

## stsci4110hw6

Nick Gembs

11/29/2022

1.

a.  $\pi_{+1} - \pi_{1+}$  is the probability of Bryant making the second free throw minus the probability of Bryant making the first free throw. A larger absolute value of this number indicates dependence of the first on the second free throw.

b.

$H_0: \pi_{12} = \pi_{21} \equiv H_0: \pi_{+1} = \pi_{1+}$   $H_A: \pi_{12} \neq \pi_{21} \equiv H_0: \pi_{+1} \neq \pi_{1+}$

```
M = (33-37)^2/(33+37)
1-pchisq(M,1,lower.tail = T)
```

```
## [1] 0.6325851
```

p-value > .05, fail to reject the null.

Conclude: Table margins are homogeneous.  $\pi_{12} = \pi_{21} \equiv H_0: \pi_{+1} = \pi_{1+}$  The probability of Bryant making the second free throw does not change based on the outcome of the first free throw

c.

```
library(PropCIs)
diffpropci.mp(33, 37, 230, conf.level = 0.95)
```

```
##
##
##
## data:
##
## 95 percent confidence interval:
## -0.05390917 0.08839193
## sample estimates:
## [1] 0.01724138
```

0 is included in the 95% CI. Therefore, there is not enough evidence to prove non-homogeneity.

## 2.

a.

```
df = read.csv("C:/Users/Nick/Downloads/afterlife2.csv")

df$belief = factor(df$belief, ordered = F)
df$race = factor(df$race, ordered = F)
df$gender = factor(df$gender, ordered = F)
df$belief = relevel(df$belief, ref = "No")

library(tidyr)

df = uncount(df, count, .remove = TRUE, .id = NULL)

library(nnet)

## Warning: package 'nnet' was built under R version 4.2.2

bcl = multinom(belief ~ race+gender+religiosity, data = df)

## # weights: 15 (8 variable)
## initial value 1231.544376
## iter 10 value 754.526566
## final value 751.751592
## converged

summary(bcl)

## Call:
## multinom(formula = belief ~ race + gender + religiosity, data = df)
##
## Coefficients:
## (Intercept) raceWhite genderMale religiosity
## Undecided -0.6942144 0.1944430 -0.1445318 0.06299078
## Yes -1.0998846 0.3690174 -0.4902732 0.89822127
##
## Std. Errors:
## (Intercept) raceWhite genderMale religiosity
## Undecided 0.3194276 0.2635508 0.2306095 0.10141679
## Yes 0.2517191 0.2006170 0.1766465 0.07473111
##
## Residual Deviance: 1503.503
## AIC: 1519.503
```

b.

```
# Undecided = -0.6942144 + 0.1944430*raceWhite + -0.1445318*genderMale +  
0.06299078*religiosity
```

```
# Yes = -1.0998846 + 0.3690174*raceWhite + -0.4902732*genderMale +  
0.89822127*religiosity
```

```
# odds of "yes" versus "no" for the two genders
```

```
odds = exp(-0.4902732)
```

```
odds
```

```
## [1] 0.612459
```

```
cat("being male has a 0.612459 multiplicative effect on odds of being 'yes'  
belief over 'no'\n")
```

```
## being male has a 0.612459 multiplicative effect on odds of being 'yes'  
belief over 'no'
```

c.

```
# odds of "yes" versus "no" for the two races
```

```
odds = exp(0.3690174)
```

```
odds
```

```
## [1] 1.446313
```

```
cat("being white has a 1.446313 multiplicative effect on odds of being 'yes'  
belief over 'no'\n")
```

```
## being white has a 1.446313 multiplicative effect on odds of being 'yes'  
belief over 'no'
```

```
# change in odds of "yes" versus "no" for a one unit increase in religiosity
```

```
odds = exp(0.89822127)
```

```
odds
```

```
## [1] 2.455232
```

```
cat("A one unit increase in religiosity has a 2.455232 multiplicative effect  
on odds of being 'yes' belief over 'no'\n")
```

```
## A one unit increase in religiosity has a 2.455232 multiplicative effect on  
odds of being 'yes' belief over 'no'
```

d.

```
library(lmtest)

## Loading required package: zoo

##
## Attaching package: 'zoo'

## The following objects are masked from 'package:base':
##
##      as.Date, as.Date.numeric

#H0 : gender is not significant in predicting belief
#H1 : gender is significant in predicting belief
bclg = multinom(belief ~ race+religiosity, data = df)

## # weights: 12 (6 variable)
## initial value 1231.544376
## iter 10 value 756.011597
## final value 756.008290
## converged

lrtest(bcl,bclg)

## Likelihood ratio test
##
## Model 1: belief ~ race + gender + religiosity
## Model 2: belief ~ race + religiosity
##   #Df LogLik Df  Chisq Pr(>Chisq)
## 1    8 -751.75
## 2    6 -756.01 -2  8.5134    0.01417 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# p-value<.05, reject null
```

e.

```
#H0 : race is not significant in predicting belief
#H1 : race is significant in predicting belief
bclrace = multinom(belief ~ gender+religiosity, data = df)

## # weights: 12 (6 variable)
## initial value 1231.544376
## iter 10 value 753.442157
## final value 753.432581
## converged

lrtest(bcl,bclrace)
```

```

## Likelihood ratio test
##
## Model 1: belief ~ race + gender + religiosity
## Model 2: belief ~ gender + religiosity
##   #Df  LogLik Df Chisq Pr(>Chisq)
## 1    8 -751.75
## 2    6 -753.43 -2 3.362    0.1862

# p-value>.05, fail to reject null

f.

#H0 : religiosity is not significant in predicting belief
#H1 : religiosity is significant in predicting belief
bclr = multinom(belief ~ race+gender, data = df)

## # weights:  12 (6 variable)
## initial  value 1231.544376
## iter   10 value 882.187101
## final   value 882.187030
## converged

lrtest(bcl,bclr)

## Likelihood ratio test
##
## Model 1: belief ~ race + gender + religiosity
## Model 2: belief ~ race + gender
##   #Df  LogLik Df  Chisq Pr(>Chisq)
## 1    8 -751.75
## 2    6 -882.19 -2 260.87  < 2.2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

# p-value<.05, reject null

g.

bclrace = multinom(belief ~ gender+religiosity, data = df)

## # weights:  12 (6 variable)
## initial  value 1231.544376
## iter   10 value 753.442157
## final   value 753.432581
## converged

summary(bclrace)

## Call:
## multinom(formula = belief ~ gender + religiosity, data = df)
##

```

```
## Coefficients:
##           (Intercept) genderMale religiosity
## Undecided  -0.5578253 -0.1295137  0.06309689
## Yes        -0.8341452 -0.4617498  0.89802444
##
## Std. Errors:
##           (Intercept) genderMale religiosity
## Undecided   0.2591456  0.2296061  0.10138932
## Yes         0.2049019  0.1754738  0.07460479
##
## Residual Deviance: 1506.865
## AIC: 1518.865

# Undecided = -0.5578253 -0.1295137*genderMale + 0.06309689*religiosity
# Yes =      -0.8341452 -0.4617498*genderMale + 0.89802444* religiosity

h.

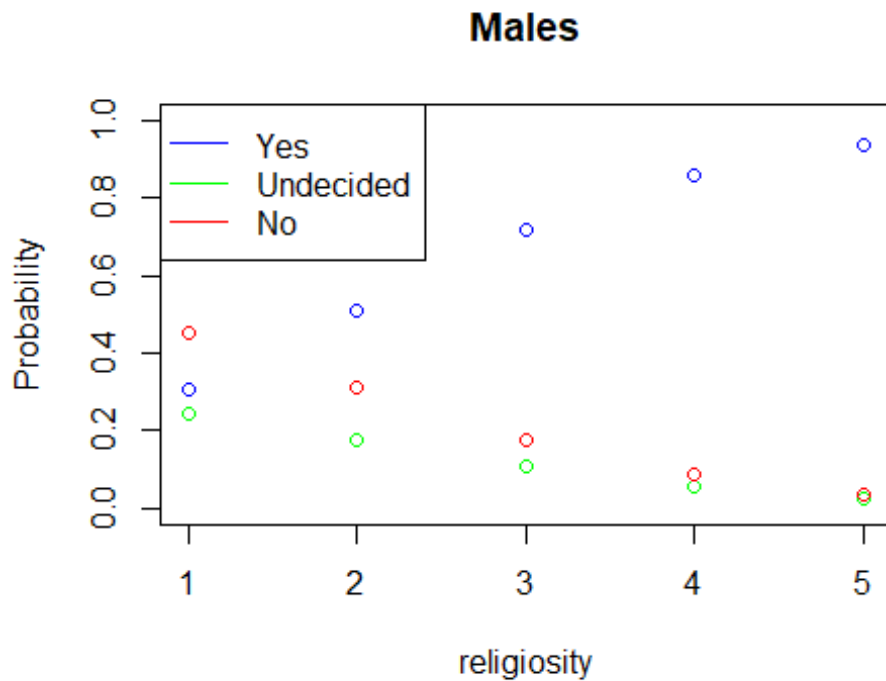
newdf = data.frame(df$religiosity, fitted(bclrace), df$gender)

plot(newdf$df.religiosity[newdf$df.gender=="Male"],
newdf$Yes[newdf$df.gender=="Male"], type = "p", col="blue", xlab =
"religiosity", ylab = "Probability", xlim = c(1,5), ylim = c(0,1), main =
"Males")

legend("topleft", lty = c(1,1,1), pch = c(-1,-1,-1), col = c("blue", "green",
"red"), legend = c("Yes","Undecided", "No"))

points(newdf$df.religiosity[newdf$df.gender=="Male"],
newdf$Undecided[newdf$df.gender=="Male"], col="green")

points(newdf$df.religiosity[newdf$df.gender=="Male"],
newdf$No[newdf$df.gender=="Male"], col="red")
```

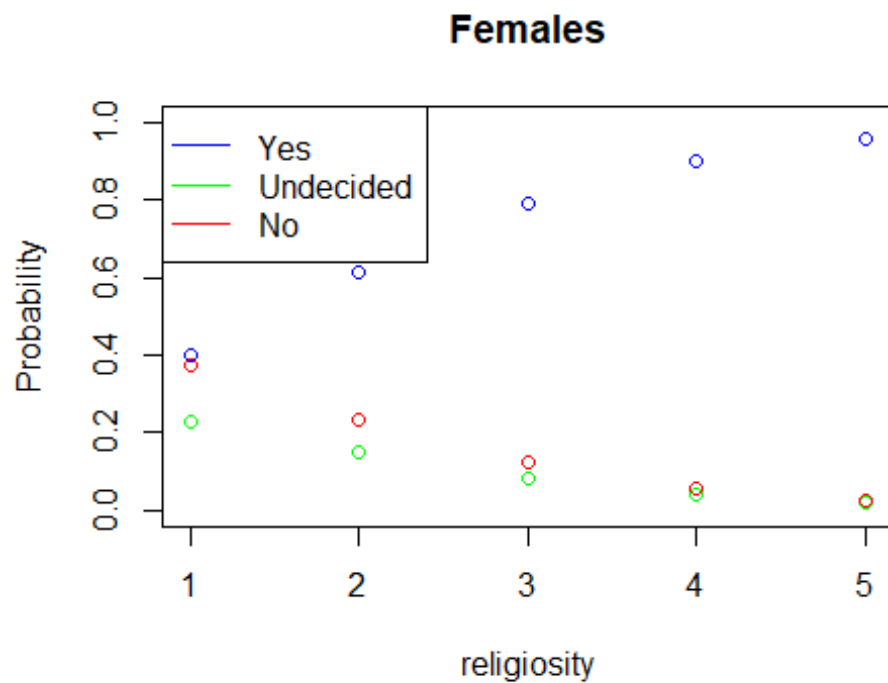


```
plot(newdf$df.religiosity[newdf$df.gender=="Female"],
newdf$Yes[newdf$df.gender=="Female"], type = "p", col="blue", xlab =
"religiosity", ylab = "Probability", xlim = c(1,5), ylim = c(0,1), main =
"Females")

legend("topleft", lty = c(1,1,1), pch = c(-1,-1,-1), col = c("blue", "green",
"red"), legend = c("Yes","Undecided", "No"))

points(newdf$df.religiosity[newdf$df.gender=="Female"],
newdf$Undecided[newdf$df.gender=="Female"], col="green")

points(newdf$df.religiosity[newdf$df.gender=="Female"],
newdf$No[newdf$df.gender=="Female"], col="red")
```



Non religious males are less likely to believe than non religious females. At levels of high religiosity, both males and females are very likely to believe.

i.

```
#sample(newdf[newdf$df.religiosity == 3,])
```

|        | Belief in Afterlife |           |     |
|--------|---------------------|-----------|-----|
| Gender | Yes                 | Undecided | No  |
| Male   | .72                 | .11       | .18 |
| Female | .79                 | .09       | .12 |



3.

- a.  $\text{logit}(P(Y \leq 1)) = \alpha_1 + \beta_1 * \text{therapyseq} + \beta_2 * \text{genderm}$   
 $\text{logit}(P(Y \leq 2)) = \alpha_2 + \beta_1 * \text{therapyseq} + \beta_2 * \text{genderm}$   
 $\text{logit}(P(Y \leq 3)) = \alpha_3 + \beta_1 * \text{therapyseq} + \beta_2 * \text{genderm}$

$$\begin{aligned}\text{logit}(P(Y \leq 1)) &= -2.3652 + 0.5587 * \text{therapyseq} + 0.5299 * \text{genderm} \\ \text{logit}(P(Y \leq 2)) &= -1.3041 + 0.5587 * \text{therapyseq} + 0.5299 * \text{genderm} \\ \text{logit}(P(Y \leq 3)) &= 0.1582 + 0.5587 * \text{therapyseq} + 0.5299 * \text{genderm}\end{aligned}$$

- b.  $e^{.5587} = 1.748$

Using sequential treatment over alternating will have a 1.748 multiplicative effect on the severity outcome.

- c. In this model, therapy type is a significant predictor in severity outcome, while gender is not. There is lower severity among those who use the alternating treatment method.

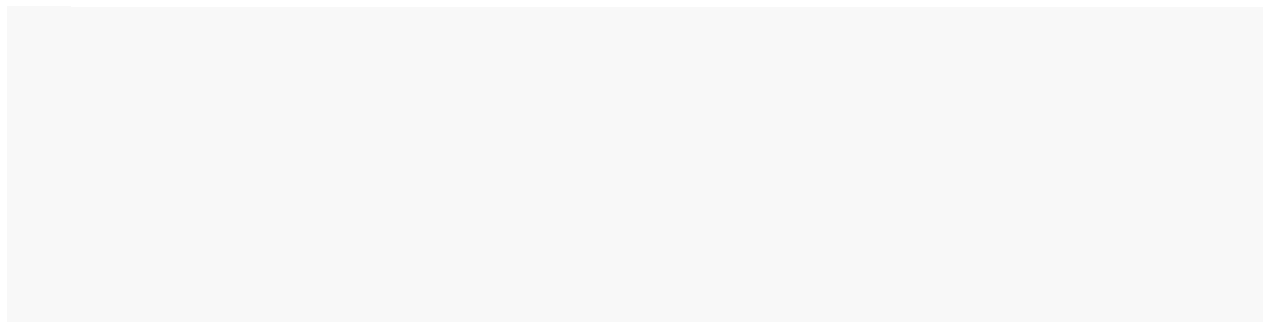
d.

.

11/29/22, 5:30 PM

Desmos | Scientific Calculator

|   |   |
|---|---|
| $p(y = 1)$ Alternating  |    |
| $\frac{e^{-2.3652 + 0.5587*0 + 0.5299*1}}{1 + e^{-2.3652 + 0.5587*0 + 0.5299*1}}$   | = 0.1376081021  |
| $p(y = 2)$  |    |
| $\frac{e^{-1.3041 + 0.5587*0 + 0.5299*1}}{1 + e^{-1.3041 + 0.5587*0 + 0.5299*1}} - \frac{e^{-2.3652 + 0.5587*0 + 0.5299*1}}{1 + e^{-2.3652 + 0.5587*0 + 0.5299*1}}$ | = 0.1779631609  |
| $p(y = 3)$  |    |
| $\frac{e^{-1.582 + 0.5587*0 + 0.5299*1}}{1 + e^{-1.582 + 0.5587*0 + 0.5299*1}} - 0.1779631609$  | = 0.4875809682  |
| $p(y = 4)$  |    |
| $1 - 0.4875809682$  | = 0.5124190318  |
| $p(y = 1)$ Sequential   |    |
| $\frac{e^{-2.3652 + 0.5587*1 + 0.5299*1}}{1 + e^{-2.3652 + 0.5587*1 + 0.5299*1}}$   | = 0.2181295346  |
| $p(y = 2)$  |  |
| $\frac{e^{-1.3041 + 0.5587*1 + 0.5299*1}}{1 + e^{-1.3041 + 0.5587*1 + 0.5299*1}} - \frac{e^{-2.3652 + 0.5587*1 + 0.5299*1}}{1 + e^{-2.3652 + 0.5587*1 + 0.5299*1}}$ | = 0.2282029991  |
| $p(y = 3)$  |  |
| $\frac{e^{-1.582 + 0.5587*1 + 0.5299*1}}{1 + e^{-1.582 + 0.5587*1 + 0.5299*1}} - 0.2282029991$  | = 0.5485424353  |
| $p(y = 4)$  |  |
| $1 - 0.5485424353$  | = 0.4514575647  |



4.

a.

```
Northeast = c(266,10,8,7)
```

```
Midwest = c(15,414,22,6)
```

```
South = c(61,50,578,27)
```

```
West = c(28,40,22,301)
```

```
df = data.frame(Northeast,Midwest,South,West, row.names = c("Northeast",  
"Midwest", "South", "West"))
```

```
mat = data.matrix(df)
```

```
mat
```

```
##           Northeast Midwest South West  
## Northeast      266      15    61   28  
## Midwest        10     414    50   40  
## South           8      22   578   22  
## West            7       6    27  301
```

```
sixteen = c()  
for (i in 1:4){  
  sixteen[i] = (sum(mat[i,]))/(sum(mat))  
}
```

```
twentyten = c()  
for (i in 1:4){  
  twentyten[i] = (sum(mat[,i]))/(sum(mat))  
}
```

```
sixteen
```

```
## [1] 0.1994609 0.2770889 0.3396226 0.1838275
```

```
twentyten
```

```
## [1] 0.1568733 0.2463612 0.3859838 0.2107817
```

b.

*#H0:  $\pi_{i+} = \pi_{+i}$*

*#for all  $i = 1, \dots, 4$ .*

*#HA: For at least one pair,  $\pi_{i+} \neq \pi_{+i}$*

```
library(coin)
```

```
## Warning: package 'coin' was built under R version 4.2.2
```

```
## Loading required package: survival

library(mvtnorm)
library(modeltools)

## Loading required package: stats4

mh_test(as.table(mat))

##
## Asymptotic Marginal Homogeneity Test
##
## data: response by
## conditions (Var1, Var2)
## stratified by block
## chi-squared = 86.236, df = 3, p-value < 2.2e-16

# p - value is less than .05, reject the null. The marginal probabilities are
# not all the same. That is, region
# preferences have shifted from age 16 to 2010.
```

## 5.

a.

```
ten2 = read.csv("C:/Users/Nick/Downloads/tennis-18-20v2.csv")

library(BradleyTerry2)

## Warning: package 'BradleyTerry2' was built under R version 4.2.2

tennisModel <- BTm(outcome = cbind(win1,win2), as.factor(first.player),
as.factor(second.player),
formula = ~ player, id = "player", data=ten2, refcat = "Kenin")
tennisModel

## Bradley Terry model fit by glm.fit
##
## Call: BTm(outcome = cbind(win1, win2), player1 = as.factor(first.player),
## player2 = as.factor(second.player), formula = ~player, id = "player",
## refcat = "Kenin", data = ten2)
##
## Coefficients:
## playerBarty playerHalep playerOsaka playerSWilliams
## 0.3771 1.2245 0.6105 0.9643
##
## Degrees of Freedom: 10 Total (i.e. Null); 6 Residual
## Null Deviance: 6.565
## Residual Deviance: 4.491 AIC: 23.84
```

```
cat("\nPlayers ranked:\n 1. Halep - 1.2245\n 2. Williams - .9643\n 3. Osaka - .6105\n 4. Barty - .3771\n 5. Kenin - 0(ref)")
```

```
##
```

```
## Players ranked:
```

```
## 1. Halep - 1.2245
```

```
## 2. Williams - .9643
```

```
## 3. Osaka - .6105
```

```
## 4. Barty - .3771
```

```
## 5. Kenin - 0(ref)
```

b.

```
# P(Halep beats Barty)
```

```
p = (exp(1.2245-.3771))/(1+exp(1.2245-.3771))
```

```
p
```

```
## [1] 0.7000214
```

```
cat("There is a 70% chance that Halep beats Barty")
```

```
## There is a 70% chance that Halep beats Barty
```

c.

```
# P(Kenin beats Williams)
```

```
p = (exp(0-.9643))/(1+exp(0-.9643))
```

```
p
```

```
## [1] 0.2760181
```

```
cat("There is a 27.6% chance that Kenin beats Williams")
```

```
## There is a 27.6% chance that Kenin beats Williams
```