Multinomial logistic regression

- We use the multinom function from the nnet package to estimate a multinomial logistic regression model.
- Remark There are other functions in other R packages capable of multinomial regression, such as the **mlogit** package.
- The multinom function does not require the data to be reshaped (as the mlogit package does)
- ► (Similar format to example code found in Hilbe's *Logistic Regression Models*).

Multinomial logistic regression

- We must choose the level of our outcome that we wish to use as our **baseline** and specify this in the relevel function
 - Let's choose "academic".
- Then, we run our model using multinom.
- ▶ The multinom command does not include p-value calculation for the regression coefficients
- (We can calculate p-values using Wald tests or z-tests).

```
ml$prog2 <- relevel(ml$prog, ref = "academic")
test <- multinom(prog2 ~ ses + write, data = ml)

# weights: 15 (8 variable)
initial value 219.722458
iter 10 value 179.982880
final value 179.981726
converged</pre>
```

```
. . . .
```

```
Std. Errors:
```

```
(Intercept) sesmiddle seshigh write
general 1.166 0.4437 0.5142 0.02141
vocation 1.164 0.4764 0.5956 0.02222
```

Residual Deviance: 360

AIC: 376

Wald Test

- ▶ The Wald test in the context of logistic regression is used to determine whether a certain predictor variable X is significant or not. It rejects the null hypothesis of the corresponding coefficient being zero.
- ► The test consists of dividing the value of the coefficient by standard error

```
# Coefficients Divided by Standard Errors
# Then Compute p-values.
# 2-tailed z test
# p.values
```

```
(Intercept) sesmiddle seshigh write
general 1.448e-02 0.2294 0.02374 6.819e-03
vocation 7.299e-06 0.5408 0.09895 3.176e-07
```

- Remark: Some output is generated by running the model, even though we are assigning the model to a new R object.
- ► This model-running output includes some iteration history and includes the final **negative log-likelihood** (+ 179.981726).
- ► This value multiplied by two is then seen in the model summary as the **Residual Deviance** and it can be used in comparisons of nested models (360).

- As with many summary outputs, the output contains a column of coefficients and a column of standard errors.
- ► Each of these blocks has one row of values corresponding to a model equation.
- Focusing on the block of coefficients, we can look at the first row comparing prog = "general" to our baseline prog = "academic" and the second row comparing prog = "vocation" to our baseline prog = "academic".

▶ If we consider our coefficients from the first row to be b_1 and our coefficients from the second row to be b_2 , we can write our model equations:

$$In\left(\frac{P(prog = gen.)}{P(prog = acad.)}\right) = b_{10} + b_{11}(ses = 2) + b_{12}(ses = 3) + b_{13}write$$

$$In\left(\frac{P(prog = voc.)}{P(prog = acad.)}\right) = b_{20} + b_{21}(ses = 2) + b_{22}(ses = 3) + b_{23}write$$

- A one-unit increase in the variable write is associated with the decrease in the log odds of being in general program vs. academic program in the amount of 0.058 (b₁₃).
- A one-unit increase in the variable write is associated with the decrease in the log odds of being in vocation program vs. academic program in the amount of 0.1136 (b₂₃).

- ► The log odds of being in general program vs. in academic program will decrease by 1.163 if moving from ses="low" to ses="high" b₁₂.
- ► The log odds of being in general program vs. in academic program will decrease by 0.533 if moving from ses="low" to ses="middle" b₁₁, although this coefficient is not significant.

- ► The log odds of being in vocation program vs. in academic program will decrease by 0.983 if moving from ses="low" to ses="high"(b₂₂).
- ► The log odds of being in vocation program vs. in academic program will increase by 0.291 if moving from ses="low" to ses="middle"(b₂₁), although this coefficient is not signficant.