MA678 homework 05

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 7-point scale) using ideology and demographics with an ordered multinomial logit model.

1. Summarize the parameter estimates numerically and also graphically.

```
x = nes_data_comp$partyid7
nes_data_comp <- nes_data_comp[!is.na(levels(x)[x]),]</pre>
nes.order <- polr(factor(partyid7) ~ ideo + age + gender + race + south, Hess = TRUE, data = nes_data_c
summary(nes.order)
## Call:
## polr(formula = factor(partyid7) ~ ideo + age + gender + race +
##
       south, data = nes data comp, Hess = TRUE)
##
## Coefficients:
##
                           Value Std. Error t value
## ideomoderate
                        0.95339
                                  0.330586 2.8840
## ideoconservative
                        1.94046
                                   0.181464 10.6933
                                   0.004937 -2.6676
                       -0.01317
## genderfemale
                       -0.38543
                                  0.155547 -2.4779
## raceblack
                       -1.79583
                                   0.277242 -6.4775
                                   0.544657 0.2303
## raceasian
                        0.12546
## racenative american -0.13670
                                   0.368338 -0.3711
## racehispanic
                       -0.62635
                                   0.297434 -2.1058
## south1
                        0.21547
                                   0.175126 1.2304
##
## Intercepts:
##
                                                          Value
                                                                  Std. Error
                                                          -1.3844
## 1. strong democrat | 2. weak democrat
                                                                  0.3053
## 2. weak democrat | 3. independent-democrat
                                                          -0.5338
                                                                   0.2973
## 3. independent-democrat | 4. independent-independent
                                                           0.2580
                                                                   0.2969
## 4. independent-independent|5. independent-republican 0.6528
                                                                  0.2995
## 5. independent-republican | 6. weak republican
                                                           1.4496
                                                                  0.3046
## 6. weak republican | 7. strong republican
                                                           2.4281
                                                                  0.3159
##
                                                          t value
## 1. strong democrat | 2. weak democrat
                                                          -4.5350
                                                          -1.7958
## 2. weak democrat|3. independent-democrat
## 3. independent-democrat | 4. independent-independent
                                                           0.8689
## 4. independent-independent|5. independent-republican 2.1799
## 5. independent-republican | 6. weak republican
                                                           4.7597
## 6. weak republican | 7. strong republican
                                                           7.6864
```

```
## Residual Deviance: 1889.52
## AIC: 1919.52
## (9 observations deleted due to missingness)
```

round(summary(nes.order)\$coef,2)

##		Value Std.	Error
##	ideomoderate	0.95	0.33
##	ideoconservative	1.94	0.18
##	age	-0.01	0.00
##	genderfemale	-0.39	0.16
##	raceblack	-1.80	0.28
##	raceasian	0.13	0.54
##	racenative american	-0.14	0.37
##	racehispanic	-0.63	0.30
##	south1	0.22	0.18
##	1. strong democrat 2. weak democrat	-1.38	0.31
##	2. weak democrat 3. independent-democrat	-0.53	0.30
##	3. independent-democrat 4. independent-independent	0.26	0.30
##	${\tt 4. independent-independent 5. independent-republican}$	0.65	0.30
	5. independent-republican 6. weak republican	1.45	0.30
##	6. weak republican 7. strong republican	2.43	0.32
##		t value	
##	ideomoderate	2.88	
##	ideoconservative	10.69	
##	age	-2.67	
##	genderfemale	-2.48	
##	raceblack	-6.48	
##	raceasian	0.23	
##	racenative american	-0.37	
##	racehispanic	-2.11	
##	south1	1.23	
##	1. strong democrat 2. weak democrat	-4.54	
	2. weak democrat 3. independent-democrat	-1.80	
	3. independent-democrat 4. independent-independent	0.87	
	${\tt 4. independent-independent 5. independent-republican}$		
	5. independent-republican 6. weak republican	4.76	
##	6. weak republican 7. strong republican	7.69	

2. Explain the results from the fitted model.

confint(nes.order)

Waiting for profiling to be done...

```
2.5 %
                                        97.5 %
## ideomoderate
                       0.30274518 1.602885103
## ideoconservative
                      1.58819102 2.299908250
                      -0.02288059 -0.003516047
## age
## genderfemale
                      -0.69089528 -0.080899207
## raceblack
                      -2.34876169 -1.259970216
## raceasian
                      -0.94682823 1.212855171
## racenative american -0.86519074 0.585344750
## racehispanic
                      -1.21378606 -0.045074705
## south1
                      -0.12813859 0.558798934
```

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

```
nes <- cbind(partyid7 = nes_data_comp$partyid7, ideo = nes_data_comp$ideo, race = nes_data_comp$race, a
nes <- data.frame(na.omit(nes))</pre>
resid <- model.matrix(~ factor(partyid7) - 1, data = nes) - fitted(nes.order)
par(mfrow = c(3, 2))
for (i in 1:6) {
  binnedplot(fitted(nes.order)[, i], resid[, i], cex.main = 1.3, main = "Binned residual plot")
}
                                                                                Binned residual plot
                        Binned residual plot
                                                             Average residual
      Average residual
                       0.2
                                  0.4
                                            0.6
                                                                    0.05
                                                                                 0.10
                                                                                             0.15
                                                                                                          0.20
                            Expected Values
                                                                                    Expected Values
                        Binned residual plot
                                                                                Binned residual plot
      Average residual
                                                             Average residual
                                                                                 0.04
                         0.10
                                                                       0.02
              0.06
                    0.08
                              0.12 0.14
                                          0.16
                                                0.18
                                                                                          0.06
                                                                                                    0.08
                                                                                                             0.10
                                                                                    Expected Values
                            Expected Values
                        Binned residual plot
                                                                                Binned residual plot
                                                             Average residual
      Average residual
                    0.05
                                          0.15
                                                     0.20
                                                                            0.05
                                                                                     0.10
                                                                                             0.15
                                                                                                      0.20
                               0.10
                            Expected Values
                                                                                    Expected Values
```

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).

```
library(nnet)
reg.hs <- multinom(prog ~ read + write + math + science + race,data = hsb,trace = FALSE,HESS = TRUE)
summary(reg.hs)

## Call:
## multinom(formula = prog ~ read + write + math + science + race,
## data = hsb, trace = FALSE, HESS = TRUE)</pre>
```

```
##
## Coefficients:
##
            (Intercept)
                                read
                                           write
               4.924957 -0.05388450 -0.03946933 -0.1071044 0.09229507
## general
## vocation
               8.777829 -0.05594167 -0.06281609 -0.1253231 0.05262485
             raceasian racehispanic
##
                                       racewhite
## general 1.11489221 -0.60687283 -0.01313942
## vocation 0.08636574
                         0.07298783 0.42373684
##
## Std. Errors:
            (Intercept)
                               read
                                         write
                                                      math
                                                              science raceasian
               1.528744 0.02853999 0.02864533 0.03391490 0.03053422 0.9950814
## general
               1.629837 0.03052243 0.02855810 0.03616922 0.03106921 1.3388885
## vocation
##
            racehispanic racewhite
               0.8707214 0.6995466
## general
## vocation
               0.7864713 0.6836971
##
## Residual Deviance: 332.6696
## AIC: 364.6696
  2. For the student with id 99, compute the predicted probabilities of the three possible choices.
predict(reg.hs,newdata=hsb[hsb$id==99,],type="probs")
## academic
               general vocation
## 0.3756043 0.4338602 0.1905356
```

Happiness

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.

```
library(nnet)
reg.happy <- polr(factor(happy) ~ money + sex + love + work,data = happy)
summary(reg.happy)
##
## Re-fitting to get Hessian
## Call:
## polr(formula = factor(happy) ~ money + sex + love + work, data = happy)
##
## Coefficients:
##
            Value Std. Error t value
## money 0.02246
                     0.01066 2.1064
## sex
         -0.47344
                     0.79498 -0.5955
## love
          3.60765
                     0.80114 4.5031
## work
         0.88751
                     0.40826 2.1739
##
## Intercepts:
```

```
##
        Value
                Std. Error t value
                 1.9891
         5.4708
                             2.7504
## 2|3
                 1.9223
## 3|4
         6.4684
                             3.3650
                             4.2212
## 4|5
         9.1591
                 2.1698
  516
        10.9725
                 2.3213
                             4.7268
  617
        11.5113
                 2.3720
                             4.8530
        13.5433
                             5.0776
## 718
                 2.6673
        17.2909
## 8|9
                 3.1454
                             5.4971
## 9|10 19.0112
                 3.3270
                             5.7142
##
## Residual Deviance: 94.86029
## AIC: 118.8603
```

confint(reg.happy)

2. Interpret the parameters of your chosen model.

```
## Waiting for profiling to be done...

##

## Re-fitting to get Hessian

## 2.5 % 97.5 %

## money 0.002276811 0.04490097

## sex -2.068912556 1.07918378

## love 2.168908570 5.37172936

## work 0.123787534 1.74622976
```

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

```
predict(reg.happy,newdata = data.frame(love = 1,sex = 0,work = 1,money = 30),type = "probs")
## 2 3 4 5 6
## 5.749087e-01 2.108348e-01 1.960962e-01 1.515266e-02 1.250656e-03
## 7 8 9 10
## 1.526336e-03 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
```

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:
##
##
                            Library
     Package
     VGAM
                            /Library/Frameworks/R.framework/Versions/3.4/Resources/library
##
##
                            /Library/Frameworks/R.framework/Versions/3.4/Resources/library
     faraway
##
##
## 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.
pneumo.reg <- multinom(status ~ year, data = pneumo)</pre>
## # weights: 9 (4 variable)
## initial value 26.366695
## final value 26.366695
## converged
summary(pneumo.reg)
## Call:
## multinom(formula = status ~ year, data = pneumo)
## Coefficients:
           (Intercept)
##
                                year
## normal 2.109424e-15 2.486900e-14
## severe 2.664535e-15 3.552714e-14
##
## Std. Errors:
##
        (Intercept)
                             year
## normal 1.142515 0.03420049
           1.142515 0.03420049
## severe
##
## Residual Deviance: 52.73339
## AIC: 60.73339
predict(pneumo.reg, data.frame (year = 25), type = "probs")
##
        mild
                normal
                           severe
## 0.3333333 0.3333333 0.3333333
  2. Repeat the analysis with the pneumonoconiosis status being treated as ordinal.
pneumo.order <- polr(factor(status) ~ year, data = pneumo, Hess = TRUE)</pre>
summary(pneumo.order)
## Call:
## polr(formula = factor(status) ~ year, data = pneumo, Hess = TRUE)
##
## Coefficients:
            Value Std. Error t value
## year 4.341e-11 0.02565 1.692e-09
```

```
##
## Intercepts:
##
                  Value
                          Std. Error t value
## mild|normal
                 -0.6931 0.8838
                                      -0.7842
## normal|severe 0.6931 0.8838
                                       0.7842
##
## Residual Deviance: 52.73339
## AIC: 58.73339
predict(pneumo.order, data.frame (year = 25), 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.
pneumo$status.h <- ifelse(pneumo$status == "normal", 0, 1)</pre>
pneumo.sub <- as.data.frame(cbind(Freq = pneumo$Freq, normal = ifelse(pneumo$status == "normal",1,0), m
pneumo.hie <- multinom(cbind(normal,mild,severe) ~ year, data = pneumo.sub)</pre>
## # weights: 9 (4 variable)
## initial value 26.366695
## final value 26.366695
## converged
summary(pneumo.hie)
## Call:
## multinom(formula = cbind(normal, mild, severe) ~ year, data = pneumo.sub)
##
## Coefficients:
           (Intercept)
          1.221245e-15 2.486900e-14
## mild
## severe 2.664535e-15 3.552714e-14
##
## Std. Errors:
##
          (Intercept)
                             year
## mild
             1.142515 0.03420049
             1.142515 0.03420049
## severe
##
## Residual Deviance: 52.73339
## AIC: 60.73339
predict(pneumo.hie, data.frame(year = 25), type = "probs")
      normal
                   mild
                           severe
## 0.3333333 0.3333333 0.3333333
```

4. Compare the three analyses. The results from these three analyses are quite similar, and they are with mild around 0.08-0.10, normal around 0.78-0.83, and severe around 0.08-0.12.

(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
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
DAM	screen actor's guild mate will

name	description
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	DD*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.