv` is a symlink to the original file in the "../3 + 4" directory in t
raw_data <- read_csv("31118130.csv")</pre>
##
## -- Column specification -----
## cols(
##
     .default = col_character(),
##
     end_state1 = col_double(),
     acarot = col_number(),
##
##
     endq1 = col_double(),
##
     q2rot = col_logical(),
     endq2 = col_double(),
##
     endq10 = col_double(),
##
     q17rot = col_number(),
##
     endq22 = col_double(),
##
     partyrot = col_number(),
##
##
     length = col_double(),
     reprostat = col_logical(),
##
##
     qctr = col double(),
     xctr = col_double(),
##
##
     wt1 = col_double(),
##
     weight = col_double(),
##
     standwt = col_double(),
##
     weight_ssrs = col_double(),
     cdc2013 = col_double()
##
## )
## i Use `spec()` for the full column specifications.
First step: get the frequency distribution for "ideology" for the whole set, which is 1676 rows.
ideology_count <- table(raw_data$ideology)</pre>
ideology_count
##
                                                  Moderate
                                                                 Refused
## Conservative
                   Don't Know
                                     Liberal
##
             527
                            68
                                         424
                                                       617
Then extract the subset of rows we want to examine, based on the answer to one question, and get the
frequency distribution of that subset, to compare for proportionality with the full set.
```

ideology_by_refused_voted2 <- raw_data %>%

select(ideology, voted2) %>%
filter(voted2 == "Refused")

```
voted2_refused_count <- table(ideology_by_refused_voted2$ideology)</pre>
voted2_refused_count
##
## Conservative
                   Don't Know
                                     Liberal
                                                  Moderate
                                                                 Refused
##
                                          16
                                                        37
                                                                       14
Then run a chisq.test on the two distributions. First off: is this comparison valid? The objects we're
comparing are tables. Not sure if that's an acceptable input type for chisq.test().
class(ideology_count)
## [1] "table"
class(voted2_refused_count)
## [1] "table"
chisq.test(ideology_count,voted2_refused_count)
## Warning in chisq.test(ideology_count, voted2_refused_count): Chi-squared
## approximation may be incorrect
##
##
    Pearson's Chi-squared test
##
## data: ideology_count and voted2_refused_count
## X-squared = 20, df = 16, p-value = 0.2202
The function doesn't complain about its inputs, and returns a p-value of 0.2202, meaning we can't reject
our null hypothesis and the differences in the distributions are explainable by chance. But I don't know if
this result is meaningful, because I don't know if the call made sense.
To force it another way, build a manual data set with these numbers, using xtabs to build cross-tabs, manually
entering the data from the above tables, which (are? aren't?) the same thing as a frequency distribution:
group <- c("all", "all", "all", "all", "refused", "refused", "refused", "refused", "refused")</pre>
ideology <- c("Moderate", "Conservative", "Liberal", "Don't Know", "Refused", "Moderate", "Conservativ
totals \leftarrow c(617,527,424,68,40,37,38,16,6,14)
A <- data.frame(group,ideology,totals)
Α
##
        group
                   ideology totals
## 1
          all
                   Moderate
                                617
## 2
                                527
          all Conservative
## 3
          all
                    Liberal
                                424
## 4
           all
                 Don't Know
                                 68
## 5
           a11
                    Refused
                                 40
## 6
     refused
                   Moderate
                                 37
                                 38
## 7
      refused Conservative
      refused
                    Liberal
                                  16
## 9 refused
                 Don't Know
                                  6
## 10 refused
                    Refused
                                 14
A_xtabs <- xtabs(totals~group+ideology,data=A)
A_xtabs
##
             ideology
## group
              Conservative Don't Know Liberal Moderate Refused
##
     all
                        527
                                     68
                                            424
                                                      617
                                                                40
```

```
## refused 38 6 16 37 14
```

Now run the chisq.test() on this xtab object:

```
chisq.test(A_xtabs,correct=F)
```

```
## Warning in chisq.test(A_xtabs, correct = F): Chi-squared approximation may be
## incorrect
##
## Pearson's Chi-squared test
##
## data: A_xtabs
## X-squared = 42.092, df = 4, p-value = 1.596e-08
```

This gives a very different p-value, showing that the difference in the distributions IS statistically significant and we can reject the null hypothesis.

The input this time is still a table, and also an "xtabs" object:

```
class(A_xtabs)
```

```
## [1] "xtabs" "table"
```

Finally, run a chisq.test() with a manual version of the dataset and a calculated expected values (as ratios) list, based on the subset as the observed values, and the expected values as ratios of the full set values to the size of the set. I would expect this test to give the same p-value as the above test:

```
obs <- c(37,38,16,6,14)
exp <- c(617/1676,527/1676,424/1676,68/1676,40/1676)
chisq.test(obs, p=exp)
```

```
## Warning in chisq.test(obs, p = exp): Chi-squared approximation may be incorrect
##
## Chi-squared test for given probabilities
##
## data: obs
## X-squared = 54.97, df = 4, p-value = 3.297e-11
```

It returns 3.297e-11... basically zero, but still 3 orders of magnitude off the other very tiny p-value of 1.596e-08. It also complains that the X-squared approximation may be off.

Running the Fisher test gives a different, but still very tiny P-value:

```
fisher.test(A_xtabs)
```

```
##
## Fisher's Exact Test for Count Data
##
## data: A_xtabs
## p-value = 5.339e-06
## alternative hypothesis: two.sided
...what am I doing wrong?
```