MAST90104: A First Course in Statistical Learning

Week 12 Lab and Workshop

1 Practical questions

AIC: 357.8705

- 1. In the multinom function from the nnet package, the response should be a factor with J levels or a matrix with J columns, which will be interpreted as counts for each of J classes. The first case is a short hand for responses of the form multinomial(1, p). The hsb data from the faraway package was collected as a subset of the "High School and Beyond" study, conducted by the National Education Longitudinal Studies program of the U.K. 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.
 - (a) Fit a trinomial response model with the other relevant variables as predictors (untransformed). **Solution:**

```
> library(faraway)
> data(hsb)
 library(nnet)
 mmod <- multinom(prog ~ gender + race + ses + schtyp + read + write + math +</pre>
                     science + socst, hsb, trace = FALSE)
> summary(mmod)
multinom(formula = prog ~ gender + race + ses + schtyp + read +
write + math + science + socst, data = hsb, trace = FALSE)
Coefficients:
         (Intercept)
                      gendermale raceasian racehispanic racewhite
            3.631901 -0.09264717 1.352739
                                              -0.6322019 0.2965156 1.09864111
general
vocation
            7.481381 -0.32104341 -0.700070
                                              -0.1993556 0.3358881 0.04747323
          sesmiddle schtyppublic
                                      read
                                                 write
                                                             math
general 0.7029621
                      0.5845405 - 0.04418353 - 0.03627381 - 0.1092888 0.10193746
                      2.0553336 -0.03481202 -0.03166001 -0.1139877 0.05229938
vocation 1.1815808
socst
general -0.01976995
vocation -0.08040129
Std. Errors:
         (Intercept) gendermale raceasian racehispanic racewhite
                                                                      seslow sesmiddle
general
            1.823452 0.4548778 1.058754
                                              0.8935504 0.7354829 0.6066763 0.5045938
            2.104698 0.5021132
                                 1.470176
                                              0.8393676 \ 0.7480573 \ 0.7045772 \ 0.5700833
vocation
           schtyppublic
                                      write
                                                  math
                                                          science
                           read
            0.5642925 0.03103707 0.03381324 0.03522441 0.03274038 0.02712589
general
vocation
            0.8348229 0.03422409 0.03585729 0.03885131 0.03424763 0.02938212
Residual Deviance: 305.8705
```

(b) Use either backward elimination with χ^2 tests (using the anova command), or the AIC (using step), to produce a parsimonious model. Give an interpretation of the resulting model. Solution: Use backward selection with AIC

```
> mmod2 <- step(mmod, scope=~., trace = FALSE, direction = "backward")
> summary(mmod2)
multinom(formula = prog ~ ses + schtyp + math + science + socst,
data = hsb, trace = FALSE)
Coefficients:
           (Intercept)
                           seslow sesmiddle schtyppublic
                                                                 math
             2.587029 0.87607389 0.6978995
                                                  0.6468812 -0.1212242 0.08209791
general
             6.687272 -0.01569301 1.2065000
                                                  1.9955504 -0.1369641 0.03941237
vocation
socst
general -0.04441228
vocation -0.09363417
Std. Errors:
          (Intercept)
                          seslow sesmiddle
                                               schtyppublic
                                                                 math
                                                                          science
             1.686492 0.5758781 0.4930330
                                                 0.545598 0.03213345 0.02787694
general
                                                 0.812881 0.03591701 0.02864929
vocation
             1.945363 0.6690861 0.5571202
socst
general 0.02344856
vocation 0.02586717
Residual Deviance: 315.5511
AIC: 343.5511
Compared to students from a high socioeconomic class, students from a low socioeconomic
class are more likely to choose a general high school program, while students from a middle
socioeconomic class are more likely to choose a general program but even more likely to choose
a vocational program. It is interesting that students from a low socioeconimic class do not
show more of an interest in vocational programs.
Students from public schools are are more likely to choose a general program and much more
likely to choose a vocational program, than students from private schools.
High scores in maths and social sciences indicate a higher chance of choosing an academic pro-
gram, while (curiously) high scores in science indicate a lower chance of choosing an academic
program.
If you wish to use a chisquared test instead of the AIC, then you will have to separately fit
all the candidate models, and then compare them using anova. For example:
> mmodXgender <- multinom(prog ~ race + ses + schtyp + read + write + math +
                              science + socst, hsb, trace = FALSE)
> anova(mmod, mmodXgender)
Likelihood ratio tests of Multinomial Models
Response: prog
```

Model

```
1
           race + ses + schtyp + read + write + math + science + socst
2 gender + race + ses + schtyp + read + write + math + science + socst
     Resid. df Resid. Dev
                             Test
                                      Df
                                           LR stat.
                                                         Pr(Chi)
        376
1
                 306.28565
        374
                                           0.41514199
                 305.87051
                             1 vs 2
                                       2
                                                        0.81255555
```

(c) For the student with id 99, compute the predicted probabilities of the three possible choices. Solution:

```
> hsb[hsb$id==99,]
    id gender race ses schtyp
                                   prog read write math science socst
102 99 female white high public general
                                          47
                                                59
                                                      56
                                                              66
                                                                    61
> predict(mmod2, newdata = hsb[hsb$id==99,], type="probs")
```

```
academic general vocation 0.644263088 0.276656088 0.079080824
```

- 2. The pneumo data from the faraway package 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.
 - (a) 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.

Solution: First we have a look at the data. Then the data needs to be reformatted before we can use the multinom function to fit a model. The fit looks quite good.

```
> data(pneumo)
> counts <- xtabs(Freq ~ status + year, pneumo)</pre>
> (props <- prop.table(counts, 2))</pre>
year
                                     21.5
status
            5.8
                                                 27.5
       0.00000000 0.037037037 0.139534884 0.104166667 0.196078431
mild
normal 1.000000000 0.944444444 0.790697674 0.729166667 0.627450980
severe 0.000000000 0.018518519 0.069767442 0.166666667 0.176470588
year
status
           39.5
                          46
                                    51 5
mild
       0.184210526 0.214285714 0.181818182
normal 0.605263158 0.428571429 0.363636364
severe 0.210526316 0.357142857 0.454545455
> years <- c(5.8, 15, 21.5, 27.5, 33.5, 39.5, 46, 51.5)
> par(mfrow=c(1,1))
> plot(years, props[1,], col="red", ylim=c(0,1))
> points(years, props[2,], col="blue")
> points(years, props[3,], col="green")
> mmod <- multinom(t(counts) ~ years, trace=FALSE)</pre>
> summary(mmod)
Call:
multinom(formula = t(counts) ~ years, trace = FALSE)
Coefficients:
        (Intercept)
                         years
normal 4.29167227 -0.083565062
severe -0.76817058 0.025720269
Std. Errors:
        (Intercept)
                         years
normal 0.52141098 0.015280443
severe 0.73771918 0.019766616
Residual Deviance: 417.44956
AIC: 425.44956
For a miner with 25 year down pit we have the following fitted probabilities
> predict(mmod, newdata=list(years=25), type="probs")
```

In the model above we had eight multinomial observations, with the number of trials equal to 98, 54, 43, 48, 51, 38, 28, 11. Each of these multinomials can be regarded as the sum of a number of independent multinomials each based on a single trial (just as a binomial is a sum of independent Bernoulli random variables). If we treat the data this way and fit a multinomial logistic regression, we get the same model.

severe

mild

normal

0.091488209 0.827786959 0.080724832

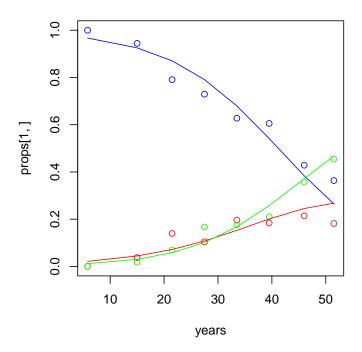


Figure 1: Fitted probability of the three outcomes vs length of service

```
> pneumo2 <- data.frame(status = rep(pneumo$status, pneumo$Freq),</pre>
year = rep(pneumo$year, pneumo$Freq))
> mmod2 <- multinom(status ~ year, data = pneumo2, trace = FALSE)
> summary(mmod2)
Call:
multinom(formula = status ~ year, data = pneumo2, trace = FALSE)
Coefficients:
        (Intercept)
                         year
       4.29167227 -0.083565062
severe -0.76817058 0.025720269
Std. Errors:
        (Intercept)
                        year
       0.52141098 0.015280443
normal
severe 0.73771918 0.019766616
Residual Deviance: 417.44956
AIC: 425.44956
```

(b) Repeat the analysis with the pneumonoconiosis status being treated as ordinal.

Solution: First we convert status into an ordered factor (take care to get the order correct), then use the polr function.

```
> # b
> pneumo2$status <- ordered(pneumo2$status, levels=c("normal", "mild", "severe"))
> library(MASS)
> omod <- polr(status ~ year, pneumo2)
> summary(omod)
```

```
Re-fitting to get Hessian
Call:
polr(formula = status ~ year, data = pneumo2)
Coefficients:
      Value Std. Error t value
year 0.095904 0.011938 8.0336
Intercepts:
              Value Std. Error t value
normal|mild 3.95584 0.40969
                                9.65576
mild|severe 4.86905 0.44110
                                  11.03833
Residual Deviance: 416.91883
AIC: 422.91883
Note: An alternative solution that will give the same answer using the original data is
pneumo$status <- ordered(pneumo$status, levels = c('normal', 'mild', 'severe'))</pre>
omod2 <- polr(status~year, data = pneumo, weights = Freq)</pre>
The fit looks good (Figure 2), and the AIC for this model is slightly smaller than that for the
multinomial logistic regression model, so we prefer it.
> plot(years, props[1,], col="red", ylim=c(0,1))
> points(years, props[2,], col="blue")
> points(years, props[3,], col="green")
> fitted <- predict(omod, newdata=list(year=years), type="probs")</pre>
> lines(years, fitted[,1], col="blue")
> lines(years, fitted[,2], col="red")
> lines(years, fitted[,3], col="green")
For a miner with 25 years exposure we have the following fitted probabilities
> predict(omod, newdata=list(year=25), type="probs")
                  mild
    normal
                             severe
```

5

0.826100955 0.096014743 0.077884302

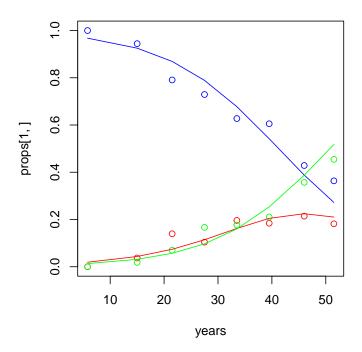


Figure 2: Fitted probability of the three outcomes vs length of service from the ordinal model