S632 HW5

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1. The dataset melanoma in the library faraway gives data on a sample of patients suffering from

melanoma (skin cancer) cross-classified by the type of cancer and the location on the body.

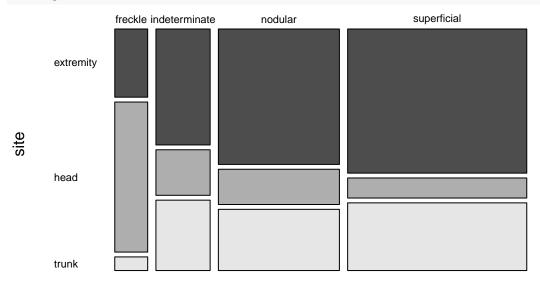
```
rm(list = 1s())

library(ggplot2)
library(faraway)
library(alr4)
library(dplyr)

cancer <- melanoma</pre>
```

a) Display the data in a contingency table and obtain a mosaic plot to visually check if *tumor* and *site* are independent. In addition, use a poisson model to determine if *tumor* and *site* are independent. Does your test agree with your visual conclusions?

```
ct <- xtabs(count ~ tumor + site, cancer)</pre>
ct
##
## tumor
                    extremity head trunk
##
     freckle
                            10
                                 22
                                         2
                                        17
##
     indeterminate
                            28
                                 11
                            73
##
     nodular
                                  19
                                        33
     superficial
                           115
                                 16
                                        54
##
mosaicplot(ct, color = TRUE, main = NULL, las = 1)
```



tumor

```
m1 <- glm(count ~ tumor + site, cancer, family = poisson)
pchisq(deviance(m1), df.residual(m1), lower = FALSE)</pre>
```

```
## [1] 2.050453e-09
```

Based on the mosaic plot, it appears that *tumor* and *site* are not independent because the elements of the grid are not all proportional and symmetric. Furthermore, when a chi-square test based on a poisson model is performed to test for independence it results in very low p-value allowing rejection of the null hypothesis that the two variables are independent, and supporting the earlier visual interpretation.

b) Make a two-way table of the deviance residuals from your model in a). Are there any larger residuals? Comment.

```
round(xtabs(residuals(m1) ~ tumor + site, cancer), 3)
##
                  site
## tumor
                   extremity
                               head trunk
##
     freckle
                      -2.316 5.135 -2.828
                      -0.660 0.468 0.548
##
     indeterminate
##
    nodular
                       0.281 -0.497 -0.022
##
     superficial
                       1.008 -3.045 0.699
```

Based on this two-way table of deviance residuals, we see that there are a number of larger residuals. All entries in the *freckle* row have quite large residuals, as do the *superficial* tumors found on the *head* and *extremity* regions. These factor levels with large residuals correspond to boxes in the mosaic plot from part a) that appeared quite different from the others - so I suppose it makes sense that they would be found to have relatively large deviance residuals.

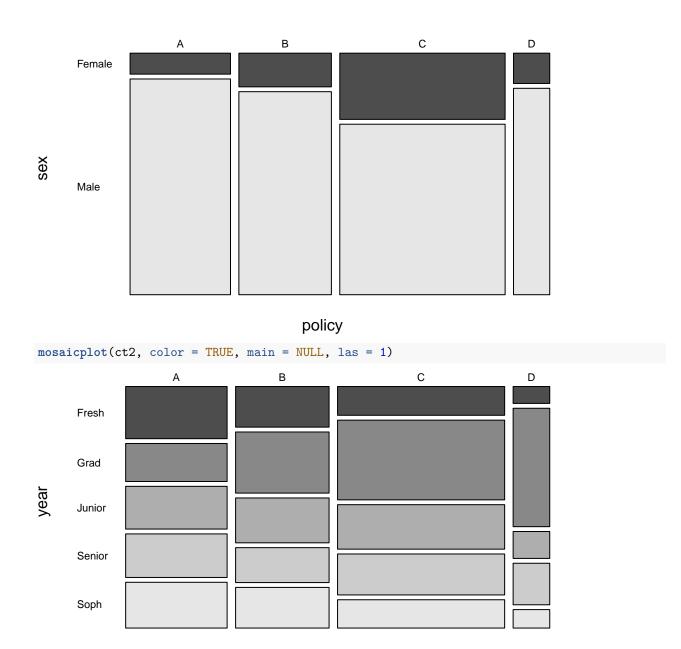
- 2. A student newspaper conducted a survey of student opinions about the Vietnam War in May 1967. Responses were classified by sex, year in the program and one of four opinions. The survey was voluntary. The data may be found in the dataset *uncviet* in the library *faraway*.
- a) Conduct an analysis of the patterns of dependence in the data assuming that all variables are nominal.

```
war <- uncviet

ct1 <- xtabs(y ~ policy + sex, war)

ct2 <- xtabs(y ~ policy + year, war)

mosaicplot(ct1, color = TRUE, main = NULL, las = 1)</pre>
```



policy

```
m2 <- glm(y ~ policy + sex, family = poisson, war)
pchisq(deviance(m2), df.residual(m2), lower = FALSE)
## [1] 2.351976e-112
m3 <- glm(y ~ policy + year, family = poisson, war)
pchisq(deviance(m3), df.residual(m3), lower = FALSE)</pre>
```

[1] 0

From the mosaic plots, and the poisson tests for independence, we see clearly that the variables sex and year are not independent from policy, when all variables are treated as nominal.

b) Assign scores to year and policy and fit an appropriate model. Interpret the trends in opinion over the years. Check the sensitivity of your conclusions to the assignment of the

```
scores (by trying other sensible alternatives).
war\gamma (c(1, 2, 3, 4, 5), each = 4)
war$policyo <- unclass(war$policy)</pre>
m1 <- glm(y ~ year + policy + sex, family = poisson, war)</pre>
Anova(m1)
## Analysis of Deviance Table (Type II tests)
## Response: y
         LR Chisq Df Pr(>Chisq)
##
## year
           216.56 4 < 2.2e-16 ***
## policy 718.59 3 < 2.2e-16 ***
           1349.11 1 < 2.2e-16 ***
## sex
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
m2 <- glm(y ~ year + policy + sex + I(yearo * policyo), family = poisson, war)
anova(m1, m2, test = "Chi")
## Analysis of Deviance Table
## Model 1: y ~ year + policy + sex
## Model 2: y ~ year + policy + sex + I(yearo * policyo)
## Resid. Df Resid. Dev Df Deviance Pr(>Chi)
## 1
            31
                   423.83
## 2
            30
                   246.13 1 177.69 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
# So there is evidence that there is an association between year and policy
# when they are treated as ordinal variables.
summary(m2)$coef["I(yearo * policyo)", ]
                  Std. Error
                                              Pr(>|z|)
##
       Estimate
                                  z value
## 1.751092e-01 1.354426e-02 1.292867e+01 3.101548e-38
# This coefficient is positive, indicating based on how the data is coded,
# that a higher year in college is associated with a greater probability of
# supporting less involvement in the Vietnam war.
# now to check sensitivity of the scores, I will assign different ones.
ayear \leftarrow c(1, 2, 3, 4, 8)
apolicy <-c(1, 5, 6, 10)
# Larger difference between senior and grad, as there can be way more than
# one year seperation between those two levels. Also, larger differences
# between the policy standings, as A seems really different from B and B and
# C are pretty similar, and D is quite different from C.
m2a <- glm(y ~ year + policy + sex + I(ayear[yearo] * apolicy[policyo]), family = poisson,</pre>
    war)
```

```
anova(m1, m2a, test = "Chi")
## Analysis of Deviance Table
##
## Model 1: y ~ year + policy + sex
## Model 2: y ~ year + policy + sex + I(ayear[yearo] * apolicy[policyo])
    Resid. Df Resid. Dev Df Deviance Pr(>Chi)
## 1
           31
                  423.83
## 2
           30
                  239.48 1
                              184.35 < 2.2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
# Same conclusion using different scores.
```

Based on the above data and code, we see that a higher year in college is associated with a greater probability of supporting reduced US involvement in the Vietnam war among UNC students in 1967. This conclusion is not dependent on a particular assignment of ordinal scores, as an alternative assignment strategy also leads to the same conclusion that there is an association between *year* and *policy* (both coding schemes tested resulted in a LR test p-value of 2.2e-16).

- 4. The hsb data from the library faraway was collected as a subset of the High School and Beyond study conducted by the National Education Longitudinal Studies program of the National Center for Education Statistics. The variables are gender; race; socioeconomic status (SES); school type; chosen high school program type; scores on reading, writing, math, science, and social studies. We want to determine which factors are related to the choice of the type of program academic, vocational or general that the students pursue in high school. The response is multinomial with three levels.
- a) Make a table showing the proportion of males and females choosing the three different programs. Comment on the difference. Repeat this comparison but for *SES* rather than gender.

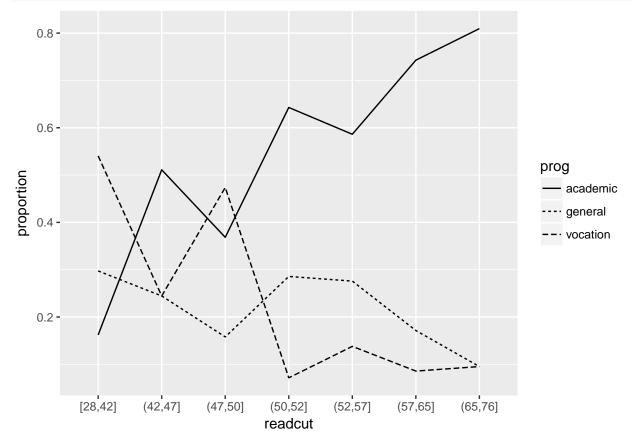
```
school <- hsb
schoolg <- group_by(school, gender, prog) %>% summarise(count = n()) %>% group_by(gender) %>%
    mutate(total = sum(count), proportion = count/total)
schools <- group_by(school, ses, prog) %>% summarise(count = n()) %>% group_by(ses) %>%
    mutate(total = sum(count), proportion = count/total)
xtabs(proportion ~ gender + prog, schoolg)
##
           prog
## gender
                        general vocation
             academic
##
     female 0.5321101 0.2201835 0.2477064
##
           0.5164835 0.2307692 0.2527473
xtabs(proportion ~ ses + prog, schools)
##
           prog
## ses
                        general vocation
             academic
##
            0.7241379 0.1551724 0.1206897
##
            0.4042553 0.3404255 0.2553191
     middle 0.4631579 0.2105263 0.3263158
##
```

There doesn't appear to be any large difference between the number of male and female students who choose different academic programs. On the other hand, there does seem to be a real difference between the academic program choices between different socioeconomic levels. Individuals from high ses backgrounds overwhelmingly choose to enroll in academic programs, while students from low and middle ses levels enroll more in general and vocational programs.

b) Construct a plot like the right panel of Figure 7.1 in ELM that shows the relationship between program choice and reading score. Comment on the plot.

```
schoolr <- mutate(school, readcut = cut_number(read, 7)) %>% group_by(readcut,
    prog) %>% summarise(count = n()) %>% group_by(readcut) %>% mutate(total = sum(count),
    proportion = count/total)

ggplot(schoolr, aes(x = readcut, y = proportion, group = prog, linetype = prog)) +
    geom_line()
```



The above plot shows that in general, as reading scores increase, the proportion of students choosing to enroll in academic programs also increases. Students with lower reading scores generally enroll in either general or vocational programs, though there is some variation to this pattern.

c) Fit a multinomial response model for the program choice and examine the fitted coefficients. Interpret at least two coefficients. In addition, observe that of the five subjects, one gives unexpected coefficients. Why do you think this happens?

```
library(nnet)
```

```
mmod <- multinom(prog ~ gender + race + ses + schtyp + read + write + math +
    science + socst, data = school)
## # weights: 42 (26 variable)
## initial value 219.722458
## iter 10 value 171.814970
## iter
        20 value 153.793692
## iter 30 value 152.935260
## final value 152.935256
## converged
summary(mmod)
## Call:
## multinom(formula = prog ~ gender + race + ses + schtyp + read +
##
       write + math + science + socst, data = school)
##
##
  Coefficients:
##
            (Intercept) gendermale raceasian racehispanic racewhite
               3.631901 -0.09264717 1.352739
                                                -0.6322019 0.2965156
## general
               7.481381 -0.32104341 -0.700070
##
  vocation
                                                -0.1993556 0.3358881
##
                seslow sesmiddle schtyppublic
                                                      read
                                                                 write
  general 1.09864111 0.7029621
                                    0.5845405 -0.04418353 -0.03627381
                                    2.0553336 -0.03481202 -0.03166001
  vocation 0.04747323 1.1815808
                  math
                          science
                                        socst
## general -0.1092888 0.10193746 -0.01976995
## vocation -0.1139877 0.05229938 -0.08040129
##
## Std. Errors:
##
            (Intercept) gendermale raceasian racehispanic racewhite
               1.823452 0.4548778
                                   1.058754
                                                0.8935504 0.7354829 0.6066763
## general
##
               2.104698
                         0.5021132
                                    1.470176
                                                0.8393676 0.7480573 0.7045772
##
            sesmiddle schtyppublic
                                         read
                                                    write
                                                                math
  general 0.5045938
                         0.5642925 0.03103707 0.03381324 0.03522441
                         0.8348229 0.03422409 0.03585729 0.03885131
## vocation 0.5700833
##
               science
                            socst
## general 0.03274038 0.02712589
## vocation 0.03424763 0.02938212
##
## Residual Deviance: 305.8705
## AIC: 357.8705
```

In the above model, I will interpret the coefficients for the program levels general and vocation for the regressor gender. First, the log-odds of moving from the baseline program category of academic to the category general decrease by 0.093, or $(1-e^{-0.093}) \times 100 = 8.85\%$, when we move from considering female, to male students. Similarly, the log-odds of moving from the category academic to vocation decrease by 0.321, or $(1-e^{-0.321}) \times 100 = 27.5\%$, when we move from female to male students.

Furthermore, we can see from the summary above that of the five subject tests, *science* gives coefficients that are in the opposite direction of the others. Where the other four subjects have negative coefficients, saying that increasing scores on those tests results in lower odds of attending either a general or vocational school compared to an academic one, higher scores on the science test lead to *increased* odds of attending either general or vocational schools. One possible reason for this observation could be that students with parents who work in vocational, or otherwise "non-academic" jobs (like plumbers, carpenters, electricians, etc.) may have more exposure to

real-life scientific concepts found in physics and chemistry and so find that those concepts are easier to understand, leading to them doing better on that subject test. These same students may then choose to enter a general or vocational high school, not because of their test scores, but because of their family history in those situations. Alternatively, maybe the science subject test was the last test taken, and the high-achieving students, who would later go on to enter academic programs, were tired from their high effort on the other tests and so did comparitively worse on that one.