# BST 210 Homework 6

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```
library(foreign)
library(dplyr)

##

## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':

##

## filter, lag

## The following objects are masked from 'package:base':

##

## intersect, setdiff, setequal, union

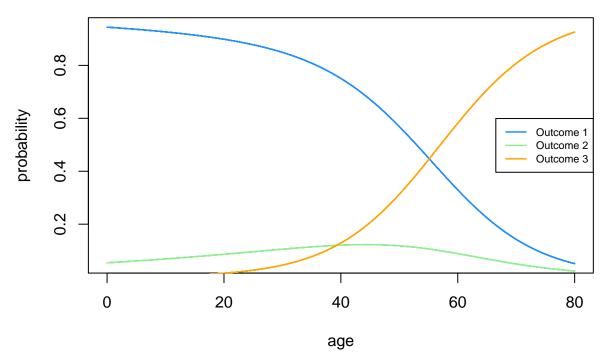
library(nnet)
```

### Problem 1

```
# Data cleaning
library(haven)
framingham = read_dta("Data and Programs/framingham.dta")
framingham = framingham[framingham$prevchd == 0,]
framingham$outcome = (framingham$death == 0 & framingham$anychd == 0)*1 +
  (framingham$death == 0 & framingham$anychd == 1)*2 + (framingham$death == 1)*3
framingham$prevchd = NULL
framingham$sex = framingham$sex -1
summ.MNfit <- function(fit, digits=3){</pre>
  s <- summary(fit)</pre>
  for(i in 2:length(fit$lev))
    cat("\nLevel", fit$lev[i], "vs. Level", fit$lev[1], "\n")
    betaHat <- s$coefficients[(i-1),]</pre>
    se <- s$standard.errors[(i-1),]</pre>
    zStat <- betaHat / se
    pval <- 2 * pnorm(abs(zStat), lower.tail=FALSE)</pre>
    RRR <- exp(betaHat)</pre>
    RRR.lo <- exp(betaHat - qnorm(0.975)*se)
    RRR.up <- exp(betaHat + qnorm(0.975)*se)
    results <- cbind(betaHat, se, pval, RRR, RRR.lo, RRR.up)
    print(round(results, digits=digits))
  }
```

```
model.sex = multinom(outcome~sex, data = framingham)
## # weights: 9 (4 variable)
## initial value 4658.116104
## iter 10 value 3904.409463
## iter 10 value 3904.409444
## final value 3904.409444
## converged
model.age = multinom(outcome~age, data = framingham)
## # weights: 9 (4 variable)
## initial value 4658.116104
## iter 10 value 3581.808246
## iter 10 value 3581.808246
## final value 3581.808246
## converged
model.age.sex = multinom(outcome~age+sex, data = framingham)
## # weights: 12 (6 variable)
## initial value 4658.116104
## iter 10 value 3509.585533
## final value 3509.484236
## converged
model.age.sex.int = multinom(outcome~age+sex+age*sex, data = framingham)
## # weights: 15 (8 variable)
## initial value 4658.116104
## iter 10 value 3508.375999
## final value 3505.913226
## converged
1(a)
age.seq = seq(0,80,0.01)
prob.model.age = predict(model.age,list(age = age.seq), type = 'probs')
par(mfrow = c(1,1))
plot(age.seq, prob.model.age[,1], cex = 0.05, main = "Probability of Outcomes",
     xlab ="age",ylab = "probability", col = "dodgerblue")
points(age.seq, prob.model.age[,2], cex = 0.05, col = "lightgreen")
points(age.seq, prob.model.age[,3], cex = 0.05, col = "orange")
legend("right",c("Outcome 1", "Outcome 2", "Outcome 3"),
       col = c("dodgerblue","lightgreen","orange"), lwd = 1.5, cex = 0.7)
```

# **Probability of Outcomes**



As age goes up, the estimated probability of outcome 1d (no death or chronic heart disease in the follow-up period) decreases. The probability of outcome 2 (chronic heart disease, but remained alive) follows a parabolic pattern, and reaches maximum at around 40 yo. The probability of outcome 3 (death) gets larger as age goes up. According to the fitted probability curves, the probability of outcome 2 (CHD) is overall lower than outcome 1 (no chd or death) and outcome 3 (death). For people older than 55 years old, the prevalent outcome is death and for people younger than 55 years old, the prevalent outcome is no death or chd.

```
summ.MNfit(model.age)
```

```
##
## Level 2 vs. Level 1
##
               betaHat
                           se pval
                                     RRR RRR.lo RRR.up
                -2.859 0.312
##
   (Intercept)
                                 0 0.057
                                          0.031
                                                 0.106
## age
                 0.026 0.006
                                 0 1.026
                                          1.013
                                                 1.039
##
## Level 3 vs. Level 1
##
               betaHat
                           se pval
                                     RRR RRR.lo RRR.up
## (Intercept)
                -6.424 0.244
                                 0 0.002
                                          0.001
                                                 0.003
## age
                 0.117 0.005
                                 0 1.124
                                          1.113
                                                 1.134
confint(model.age)
##
##
```

```
## (Intercept) -6.9032729 -5.9455880
                0.1073283 0.1257421
## age
vcov(model.age)
##
                 2:(Intercept)
                                        2:age 3:(Intercept)
## 2:(Intercept) 0.0972763186 -1.965591e-03 0.0180569339 -3.734082e-04
## 2:age
                 -0.0019655907 4.080459e-05 -0.0003756535 7.955112e-06
## 3:(Intercept) 0.0180569339 -3.756535e-04 0.0596882838 -1.134172e-03
                 -0.0003734082 7.955112e-06 -0.0011341722 2.206629e-05
## 3:age
beta3 2 std = sqrt(4.080459e-05 + 2.206629e-05 - 2*7.955112e-06)
beta3 2 std
## [1] 0.006852785
lower = \exp(0.117*10 - 0.026*10 - 1.96 * beta3_2_std *10)
upper = \exp(0.117*10 - 0.026*10+ 1.96 * beta3_2_std *10)
sprintf("The estimated relative risk ratio of having outcome 2 to having outcome 1 for a population is
        \exp(0.026*10), \exp(0.01326*10), \exp(0.0383*10))
## [1] "The estimated relative risk ratio of having outcome 2 to having outcome 1 for a population is 1
sprintf("The estimated relative risk ratio of having outcome 3 to having outcome 1 for a population is
        \exp(0.117*10), \exp(0.1073*10), \exp(0.12574*10))
## [1] "The estimated relative risk ratio of having outcome 3 to having outcome 1 for a population is 3
sprintf("The estimated relative risk ratio of having outcome 3 to having outcome 2 for a population is '
        \exp(0.117*10)/\exp(0.026*10), lower, upper)
## [1] "The estimated relative risk ratio of having outcome 3 to having outcome 2 for a population is 2
1(b)
female = fitted(model.sex)[framingham$sex == 1,][1,]
male = fitted(model.sex)[framingham$sex == 0,][1,]
fitted_prob_table_sex = rbind(male,female)
outcome_sex_table = table(framingham$sex, framingham$outcome)
outcome_sex_table_prop = prop.table(outcome_sex_table, 1)
fitted_prob_table_sex
##
                                       3
                            2
                  1
        0.4697938 0.1236231 0.4065831
## female 0.6260492 0.0987825 0.2751683
outcome_sex_table_prop
##
##
                                       3
     0 0.46978022 0.12362637 0.40659341
##
     1 0.62603306 0.09876033 0.27520661
According to the tables above, we can confirm that for the model with sex alone, the fitted probabilities
match the outcome-sex tabulation exactly.
summ.MNfit(model.sex)
```

```
##
## Level 2 vs. Level 1
##
                                                   betaHat
                                                                                          se pval
                                                                                                                            RRR RRR.lo RRR.up
## (Intercept) -1.335 0.075
                                                                                                              0 0.263 0.227 0.305
## sex
                                                       -0.511 0.102
                                                                                                              0 0.600 0.491 0.733
##
## Level 3 vs. Level 1
##
                                                   betaHat
                                                                                          se pval
                                                                                                                                RRR RRR.lo RRR.up
## (Intercept) -0.145 0.050 0.004 0.865
                                                                                                                                                0.784
                                                                                                                                                                      0.955
                                                      -0.678 0.068 0.000 0.508 0.444 0.581
outcome_sex_table_prop
##
                                                                                             2
##
                                                                                                                                    3
                                                       1
                 0 0.46978022 0.12362637 0.40659341
##
                 1 0.62603306 0.09876033 0.27520661
##
RRR\_21 = (P(Y=2|female)/P(Y=1|female))/(P(Y=2|male)/P(Y=1|male)) = (0.09876033/0.62603306)/(0.12362637/0.4697802)
RRR\_31 = (P(Y=3|female)) / P(Y=1|female)) / (P(Y=3|male)) / P(Y=1|male)) = (0.27520661/0.62603306) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.40659341/0.4697802) / (0.406597802) / (0.406597802) / (0.406597802) / (0.406597802) / (0.4
# calculated RRRs from the tabulation
RRR_21 = (0.09876033/0.62603306)/(0.12362637/0.46978022)
RRR_21
## [1] 0.599472
RRR_31 = (0.27520661/0.62603306)/(0.40659341/0.46978022)
RRR_31
```

## [1] 0.5079208

According to the summary of the model with sex alone, relative risk ratio of outcome 2 to outcome 1 is 0.600, and relative risk ratio of outcome 3 to outcome 1 is 0.508. The calculated RRRs from the tabulation match the results.

## 1(c)

```
anova(model.age.sex, model.age.sex.int, test = "Chisq")
##
                     Model Resid. df Resid. Dev
                                                    Test
                                                            Df LR stat.
## 1
                                        7018.968
                                 8474
                                                            NΑ
                                                                     NΑ
                 age + sex
                                 8472
                                        7011.826 1 vs 2
                                                             2 7.14202
## 2 age + sex + age * sex
##
        Pr(Chi)
## 1
             NA
## 2 0.02812743
```

The LRT statistic has a p-value = 0.028 (p<0.05). Therefore, we can reject the reduced model and conclude that the model including age,sex, and their interaction performs better than the one without interaction. We can consider fitting models with non-linear age terms in our next step.

#### Problem 2

```
library(VGAM)

## Loading required package: stats4

## Loading required package: splines
```

```
library(stats4)
library(splines)
ord.age = vglm(outcome~age,cumulative(parallel=TRUE, reverse=TRUE), data = framingham)
ord.sex = vglm(outcome~sex,cumulative(parallel=TRUE, reverse=TRUE), data = framingham)
ord.age.sex = vglm(outcome~age + sex,cumulative(parallel=TRUE, reverse=TRUE), data = framingham)
ord.age.sex.int = vglm(outcome~age + sex + age*sex,cumulative(parallel=TRUE, reverse=TRUE), data = fram
2(a)
# model with age alone
summary(ord.age)
##
## Call:
## vglm(formula = outcome ~ age, family = cumulative(parallel = TRUE,
      reverse = TRUE), data = framingham)
##
## Pearson residuals:
##
                                  1Q Median
                         Min
                                                 30
                                                      Max
## logitlink(P[Y>=2]) -2.036 -0.6986 -0.4423 0.4974 3.733
## logitlink(P[Y>=3]) -2.827 -0.4269 -0.2742 0.7561 2.979
## Coefficients:
##
                  Estimate Std. Error z value Pr(>|z|)
## (Intercept):1 -5.170906
                             0.202854 -25.49
                                                <2e-16 ***
## (Intercept):2 -5.713239
                             0.206504 -27.67
                                                <2e-16 ***
                  0.099351
                             0.003977
                                        24.98
## age
                                                <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Names of linear predictors: logitlink(P[Y>=2]), logitlink(P[Y>=3])
##
## Residual deviance: 7208.382 on 8477 degrees of freedom
##
## Log-likelihood: -3604.191 on 8477 degrees of freedom
## Number of Fisher scoring iterations: 4
## No Hauck-Donner effect found in any of the estimates
##
## Exponentiated coefficients:
        age
## 1.104454
confint(ord.age)
##
                      2.5 %
                                97.5 %
## (Intercept):1 -5.5684928 -4.7733183
## (Intercept):2 -6.1179799 -5.3084974
                  0.0915555 0.1071468
## age
sprintf("The estimated odds ratio for the effect of 10 years comparing outcome 3 vs. outcome 1 and 2 (
        \exp(0.099351*10), \exp(0.0915555*10), \exp(0.1071468*10))
```

```
## [1] "The estimated odds ratio for the effect of 10 years comparing outcome 3 vs. outcome 1 and 2 (c
sprintf("The estimated odds ratio for the effect of 10 years comparing outcome 2 and 3(combined) vs. o
        \exp(0.099351*10), \exp(0.0915555*10), \exp(0.1071468*10))
## [1] "The estimated odds ratio for the effect of 10 years comparing outcome 2 and 3(combined) vs. ou
2(b)
framingham$outcome1 = 1*(framingham$outcome == 3)
framingham$outcome2 = 1*(framingham$outcome == 2 | framingham$outcome == 3)
log.outcome1 = glm(outcome1~age, family = binomial(), data = framingham)
log.outcome2 = glm(outcome2~age, family = binomial(), data = framingham)
summary(log.outcome1)
##
## Call:
## glm(formula = outcome1 ~ age, family = binomial(), data = framingham)
## Deviance Residuals:
##
      Min
                1Q
                     Median
                                          Max
## -1.7719 -0.8255 -0.5540
                             0.9966
                                       2.3034
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -6.384296
                          0.237005 -26.94
                                             <2e-16 ***
               0.111896
                          0.004524
                                    24.73
                                             <2e-16 ***
## age
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 5387.4 on 4239 degrees of freedom
## Residual deviance: 4652.9 on 4238 degrees of freedom
## AIC: 4656.9
##
## Number of Fisher Scoring iterations: 4
summary(log.outcome2)
##
## Call:
## glm(formula = outcome2 ~ age, family = binomial(), data = framingham)
##
## Deviance Residuals:
      Min
                1Q
                    Median
                                   3Q
                                          Max
## -1.8075 -0.9741 -0.6882
                             1.0812
                                       1.9261
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -4.78576
                          0.20846 -22.96
                                            <2e-16 ***
## age
               0.09121
                          0.00411
                                    22.19
                                            <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
```

```
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 5818.8 on 4239
                                       degrees of freedom
## Residual deviance: 5256.7 on 4238
                                       degrees of freedom
## AIC: 5260.7
##
## Number of Fisher Scoring iterations: 4
confint(log.outcome1)
## Waiting for profiling to be done...
                    2.5 %
##
                              97.5 %
## (Intercept) -6.8535984 -5.9243745
## age
                0.1031075 0.1208437
confint(log.outcome2)
## Waiting for profiling to be done...
                     2.5 %
## (Intercept) -5.19745098 -4.38019036
                0.08320882 0.09932408
## age
```

The two beta coefficients for age in the two logistic regression models are 0.111896 and 0.09121 respectively, which are not close to each other. And the 95% CIs are (0.1031075, 0.1208437) and (0.08320882, 0.09932408) respectively, which do not overlap. Therefore, I suggest that proportional odds assumption doesn't hold for the ordinal logistic regression model with age alone.

# **2**(c)

```
summary(ord.sex)
```

```
##
## Call:
## vglm(formula = outcome ~ sex, family = cumulative(parallel = TRUE,
      reverse = TRUE), data = framingham)
##
## Pearson residuals:
##
                          Min
                                   1Q Median
                                                  3Q
                                                       Max
## logitlink(P[Y>=2]) -0.9474 -0.6896 -0.6896 0.6873 2.359
## logitlink(P[Y>=3]) -1.8868 -0.4672 -0.3546 1.0803 1.483
##
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept):1 0.10961
                          0.04578
                                      2.394
                                              0.0166 *
## (Intercept):2 -0.36488
                            0.04612 -7.912 2.53e-15 ***
                -0.61834
                            0.06081 -10.168 < 2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Names of linear predictors: logitlink(P[Y>=2]), logitlink(P[Y>=3])
##
## Residual deviance: 7810.012 on 8477 degrees of freedom
##
## Log-likelihood: -3905.006 on 8477 degrees of freedom
##
```

```
## Number of Fisher scoring iterations: 3
##
## No Hauck-Donner effect found in any of the estimates
##
##
## Exponentiated coefficients:
##
         sex
## 0.5388392
# model with sex alone
tab1 = table(framingham$outcome1, framingham$sex)
tab2 = table(framingham$outcome2, framingham$sex)
##
##
          0
                1
##
     0 1080 1754
##
     1 740 666
tab2
##
##
          0
                1
##
        855 1515
##
     1
       965 905
OR_{12_3} = (666/1754)/(740/1080)
OR 12 3
## [1] 0.5541619
OR_1_23 = (905/1515)/(965/855)
OR_1_23
## [1] 0.5292669
The associated odds ratio estimates are 0.554169 ((1,2) vs. 3) and 0.5292669 (1 vs (2,3)) are close to the ordinal
```

The associated odds ratio estimates are 0.554169 ((1,2) vs. 3) and 0.5292669 (1 vs. (2,3)) are close to the ordinal logistic regression-based odds ratio estimate for sex: 0.5388392. Therefore, I suggest that the proportional odds model assumption holds for the ordinal logistic regression model with sex alone. However, in the ordinal logistic regression model, there are 2 covariate patterns (female or male) and 3 parameters, indicating that the ordinal logistic regression model for sex alone is not saturated.

```
2(d)
pchisq(deviance(ord.age.sex)-deviance(ord.age.sex.int),
       df = df.residual(ord.age.sex)-df.residual(ord.age.sex.int),lower.tail=F)
## [1] 0.235037
summary(ord.age.sex.int)
##
## Call:
  vglm(formula = outcome ~ age + sex + age * sex, family = cumulative(parallel = TRUE,
##
       reverse = TRUE), data = framingham)
##
## Pearson residuals:
##
                         Min
                                  1Q Median
## logitlink(P[Y>=2]) -2.398 -0.6834 -0.4094 0.4837 4.155
```

```
## logitlink(P[Y>=3]) -3.177 -0.4147 -0.2630 0.7150 3.617
##
## Coefficients:
                  Estimate Std. Error z value Pr(>|z|)
##
## (Intercept):1 -5.250835
                             0.306883 -17.110
                                                <2e-16 ***
  (Intercept):2 -5.811767
                             0.309567 - 18.774
                                                <2e-16 ***
## age
                  0.110043
                             0.006173 17.828
                                                <2e-16 ***
## sex
                 -0.315679
                             0.415731
                                      -0.759
                                                 0.448
                 -0.009686
                             0.008155
                                      -1.188
                                                 0.235
## age:sex
##
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Names of linear predictors: logitlink(P[Y>=2]), logitlink(P[Y>=3])
##
## Residual deviance: 7055.417 on 8475 degrees of freedom
##
## Log-likelihood: -3527.708 on 8475 degrees of freedom
##
## Number of Fisher scoring iterations: 4
##
## No Hauck-Donner effect found in any of the estimates
##
##
## Exponentiated coefficients:
##
         age
                   sex
                         age:sex
## 1.1163259 0.7292937 0.9903608
```

By comparing the ordinal logistic regrssion model with age  $\times$  sex interaction and without age  $\times$  sex interaction with a "likelihood ratio" test, we get p-value = 0.2305 (>0.05). Furthermore, according to Wald test, in ordinal logistic regrssion model with age  $\times$  sex interaction, the interaction term is not significant. Therefore, we fail to reject the null hypothesis and conclude that the age  $\times$  sex interaction is not necessary for ordinal logistic regrssion modeling.

# **2**(e)

```
ord.po = vglm(outcome~age+sex, family = cumulative(parallel = TRUE,
    reverse = TRUE), data = framingham)
ord.npo = vglm(outcome~age+sex,family = cumulative(parallel = FALSE,
    reverse = TRUE), data = framingham)
pchisq(deviance(ord.po) - deviance(ord.npo),
    df = df.residual(ord.po) - df.residual(ord.npo), lower.tail = F)
```

#### ## [1] 1.423149e-09

By comparing the model with proportional odds assumption and the one without with a "likelihood ratio" test, we get p-value = 1.423149e-09 (<0.05). Therefore, we reject the null hypothesis and conclude that porportional odds assumption does not hold for the model including both effects of age and sex. I would recommend using multinomial logistic regression if we wanted to include continuous age in the modeling. On top of that, we can add quadratic term of age to the model and assess whether there is nonlinear relationship between age and outcome.