MA678 homework 06

Multinomial Regression

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Multinomial logit:

Using the individual-level survey data from the 2000 National Election Study (data in folder NES), predict party identification (which is on a five-point scale) using ideology and demographics with an ordered multinomial logit model.

```
## Call:
## polr(formula = factor(partyid7) ~ age + gender + race + ideo,
##
       data = nes, Hess = TRUE)
##
## Coefficients:
##
                                               ideo
           age
                    gender
                                   race
  -0.01193745 -0.17256010 -0.25333872
                                        0.42067639
##
## Intercepts:
##
                                 3|4
                                            4|5
                                                       5|6
                                                                   6|7
          1|2
                     2|3
  -1.3901593 -0.3133816 0.2740848 0.6796139
                                                1.2965932 2.3213684
##
## Residual Deviance: 44523.29
## AIC: 44543.29
## (22681 observations deleted due to missingness)
```

1. Summarize the parameter estimates numerically and also graphically.

summary(fit)

```
## Call:
  polr(formula = factor(partyid7) ~ age + gender + race + ideo,
       data = nes, Hess = TRUE)
##
##
## Coefficients:
##
             Value Std. Error t value
## age
          -0.01194
                     0.000982 - 12.156
  gender -0.17256
                     0.032103 -5.375
## race
          -0.25334
                     0.015284 -16.575
##
  ideo
           0.42068
                     0.009925 42.387
##
## Intercepts:
       Value
                Std. Error t value
## 1|2 -1.3902
                  0.0795
                            -17.4885
## 2|3
        -0.3134
                  0.0780
                             -4.0193
## 3|4
         0.2741
                  0.0780
                              3.5154
## 4|5
         0.6796
                  0.0783
                              8.6817
```

```
## 5|6 1.2966 0.0791 16.4009

## 6|7 2.3214 0.0814 28.5017

##

## Residual Deviance: 44523.29

## AIC: 44543.29

## (22681 observations deleted due to missingness)
```

2. Explain the results from the fitted model.

```
#a person is more likely to be higher in partyid if they are younger, have a lower indicator value for
```

3. Use a binned residual plot to assess the fit of the model.

(Optional) Choice models:

Using the individual-level survey data from the election example described in Section 10.9 (data available in the folder NES),

- 1. fit a logistic regression model for the choice of supporting Democrats or Republicans. Then interpret the output from this regression in terms of a utility/choice model.
- 2. Repeat the previous exercise but now with three options: Democrat, no opinion, Republican. That is, fit an ordered logit model and then express it as a utility/choice mode

Contingency table and ordered logit model

In a prospective study of a new living attenuated recombinant vaccine for influenza, patients were randomly allocated to two groups, one of which was given the new vaccine and the other a saline placebo. The responses were titre levels of hemagglutinin inhibiting antibody found in the blood six weeks after vaccination; they were categorized as "small", "medium" or "large".

treatment	small	moderate	large	Total
placebo	25	8	5	38
vaccine	6	18	11	35

The cell frequencies in the rows of table are constrained to add to the number of subjects in each treatment group (35 and 38 respectively). We want to know if the pattern of responses is the same for each treatment group.

1. Using a chisqure test and an appropriate log-linear model, test the hypothesis that the distribution of responses is the same for the placebo and vaccine groups.

```
chisq.test(data)

##

## Pearson's Chi-squared test

##

## data: data

## X-squared = 17.648, df = 2, p-value = 0.0001472

stan_glm(Freq ~ treatment + levels, data = df, refresh = 0)

## stan_glm

## family: gaussian [identity]

## formula: Freq ~ treatment + levels

## observations: 6

## predictors: 4
```

```
##
                    Median MAD SD
## (Intercept)
                    15.5
                             8.0
## treatmentvaccine -1.0
                             8.2
## levelsmoderate
                    -2.0
                            10.0
## levelslarge
                    -7.0
                             9.8
##
## Auxiliary parameter(s):
##
         Median MAD SD
## sigma 10.4
                 3.9
##
## * For help interpreting the printed output see ?print.stanreg
## * For info on the priors used see ?prior_summary.stanreg
```

2. For the model corresponding to the hypothesis of homogeneity of response distributions, calculate the fitted values, the Pearson and deviance residuals, and the goodness of fit statistics X^2 and D. Which of the cells of the table contribute most to X^2 and D? Explain and interpret these results.

```
chisq <- chisq.test(data)
chisq

##

## Pearson's Chi-squared test
##

## data: data
## X-squared = 17.648, df = 2, p-value = 0.0001472</pre>
```

3. Re-analyze these data using ordered logit model (use polr) to estiamte the cut-points of a latent continuous response variable and to estimate a location shift between the two treatment groups. Sketch a rough diagram to illustrate the model which forms the conceptual base for this analysis.

High School and Beyond

The hsb data 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; 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.

```
data(hsb)
?hsb
```

1. Fit a trinomial response model with the other relevant variables as predictors (untransformed).

```
fit <- multinom(factor(prog)~gender + race + ses + read + write + math + science + socst + schtyp, data
## # 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
fit</pre>
```

Call:

```
## multinom(formula = factor(prog) ~ gender + race + ses + read +
##
       write + math + science + socst + schtyp, data = hsb, hess = TRUE)
##
## Coefficients:
##
            (Intercept) gendermale raceasian racehispanic racewhite
               3.631901 -0.09264717 1.352739
                                                -0.6322019 0.2965156 1.09864111
## general
               7.481381 -0.32104341 -0.700070
                                                 -0.1993556 0.3358881 0.04747323
## vocation
##
            sesmiddle
                             read
                                         write
                                                     math
                                                             science
## general 0.7029621 -0.04418353 -0.03627381 -0.1092888 0.10193746 -0.01976995
## vocation 1.1815808 -0.03481202 -0.03166001 -0.1139877 0.05229938 -0.08040129
            schtyppublic
               0.5845405
## general
               2.0553336
## vocation
##
## Residual Deviance: 305.8705
## AIC: 357.8705
```

2. For the student with id 99, compute the predicted probabilities of the three possible choices.

```
subset <- subset(hsb, hsb$id==99)
fitted(fit)[102,]
## academic general vocation</pre>
```

Happiness

0.5076752 0.3753090 0.1170158

Data were collected from 39 students in a University of Chicago MBA class and may be found in the dataset happy.

```
library(faraway)
data(happy)
?happy
```

1. Build a model for the level of happiness as a function of the other variables.

```
fit <- polr(factor(happy)~money + sex + love + work, data = happy, Hess = TRUE)</pre>
fit
## Call:
## polr(formula = factor(happy) ~ money + sex + love + work, data = happy,
##
       Hess = TRUE)
##
##
  Coefficients:
##
        money
                                love
                                            work
                      sex
##
    0.0224593 -0.4734369 3.6076452 0.8875135
##
## Intercepts:
                              4|5
                                        5|6
                                                   6|7
                                                             7|8
                                                                        819
                                                                                 9|10
##
         2|3
                   3|4
    5.470845 6.468394 9.159127 10.972524 11.511333 13.543305 17.290890 19.011197
## Residual Deviance: 94.86029
## AIC: 118.8603
```

2. Interpret the parameters of your chosen model.

3. Predict the happiness distribution for subject whose parents earn \$30,000 a year, who is lonely, not sexually active and has no job.

```
newdata = data.frame(money=30, sex=0,love=1,work=1)
predict(fit, newdata=newdata, type = "probs")

## 2 3 4 5 6 7
## 5.749087e-01 2.108348e-01 1.960962e-01 1.515266e-02 1.250656e-03 1.526336e-03
## 8 9 10
## 2.252137e-04 4.465166e-06 9.736048e-07
```

newspaper survey on Vietnam War

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. Treat the opinion as the response and the sex and year as predictors. Build a proportional odds model, giving an interpretation to the estimates.

```
data(uncviet)
?uncviet
fit <- model <- polr(policy ~ sex + year, weights=y, data=uncviet)
## Call:
## polr(formula = policy ~ sex + year, data = uncviet, weights = y)
##
## Coefficients:
##
      sexMale
                yearGrad yearJunior yearSenior
                                                 yearSoph
  -0.6470352 1.1769887 0.3964211 0.5443945
##
                                               0.1315047
##
##
  Intercepts:
##
           AB
                       BIC
                                   CID
##
  -1.10979578 -0.01304875
                            2.44169665
##
## Residual Deviance: 7757.056
## AIC: 7773.056
# students are more likely to want the US to withdraw from Vietnam if they are older and if they are fe
```

pneumonoconiosis of coal miners

The pneumo data gives the number of coal miners classified by radiological examination into one of three categories of pneumonoconiosis and by the number of years spent working at the coal face divided into eight categories.

```
library(faraway)
data(pneumo,package="faraway")
?pneumo

## Help on topic 'pneumo' was found in the following packages:
##
## Package Library
## VGAM /Library/Frameworks/R.framework/Versions/4.0/Resources/library
```

```
##
     faraway
                            /Library/Frameworks/R.framework/Versions/4.0/Resources/library
##
##
## Using the first match ...
  1. Treating the pneumonoconiosis status as response variable as nominal, build a model for predicting the
     frequency of the three outcomes in terms of length of service and use it to predict the outcome for a
     miner with 25 years of service.
fit <- multinom(status ~ year, data = pneumo, Hess = TRUE)
## # weights: 9 (4 variable)
## initial value 26.366695
## final value 26.366695
## converged
fit
## Call:
## multinom(formula = status ~ year, data = pneumo, Hess = TRUE)
##
## Coefficients:
##
           (Intercept)
                                 year
## normal 2.109424e-15 2.486900e-14
## severe 2.664535e-15 3.552714e-14
## Residual Deviance: 52.73339
## AIC: 60.73339
newdata=data.frame(year=25)
predict(fit, newdata=newdata, type="probs")
##
        mild
                normal
                           severe
## 0.3333333 0.3333333 0.3333333
  2. Repeat the analysis with the pneumonoconiosis status being treated as ordinal.
fit <- polr(status ~ year, data = pneumo, Hess = TRUE)
fit
## polr(formula = status ~ year, data = pneumo, Hess = TRUE)
## Coefficients:
           year
## 4.340705e-11
##
## Intercepts:
##
     mild|normal normal|severe
##
      -0.6931472
                      0.6931472
##
## Residual Deviance: 52.73339
## AIC: 58.73339
newdata=data.frame(year=25)
predict(fit, newdata=newdata, type="probs")
##
        mild
                 normal
                           severe
```

0.3333333 0.3333333 0.3333333

3. Now treat the response variable as hierarchical with top level indicating whether the miner has the disease and the second level indicating, given they have the disease, whether they have a moderate or severe case.

4. Compare the three analyses.

the first two are the same

(optional) Multinomial choice models:

Pardoe and Simonton (2006) fit a discrete choice model to predict winners of the Academy Awards. Their data are in the folder Academy Awards.

name	description	
No	unique nominee identifier	
Year	movie release year (not ceremony year)	
Comp	identifier for year/category	
Name	short nominee name	
PP	best picture indicator	
DD	best director indicator	
MM	lead actor indicator	
FF	lead actress indicator	
Ch	1 if win, 2 if lose	
Movie	short movie name	
Nom	total oscar nominations	
Pic	picture nom	
Dir	director nom	
Aml	actor male lead nom	
Afl	actor female lead nom	
Ams	actor male supporting nom	
Afs	actor female supporting nom	
Scr	screenplay nom	
Cin	cinematography nom	
Art	art direction nom	
\cos	costume nom	
Sco	score nom	
Son	song nom	
Edi	editing nom	
Sou	sound mixing nom	
For	foreign nom	
Anf	animated feature nom	
Eff	sound editing/visual effects nom	
Mak	makeup nom	
Dan	dance nom	
AD	assistant director nom	
PrNl	previous lead actor nominations	
PrWl	previous lead actor wins	
PrNs	previous supporting actor nominations	
PrWs	previous supporting actor wins	
PrN	total previous actor/director nominations	
PrW	total previous actor/director wins	
Gdr	golden globe drama win	
Gmc	golden globe musical/comedy win	
Gd	golden globe director win	
Gm1	golden globe male lead actor drama win	

name	description
Gm2	golden globe male lead actor musical/comedy win
Gf1	golden globe female lead actor drama win
Gf2	golden globe female lead actor musical/comedy win
PGA	producer's guild of america win
DGA	director's guild of america win
SAM	screen actor's guild male win
SAF	screen actor's guild female win
PN	PP*Nom
PD	PP*Dir
DN	DD*Nom
DP	DD*Pic
DPrN	$\mathrm{DD}^*\mathrm{PrN}$
DPrW	$\mathrm{DD}^*\mathrm{PrW}$
MN	MM*Nom
MP	MM*Pic
MPrN	MM*PrNl
MPrW	MM*PrWl
FN	FF*Nom
FP	FF*Pic
FPrN	FF*PrNl
FPrW	FF*PrWl

- 1. Fit your own model to these data.
- 2. Display the fitted model on a plot that also shows the data.
- 3. Make a plot displaying the uncertainty in inferences from the fitted model.