Regression Model Comparison

Alejandro Arellano

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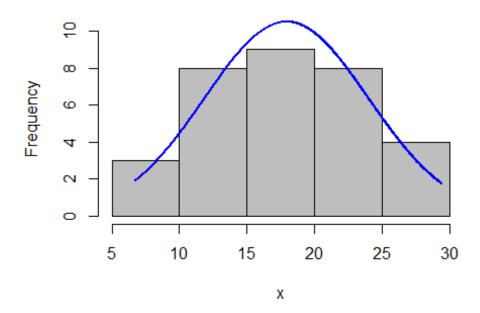
General Linear Model 1.2

```
# (a) Verify normality of the response variable, then fit the linear regressi
on model to the data. State the fitted model. Give estimates for all paramete
rs.

#Importing data for 1.2
WeightLoss<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise1.2Data.csv", header = TRUE, sep =
",")</pre>
```

#Checking for normality

plotNormalHistogram(WeightLoss\$EWL)



```
shapiro.test(WeightLoss$EWL)
##
## Shapiro-Wilk normality test
```

```
##
## data: WeightLoss$EWL
## W = 0.97424, p-value = 0.6234
#Reference Levels
drug.rel<- relevel(as.factor(WeightLoss$drug), ref="A")</pre>
gender.rel<- relevel(as.factor(WeightLoss$gender), ref="M")</pre>
#Creating general linear model
summary(fitted.model<- glm(EWL ~ drug.rel + age + gender.rel, data =</pre>
WeightLoss, family=gaussian(link=identity)))
##
## Call:
## glm(formula = EWL ~ drug.rel + age + gender.rel, family = gaussian(link =
identity),
##
       data = WeightLoss)
##
## Deviance Residuals:
       Min
                   1Q
                         Median
                                       3Q
                                                Max
                         0.0282
## -12.6861 -2.8772
                                   3.0932
                                            10.2343
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) 9.2146
                            5.6981
                                     1.617
                                             0.1171
## drug.relB
                4.8103
                            1.9988
                                     2.407
                                             0.0229 *
                                     0.967
                                             0.3420
## age
                0.1102
                            0.1140
## gender.relF 2.7235
                            1.9952
                                     1.365
                                             0.1831
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 31.44133)
##
       Null deviance: 1144.12 on 31 degrees of freedom
## Residual deviance: 880.36 on 28 degrees of freedom
## AIC: 206.88
##
## Number of Fisher Scoring iterations: 2
#To show estimated sigma
sigma(fitted.model)
## [1] 5.607257
```

(b) Which regression coefficients turn out to be significant at the 5%? Discuss goodness of fit of the #model.

Drug B is the only significant predictor in the model at the 5% significance level since the corresponding p-value is the only one under 0.05.

```
#Checking to see if the model fits using p-value and deviance test
null.model<- glm(EWL ~ 1, data=WeightLoss, family=gaussian(link=identity))
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))
## 'log Lik.' 8.386158 (df=2)
print(p.value<- pchisq(deviance, df=3, lower.tail=FALSE))
## 'log Lik.' 0.03867005 (df=2)</pre>
```

(c) Is one of the drugs more efficient for weight loss than the other? Interpret all estimated significant coefficients.

The estimated average EWL for subjects taking drug B is 4.8103 percent higher than that for subjects taking drug A, keeping all the other predictors fixed. It means that drug B is more efficient than drug A.

(d) According to the model, what is the predicted percent decrease in excess body weight for a 35- year old male who is taking drug A?

The predicted percent decrease in excess body weight for a 35-year old male who is taking drug A is computed by hand as: $EWL = 9.2146 + 0.1102 \cdot 35 = 13.0716$.

```
#Finding the predicted values using the model
print(predict(fitted.model, data.frame(drug.rel="A", age=35, gender.rel="M"))

## 1
## 13.07178
```

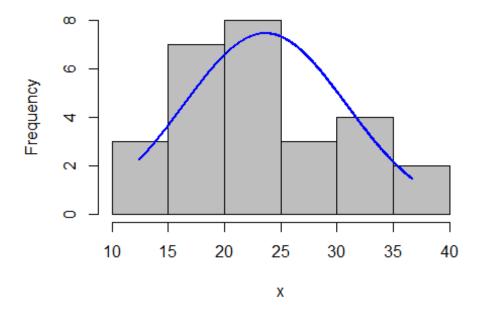
General Linear Model 1.3

```
# (a) Verify normality of the response variable, then fit the linear regressi
on model
#to the data. State the fitted model. Give estimates for all parameters.

#Importing data for 1.3
CarSales<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Ex
ercise Files/Exercise1.3Data.csv", header = TRUE, sep =",")

#Resecaling price
priceK<- CarSales$price/1000

#running normality check
plotNormalHistogram(priceK)</pre>
```



```
#Running Shapiro test
shapiro.test(priceK)
##
##
    Shapiro-Wilk normality test
##
## data: priceK
## W = 0.95482, p-value = 0.28
#Creating reference levels
bodystyle.rel<- relevel(as.factor(CarSales$bodystyle), ref="hatchback")</pre>
country.rel<- relevel(as.factor(CarSales$country), ref="Japan")</pre>
leather.rel<- relevel(as.factor(CarSales$leather), ref="no")</pre>
#Creating general linear model
summary(fitted.model<- glm(priceK ~ bodystyle.rel + country.rel + hwy + doors</pre>
leather.rel, data=CarSales, family=gaussian(link=identity)))
##
## Call:
## glm(formula = priceK ~ bodystyle.rel + country.rel + hwy + doors +
       leather.rel, family = gaussian(link = identity), data = CarSales)
##
##
## Deviance Residuals:
       Min
##
                 10
                      Median
                                    3Q
                                            Max
                       0.8821
## -4.4438 -2.4342
                                1.6428
                                         7.2350
```

```
##
## Coefficients:
##
                      Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                       5.1353
                                  5.5909
                                           0.919 0.36986
## bodystyle.relcoupe
                       2.2698
                                  2.0070
                                           1.131 0.27216
## bodystyle.relsedan
                                  1.8450
                       6.4107
                                           3.475 0.00254 **
## country.relGermany
                       3.1959
                                  2.0098
                                           1.590 0.12829
## country.relUSA
                       3.2128
                                  1.8812
                                           1.708 0.10394
                                           0.980 0.33937
## hwy
                       0.1305
                                  0.1332
## doors
                       1.5554
                                  0.7904
                                           1.968 0.06384 .
## leather.relyes
                                  1.9332
                                           6.298 4.79e-06 ***
                      12.1757
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 12.1319)
##
##
       Null deviance: 1357.08 on 26 degrees of freedom
## Residual deviance: 230.51 on 19 degrees of freedom
## AIC: 152.52
##
## Number of Fisher Scoring iterations: 2
#To show estimated sigma
sigma(fitted.model)
## [1] 3.483088
```

(b) How good is the model fit? Discuss significance of the regression coefficients.

The p-value in the deviance test is way below 0.05, indicating a good model fit. #Significant variables are sedan body style and leather interior.

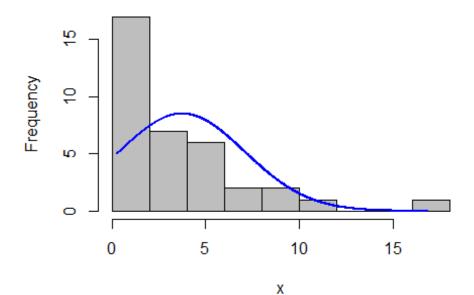
```
#Checking to see if the model fits using p-value and deviance test
null.model<- glm(priceK ~ 1, data=CarSales, family=gaussian(link=identity))
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))
## 'log Lik.' 47.86586 (df=2)
print(p.value<- pchisq(deviance, df=7, lower.tail = FALSE))
## 'log Lik.' 3.78218e-08 (df=2)</pre>
```

(c) Interpret the estimates of those regression coefficients that differ significantly from zero.

As estimated, sedan costs on average \$6,410.70 more than a hatchback, under all other equal conditions. The estimated average price of a car with leather interior is \$12,175.70 larger compared to a car withoutleather interior.

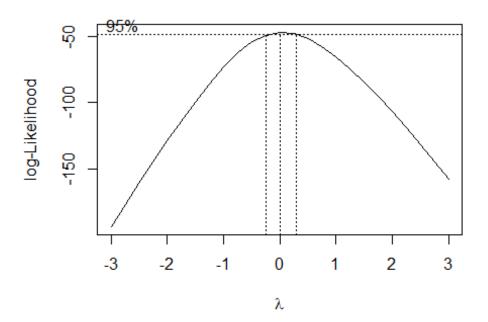
d) What is the predicted price of a sedan made in USA that has 4 doors, leather seats, and runs 30 mpg on highway? The predicted price of a sedan that is made in USA, has 4 doors, leather seats, and runs 30 mpg on highway is calculated as: price =

```
$1,000(5.1353 + 6.4107 + 3.2128 + 0.1305 \cdot 30 + 1.5554 \cdot 4 + 12.1757) =
      $37,071.10.
#Finding the predicted values using the model using fitted model for predicti
prediction<- (predict(fitted.model, data.frame(bodystyle.rel="sedan", country</pre>
.rel
="USA", hwy=30, doors=4, leather.rel="yes")))
print(prediction*1000)
##
## 37071.14
#Box Cox 2.1
#(a) Is the decrease in BMI percentile (preBMI-postBMI) normally distributed?
Plot a histogram and test for normality of the distribution.
#Importing data for 2.1
Obesity<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exe
rcise Files/Exercise2.1Data.csv",header=TRUE, sep=",")
#Finding the difference in BMI
BMIdiff<- Obesity$preBMI-Obesity$postBMI
#Checking for normality
```



plotNormalHistogram(BMIdiff)

```
#Shapiro Test for normality
shapiro.test(BMIdiff)
##
##
   Shapiro-Wilk normality test
##
## data: BMIdiff
## W = 0.79159, p-value = 1.114e-05
#(b) Find the optimal lambda for Box-Cox transformation. Transform the change
in BMI percentile (find the appropriate transformation in Table 2.1), and sho
w that the transformed variable is normally distributed. Plot the histogram a
nd do a formal testing.
#Finding indicator variables
female<- relevel(as.factor(Obesity$gender), ref="M")</pre>
control<- relevel(as.factor(Obesity$group), ref="Tx")</pre>
#Finding Lambda for box-cox
BoxCox.fit<- boxcox(BMIdiff ~ age + female + control, data=Obesity, lambda =</pre>
seq(-3,3,1/4), interp = FALSE)
```

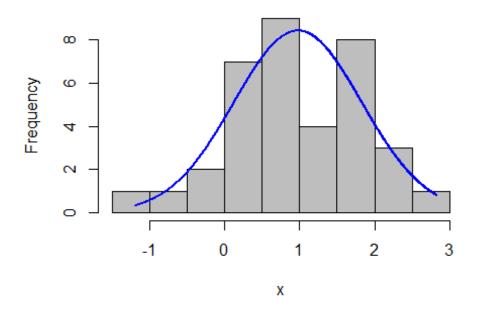


```
BoxCox.data<- data.frame(BoxCox.fit$x, BoxCox.fit$y)
ordered.data<- BoxCox.data[with(BoxCox.data, order(-BoxCox.fit.y)),]
ordered.data[1,]</pre>
```

```
## BoxCox.fit.x BoxCox.fit.y
## 13     0   -46.71658

#Applying the box-cox transformation with lambda from above
BMIdiff.tr<- log(BMIdiff)

#Running normality test again but on the transformed variables
plotNormalHistogram(BMIdiff.tr)</pre>
```



```
shapiro.test(BMIdiff.tr)
##
## Shapiro-Wilk normality test
##
## data: BMIdiff.tr
## W = 0.9877, p-value = 0.9532
#(c) Fit the general regression model to the Box-Cox transformed change in BM I percentile. Does this model have a good fit?
#Creating general linear model but on the transformed response
summary(fitted.model<- glm(BMIdiff.tr ~ age + female + control, data=Obesity,
family=gaussian(link=identity)))
##
## Call:
## glm(formula = BMIdiff.tr ~ age + female + control, family = gaussian(link)</pre>
```

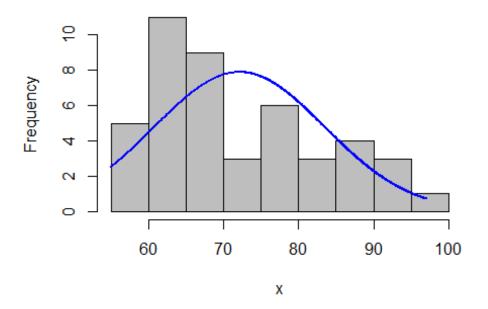
```
= identity),
       data = Obesity)
##
##
## Deviance Residuals:
        Min
                   10
                        Median
                                       3Q
                                                Max
## -1.31141 -0.41700 -0.08094
                                  0.41958
                                            1.47509
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.14375 0.45959
                                     2.489 0.018218 *
                                     1.372 0.179731
## age
               0.05008
                           0.03651
## femaleF
                           0.21708 -2.297 0.028317 *
               -0.49862
## controlCx -0.93835 0.22307 -4.207 0.000195 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 0.4187965)
##
##
       Null deviance: 25.432 on 35 degrees of freedom
## Residual deviance: 13.401 on 32 degrees of freedom
## AIC: 76.59
##
## Number of Fisher Scoring iterations: 2
#Estimated sigma
sigma(fitted.model)
## [1] 0.6471448
#Checking again to see if the model fits after transformation
null.model<- glm(BMIdiff.tr ~ 1, family=gaussian(link=identity))</pre>
#Printing deviance test for fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 23.06361 (df=2)
#Printing p-value for fit
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 3.916872e-05 (df=2)
#using fitted model for prediction
pred.BMIdiff.tr<- predict(fitted.model, data.frame(female="F", age=9,</pre>
control="Cx"))
print(exp(pred.BMIdiff.tr))
##
## 1.170609
```

```
#(a) Construct a histogram of the score. Does the distribution look normal? P
erform the test for normality. Draw conclusion.

#Importing data for 2.2
QIdata<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exer
cise Files/Exercise2.2Data.csv",header=TRUE, sep=",")

#Finding the difference in BMI
BMIdiff<- QIdata$preBMI-QIdata$postBMI

#Checking for normality
plotNormalHistogram(QIdata$score)</pre>
```



```
#Shapiro Test for normality
shapiro.test(QIdata$score)

##

## Shapiro-Wilk normality test
##

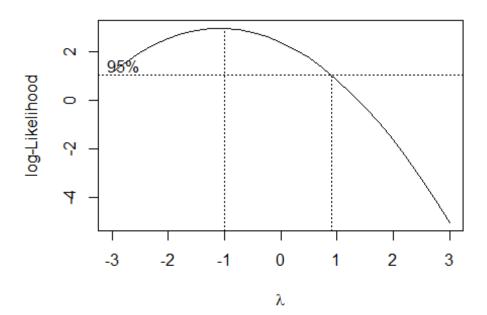
## data: QIdata$score
## W = 0.94357, p-value = 0.02913

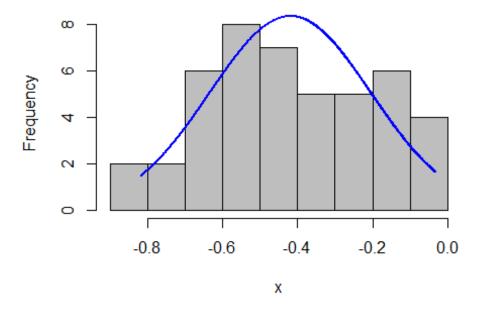
#(b) Transform the score variable using a meaningful Box-Cox transformation a
nd assure that it is now normally distributed by plotting the histogram and d
oing normality testing.

#Finding indicator variables and rescaling
```

```
desgn.rel<- relevel(as.factor(QIdata$desgn), ref="staff")
priorQI.rel<- relevel(as.factor(QIdata$priorQI), ref="no")
score<- QIdata$score/100

#Finding Lambda for box-cox
BoxCox.fit<- boxcox(score ~ desgn.rel + wrkyrs + priorQI.rel,
data=QIdata, lambda=seq(-3,3,1/4), interp = FALSE)</pre>
```





```
#Shapiro test for transformed variables
shapiro.test(score.tr)
##
##
   Shapiro-Wilk normality test
##
## data: score.tr
## W = 0.96606, p-value = 0.2073
#(c) Run the general linear regression model on the transformed score. What p
redictors are significant at the 0.05 level?
#Creating general linear model but on the transformed response
summary(fitted.model<- glm(score.tr ~ desgn.rel + wrkyrs + priorQI.rel,</pre>
data=QIdata, family=gaussian(link = identity)))
##
## Call:
## glm(formula = score.tr ~ desgn.rel + wrkyrs + priorQI.rel, family = gaussi
an(link = identity),
##
       data = QIdata)
##
## Deviance Residuals:
        Min
                         Median
                                                 Max
                   10
                                        30
## -0.39192 -0.15611 -0.00556
                                  0.14372
                                             0.36543
##
## Coefficients:
```

```
##
                   Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                  -0.609290 0.094150 -6.471 1.03e-07 ***
## desgn.reldoctor 0.179873
                             0.091507
                                        1.966
                                                0.0563 .
## desgn.relnurse 0.212200
                             0.087173
                                        2.434
                                                0.0195 *
## wrkyrs
                   0.000243
                             0.004086
                                        0.059
                                                0.9529
## priorQI.relyes 0.077263
                                                0.2649
                             0.068323
                                        1.131
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 0.04317095)
##
##
      Null deviance: 2.0290 on 44 degrees of freedom
## Residual deviance: 1.7268 on 40 degrees of freedom
## AIC: -7.0122
##
## Number of Fisher Scoring iterations: 2
#Estimated sigma
sigma(fitted.model)
## [1] 0.2077762
```

(d) Interpret the estimates of the significant beta coefficients. Does the model fit the data well? Conduct the chi-squared deviance test.

The estimated mean inverse transformed score for nurses is 0.2122 points above that for staff, and for doctors it is 0.1799 points above that for staff.

The model doesn't really fit the data well, as seen from the large p-value of the deviance test given below.

```
#checking model fit
null.model<- glm(score.tr ~ 1, family = gaussian(link=identity))

#Running deviance test for model fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))

## 'log Lik.' 7.256887 (df=2)

#Running p-value for model fit
print(p.value<- pchisq( deviance, df=4, lower.tail = FALSE))

## 'log Lik.' 0.1229198 (df=2)</pre>
```

(e) Predict the score for a nurse who has worked at the center for seven years and who had previously been a co-PI on a grant that involved quality assurance

```
We calculate score 2 = 100 \cdot (1 - (0.6093 + 0.2122 + 0.0002 \cdot 7 + 0.0773)) 2 = 75.84951.
```

```
#Predicting values using the fitted model
pscore.tr<- predict(fitted.model, data.frame(desgn.rel="nurse", wrkyrs=7,</pre>
```

```
priorQI.rel="yes"))
print(100/(1-pscore.tr))
##
         1
## 75.86528
#Gamma 2.6
#This problem uses the same data as 2.1 (a) Fit the gamma regression model wi
th the log link function. Write down the fitted model. Check its goodness of
fit.
#Importing data for 2.6 (this uses the same data as 2.1)
ObesityGamma<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 41
0/Exercise Files/Exercise2.1Data.csv",header=TRUE, sep=",")
#Finding the difference in BMI
BMIdiff<- ObesityGamma$preBMI-Obesity$postBMI
#Creating the gamma model
summary(fitted.model<- glm(BMIdiff ~ gender + age + group, data=ObesityGamma,</pre>
family=Gamma(link=log)))
##
## Call:
## glm(formula = BMIdiff ~ gender + age + group, family = Gamma(link = log),
##
      data = ObesityGamma)
##
## Deviance Residuals:
      Min
                1Q
                   Median
                                 3Q
                                         Max
## -1.2258 -0.5318 -0.2426
                             0.3200
                                      1.5279
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
0.38624
                         0.22749 1.698 0.099250 .
## genderM
               0.04703 0.03826 1.229 0.227963
## age
              0.98701 0.23376 4.222 0.000187 ***
## groupTx
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for Gamma family taken to be 0.4599244)
##
##
      Null deviance: 24.361 on 35 degrees of freedom
## Residual deviance: 13.213 on 32 degrees of freedom
## AIC: 148.76
##
## Number of Fisher Scoring iterations: 7
#Checking the fit of the model
null.model<- glm(BMIdiff ~ 1, data=ObesityGamma, family=Gamma(link=log))</pre>
```

```
#Printing deviance test for fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))
## 'log Lik.' 23.83593 (df=2)
#Printing p-value for fit
print(p.value<- pchisq(deviance,3,lower.tail = FALSE))
## 'log Lik.' 2.70297e-05 (df=2)</pre>
```

- (b) What variables are significant predictors in this model? Use the 5% significance level. Only group is a significant predictor.
- (c) Interpret estimated significant regression coefficients. Estimated average decrease in BMI percentile for patients in the intervention group is $\exp(0.9870) \cdot 100\% = 268.32\%$ of that for patients in the control group.
- (d) Predict change in BMI percentile for a 9-year old girl in the control group. Compare the prediction with the one obtained in Exercise 2.1.

We calculate the predicted value as follows: $BMIdiff \square = \exp(-0.0442 + 0.0470 \cdot 9) = 1.4605$.

```
#Using the gamma model to predict values
print(predict(fitted.model, data.frame(gender="F", age=9, group="Cx"),
type="response"))
##
          1
## 1.461058
#Gamma 2.7
#This problem uses the same data as 2.2 (a) Fit the gamma regression model wi
th the log link function. Write down the fitted model. Check its goodness of
fit.
#Importing data for 2.7 (this uses the same data as 2.2)
QIGamma<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exe
rcise Files/Exercise2.2Data.csv",header=TRUE, sep=",")
#Rescaling and relevel
desgn.rel<- relevel(as.factor(OIGamma$desgn), ref="staff")</pre>
priorQI.rel<- relevel(as.factor(QIGamma$priorQI), ref="no")</pre>
score<- QIGamma$score/100</pre>
#Creating the gamma model
summary(fitted.model<- glm(score ~ desgn.rel + wrkyrs + priorQI.rel,</pre>
data=QIGamma, family=Gamma(link=log)))
```

```
##
## Call:
## glm(formula = score ~ desgn.rel + wrkyrs + priorQI.rel, family = Gamma(lin
k = log),
       data = QIGamma)
##
##
## Deviance Residuals:
        Min
                   10
                         Median
                                       30
                                                Max
## -0.25489 -0.12255 -0.01360
                                  0.08738
                                            0.29224
##
## Coefficients:
                     Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
                    4.1427297 0.0686964 60.305
                                                   <2e-16 ***
## desgn.reldoctor 0.1339899 0.0667675
                                           2.007
                                                   0.0516 .
## desgn.relnurse 0.1540443 0.0636050
                                           2.422
                                                   0.0201 *
## wrkyrs
                  -0.0002455 0.0029813 -0.082
                                                   0.9348
## priorQI.relyes 0.0532444 0.0498513
                                           1.068
                                                   0.2919
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for Gamma family taken to be 0.02298337)
##
       Null deviance: 1.06477 on 44
                                      degrees of freedom
##
## Residual deviance: 0.90565
                              on 40 degrees of freedom
## AIC: 348.1
##
## Number of Fisher Scoring iterations: 4
#Checking the fit of the model
null.model<- glm(score ~ 1, data=QIGamma, family=Gamma(link=log))</pre>
#Printing deviance test for fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 7.310283 (df=2)
#Printing p-value for fit
print(p.value<- pchisq(deviance,4,lower.tail=FALSE))</pre>
## 'log Lik.' 0.1203719 (df=2)
```

(b) Discuss significance of the beta coefficients. Interpret the estimated significant coefficients.

Indicators of doctor and nurse are significant at the 5% level. The estimated mean score for doctors is $\exp(0.1340) \cdot 100\% = 114.34\%$ of that for staff. The estimated mean score for nurses is $\exp(0.1540) \cdot 100\% = 116.65\%$ of that for staff.

(c) Predict the score for a nurse who has worked at the center for seven years and who had previously been a co-PI on a grant that involved quality assurance component. Compare that predicted score to the one obtained in Exercise 2.2.

The predicted value is calculated as: score = $100 \cdot exp(-0.4624 + 0.1540 - 0.0002 \cdot 7 + 0.0532) = 77.3678$. The predicted score in Exercise 2.2 is 75.8653 which is smaller than what we predict here.

```
#Using the gamma model to predict values
print(predict(fitted.model, data.frame(desgn.rel="nurse", wrkyrs=7,
priorQI.rel="yes"), type="response"))
##
## 77.34687
#Logistic 3.2
#(a) Fit a binary logistic model. Write down the fitted model. Discuss signif
icance of predictor variables, and goodness of fit of the model. Use \alpha = 0.05
#Importing data for 3.2
Psoriasis<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/E
xercise Files/Exercise3.2Data.csv",header=TRUE, sep=",")
#Releveling
gender.rel<- relevel(as.factor(Psoriasis$gender), ref="F")</pre>
medication.rel<- relevel(as.factor(Psoriasis$medication), ref="B")</pre>
#Creating the logistic model
summary(fitted.model<- glm(relief ~ gender.rel + age + medication.rel,</pre>
data=Psoriasis, family=binomial(link=logit)))
##
## Call:
## glm(formula = relief ~ gender.rel + age + medication.rel, family = binomia
1(link = logit),
       data = Psoriasis)
##
##
## Deviance Residuals:
                 10
                                   30
                                           Max
##
       Min
                      Median
## -1.9823 -0.3628
                      0.1974
                               0.5294
                                        1.6865
##
## Coefficients:
                   Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                   -6.79921
                               3.11430 -2.183
                                                 0.0290 *
                                         2.156
## gender.relM
                    3.17132
                               1.47097
                                                 0.0311 *
                                         1.971
## age
                    0.17131
                               0.08691
                                                 0.0487 *
## medication.relA 3.81641
                               1.54617
                                         2.468
                                                 0.0136 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
```

```
Null deviance: 36.652 on 29 degrees of freedom
## Residual deviance: 21.008 on 26 degrees of freedom
## AIC: 29.008
##
## Number of Fisher Scoring iterations: 6
#Calculating and printing AICC
p<-4
n<-30
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 30.60842 (df=4)
#Calculating and printing BIC
BIC(fitted.model)
## [1] 34.6132
#Checking to see if this model fits
null.model<- glm(relief ~ 1, data=Psoriasis, family=binomial(link=logit))</pre>
#Deviance test for fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 15.64344 (df=1)
#P-value for fit
print(p.value<- pchisq(deviance,3,lower.tail = FALSE))</pre>
## 'log Lik.' 0.001341726 (df=1)
```

(b) Give interpretation of the estimated significant regression coefficients.

The estimated odds in favor of relief from psoriasis for male patients are $\exp(3.1713) \cdot 100\% = 2,383.85\%$ of those for female patients. As age increases by one year, the estimated odds increase by $(\exp(0.1713) - 1) \cdot 100\% = 18.68\%$. The estimated odds for patients taking medication A are $\exp(3.8164) \cdot 100\% = 4,544.03\%$ of those for patients taking medication B.

(c) Find the predicted probability of relief from psoriasis for a 50-year old woman who is administered the medication A treatment.

#Answer should be P0(relief)=0.99625

```
#Calculating predicitons using the model
print(predict(fitted.model, data.frame(gender.rel="F", age=50,
medication.rel="A"), type="response"))
## 1
## 0.9962508
```

#Logistic 3.3

#(a) Fit a binary logistic model to the data. What predictors turn out to be significant at the 5% level? How good is the fit of the model?

#Importing data for 3.3

Novel<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exerc ise Files/Exercise3.3Data.csv", stringsAsFactors=TRUE, header=TRUE, sep=",")

Novel

##		success	cover	${\sf methods}$	novels	years
##	1	yes	yes	one	many	18
##	2	no	no	one	first	7
##	3	no	yes	none	several	10
##	4	yes	yes	many	many	6
##	5	no	yes	none	several	1
##	6	no	no	one	several	1
##	7	no	no	one	first	11
##	8	yes	no	one	several	19
##	9	yes	yes	none	first	5
##	10	no	no	none	many	2
##	11	no	no	one	several	10
##	12	no	no	many	many	9
##	13	yes	no	many	several	6
##	14	yes	yes	many	many	8
##	15	no	no	one	several	12
##	16	no	no	none	many	2
##	17	yes	no	none	several	17
##	18	yes	yes	many	first	10
##	19	yes	no	none	several	7
##	20	no	no	one	first	12
##	21	no	yes	none	several	7
##	22	no	yes	none	many	4
##	23	no	no	one	several	9
##	24	yes	no	many	several	13
##	25	yes	yes	none	first	6
##	26	no	no	none	many	2
##	27	yes	yes	one	several	7
##	28	yes	yes	many	many	17
##	29	yes	yes	many	first	18
##	30	yes	yes	one	several	17
##	31	no	yes	none	several	9
##	32	no	no	one	several	11
##	33	yes	yes	many	first	17
##	34	no	no	many	many	1
##	35	no	no	many	many	6
##	36	no	yes	none	several	1
##	37	yes	yes	many	first	6
##	38	yes	yes	one	many	4
##	39	no	yes	none	many	7
##	40	no	no	one	first	12

```
## 41
           no
                 no
                        one several
## 42
                                         9
          yes
                yes
                        one several
                                         8
## 43
                        one several
           no
                 no
## 44
                                         2
                        one several
           no
                 no
#Releveling
cover.rel<- relevel(as.factor(Novel$cover), ref="no")</pre>
methods.rel<- relevel(as.factor(Novel$methods), ref="none")</pre>
novels.rel<- relevel(as.factor(Novel$novels), ref="many")</pre>
Novel <- Novel %>% mutate_each(funs(as.factor), c(cover, methods, novels))
## Warning: `funs()` was deprecated in dplyr 0.8.0.
## Please use a list of either functions or lambdas:
##
     # Simple named list:
##
     list(mean = mean, median = median)
##
##
##
     # Auto named with `tibble::lst()`:
##
    tibble::lst(mean, median)
##
##
     # Using lambdas
     list(~ mean(., trim = .2), ~ median(., na.rm = TRUE))
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
generated.
## Warning: `mutate each ()` was deprecated in dplyr 0.7.0.
## Please use `across()` instead.
## This warning is displayed once every 8 hours.
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
generated.
#Creating the logistic model
summary(fitted.model<- glm(success ~ cover.rel + methods.rel + novels.rel</pre>
+ years, data=Novel, family=binomial(link=logit)))
##
## Call:
## glm(formula = success ~ cover.rel + methods.rel + novels.rel +
       years, family = binomial(link = logit), data = Novel)
##
##
## Deviance Residuals:
##
       Min
                 10
                      Median
                                    3Q
                                            Max
## -1.5998 -0.5951 -0.1052
                                0.3947
                                         2.6358
##
## Coefficients:
##
                     Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                      -6.8762
                                   2.3656 -2.907
                                                   0.00365 **
## cover.relyes
                       3.5238
                                   1.3120
                                            2.686
                                                   0.00724 **
## methods.relmany
                       3.9286
                                  1.7917
                                            2.193 0.02834 *
```

```
## methods.relone
                       0.7008
                                  1.1914
                                           0.588
                                                   0.55642
## novels.relfirst
                       1.8768
                                  1.5784
                                           1.189
                                                   0.23442
## novels.relseveral
                       1.3992
                                  1.3254
                                            1.056 0.29113
                       0.2907
                                  0.1269
                                           2.291 0.02198 *
## years
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 60.176 on 43 degrees of freedom
## Residual deviance: 32.071 on 37 degrees of freedom
## AIC: 46.071
##
## Number of Fisher Scoring iterations: 6
#Calculating and printing AICC
p<-7
n<-44
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 49.18253 (df=7)
#Calculating and printing BIC
BIC(fitted.model)
## [1] 58.56075
#Checking to see if this model fits
null.model<- glm(success ~ 1, data=Novel, family=binomial(link=logit))</pre>
#Deviance test for fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 28.10479 (df=1)
#P-value for fit
print(p.value<- pchisq(deviance,6,lower.tail = FALSE))</pre>
## 'log Lik.' 8.97874e-05 (df=1)
```

(b) Give interpretation of the estimated significant beta coefficients.

The estimated odds in favor of financial success for a novel with catchy cover are $\exp(3.5238) \cdot 100\% = 3,391.31\%$ of those for a novel without a catchy cover. The estimated odds for a publisher with many promotional methods are $\exp(3.9286) \cdot 100\% = 5,083.58\%$ of those for a publisher with no promotional methods. For every additional year a publisher was in business prior to publication of a novel, the estimated odds in favor of financial success of the novel increase by $(\exp(0.2907) - 1) \cdot 100\% = 33.74\%$.

(c) Suppose a newly established publishing house prints a novel by some previously unknown author, and doesn't advertise the publication. Find the estimated

probability that this novel is successful financially, if it has an extremely catchy cover.

Answer should be P0(success)=0.18609

```
#Calculating predictions using the model
print(predict(fitted.model, data.frame(cover.rel="yes", methods.rel="none",
novels.rel="first", years=0), type="response"))
## 0.186086
#Probit 3.4
#(a) Run the binary logistic model, regressing on all the predictors. Identif
y variables that are significant predictors of loan default at the 5% level o
f significance. Analyze the model fit.
#Importing data for 3.4
LoanProbit<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise3.4Data.csv", stringsAsFactors=TRUE, header=TRUE, sep=",
")
#Creating probit model
summary(fitted.model<- glm(default ~ LTV + age + income, data=LoanProbit,</pre>
family=binomial(link=probit)))
##
## Call:
## glm(formula = default ~ LTV + age + income, family = binomial(link = probi
t),
       data = LoanProbit)
##
##
## Deviance Residuals:
       Min
                 10
                     Median
                                   3Q
                                           Max
## -1.5284 -0.6871 -0.1290
                               0.5551
                                        1.9530
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.60587
                          2.34383 -0.685
                                             0.4933
## LTV
               0.06200
                           0.02853
                                     2.173
                                             0.0297 *
               -0.09865
                           0.04121 -2.394
                                             0.0167 *
## age
## incomelow 0.63924
                          0.59365 1.077
                                             0.2816
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 45.004 on 34 degrees of freedom
## Residual deviance: 28.173 on 31 degrees of freedom
## AIC: 36.173
```

```
##
## Number of Fisher Scoring iterations: 7
#Calculating and printing AICC
p<-4
n<-35
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 37.50665 (df=4)
#Calculating and printing BIC
BIC(fitted.model)
## [1] 42.39471
#Checking to see if this model fits
null.model<- glm(default ~ 1, data=LoanProbit, family=binomial(link=probit))</pre>
#Deviance test for fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 16.83055 (df=1)
#P-value for fit
print(p.value<- pchisq(deviance,3,lower.tail = FALSE))</pre>
## 'log Lik.' 0.0007657748 (df=1)
```

(b) Interpret the estimated significant beta coefficients. What is your suggestion in order for the bank to decrease the default rate of home equity loans?

As loan-to-value ratio increases by one, the estimated odds in favor or default increase by $(\exp(0.1059) - 1) \cdot 100\% = 11.17\%$. As the age of a client increases by one year, the estimated odds in favor of default change by $(\exp(-0.1616) - 1) \cdot 100\% = -14.92\%$, that is, decrease by 14.92%. To decrease the default rate, the bank might want to give loans with smaller loan-to-value ratio, and/or give loans to older clients.

(c) Give a point estimate for the probability of loan default if LTV ratio is 50%, and the borrower is a 50-year old men with high income.

Answer should be P0(success)=0.0030374

#Probit 3.5

#(a) Model the probability of being a cardiac patient via the binary logistic regression. Write the fitted model explicitly. Discuss the goodness of fit of

```
the model and significance of the regression coefficients. Assume \alpha = 0.01 fo
r all tests.
#Importing data for 3.5
CardiacProbit<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 4
10/Exercise Files/Exercise3.5Data.csv", stringsAsFactors=TRUE, header=TRUE, sep
=",")
#Creating probit model
summary(fitted.model<- glm(group ~ A + W, data=CardiacProbit,</pre>
family=binomial(link=probit)))
##
## Call:
## glm(formula = group ~ A + W, family = binomial(link = probit),
       data = CardiacProbit)
##
## Deviance Residuals:
##
       Min
                 10
                      Median
                                    3Q
                                            Max
## -2.2203 -0.9519 -0.1981
                               1.1270
                                         1.6154
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -0.62995
                           0.53497 -1.178
                                              0.2390
## A
                0.24895
                           0.11125
                                      2.238
                                              0.0252 *
## W
                0.01047
                           0.07694
                                      0.136
                                              0.8918
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 66.542 on 47 degrees of freedom
## Residual deviance: 58.302 on 45 degrees of freedom
## AIC: 64.302
##
## Number of Fisher Scoring iterations: 4
#Calculating and printing AICC
p<-3
n<-48
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 64.84727 (df=3)
#Calculating and printing BIC
BIC(fitted.model)
## [1] 69.91541
#Checking to see if this model fits
null.model<- glm(group ~ 1, data=CardiacProbit, family=binomial(link=probit))</pre>
```

```
#Deviance test for fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 8.240317 (df=1)
#P-value for fit
print(p.value<- pchisq(deviance,2,lower.tail = FALSE))</pre>
## 'log Lik.' 0.01624194 (df=1)
#using fitted model for prediction
print(predict(fitted.model, data.frame(A=2, W=4), type="response"))
## 0.4640763
#Comp Log-Log 3.4
#Importing data for 3.4
LoanComp<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Ex
ercise Files/Exercise3.4Data.csv",stringsAsFactors=TRUE,header=TRUE, sep=",")
#Creating comp log log
summary(fitted.model<- glm(default ~ LTV + age + income, data=LoanComp,</pre>
family=binomial(link=cloglog)))
##
## Call:
## glm(formula = default ~ LTV + age + income, family = binomial(link = clog1
og),
##
       data = LoanComp)
##
## Deviance Residuals:
       Min
                 10
                      Median
                                   3Q
                                           Max
## -1.4972 -0.6653 -0.2425
                               0.4326
                                        1.9708
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -2.68156
                           3.27613 -0.819
                                              0.4131
## LTV
                0.07896
                           0.04114
                                    1.919
                                              0.0550 .
                           0.05111 -2.398
                                              0.0165 *
## age
               -0.12254
## incomelow
               0.89073
                           0.72016
                                    1.237
                                              0.2161
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 45.004 on 34 degrees of freedom
## Residual deviance: 28.436 on 31 degrees of freedom
## AIC: 36.436
```

```
##
## Number of Fisher Scoring iterations: 7
#Calculating and printing AICC
p<-4
n<-35
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 37.7691 (df=4)
#Calculating and printing BIC
BIC(fitted.model)
## [1] 42.65716
#Checking if the model fits
null.model<- glm(default ~ 1, data=LoanComp, family=binomial(link=cloglog))</pre>
#Deviance test for model fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 16.56811 (df=1)
#P-value for model fit
print(p.value<- pchisq(deviance,3,lower.tail = FALSE))</pre>
## 'log Lik.' 0.0008670163 (df=1)
#Predicting values using the fitted model
print(predict(fitted.model, data.frame(LTV=50, age=50, income="high"),
type="response"))
##
## 0.007713725
##Comp Log-Log 3.5
#Importing data for 3.5
CardiacComp<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410
/Exercise Files/Exercise3.5Data.csv", stringsAsFactors=TRUE, header=TRUE, sep="
,")
#Creating comp log log
summary(fitted.model<- glm(group ~ A + W, data=CardiacComp,</pre>
family=binomial(link=cloglog)))
##
## Call:
## glm(formula = group ~ A + W, family = binomial(link = cloglog),
##
       data = CardiacComp)
##
## Deviance Residuals:
##
       Min
                 10 Median
                                    3Q
                                            Max
```

```
## -2.1991 -0.9736 -0.2271
                               1.1681
                                        1.5382
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -1.03922
                           0.62874 -1.653
                                             0.0984 .
                                             0.0291 *
## A
                0.23818
                           0.10916
                                     2.182
## W
                0.01674
                           0.09199
                                     0.182
                                             0.8556
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
       Null deviance: 66.542 on 47 degrees of freedom
## Residual deviance: 59.047 on 45 degrees of freedom
## AIC: 65.047
## Number of Fisher Scoring iterations: 8
#Calculating and printing AICC
p<-3
n<-48
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 65.59256 (df=3)
#Calculating and printing BIC
BIC(fitted.model)
## [1] 70.66071
#Checking if the model fits
null.model<- glm(group ~ 1, data=CardiacComp, family=binomial(link=cloglog))</pre>
#Deviance test for model fit
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 7.495025 (df=1)
#P-value for model fit
print(p.value<- pchisq(deviance,2,lower.tail = FALSE))</pre>
## 'log Lik.' 0.02357632 (df=1)
#Predicting values using the fitted model
print(predict(fitted.model, data.frame(A=2, W=4), type="response"))
##
## 0.45612
```

Cumulative Logistic: 4.1

```
#Cumulative Logistic: 4.1
#First we import the data and fit the cumlogit model
Admission <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise4.1Data.csv", header = TRUE, sep =",")
Admission
##
      GPA GMAT
                  status
      2.96 596
## 1
                  admit
## 2 3.14 473
                  admit
## 3 3.22 482
                  admit
## 4 3.29 527
                  admit
## 5 3.69 505
                  admit
## 6 2.46 693
                  admit
## 7 3.03 626
                  admit
## 8 3.19 663
                  admit
## 9 3.63 447
                  admit
## 10 3.59 588
                  admit
## 11 3.30 563
                  admit
## 12 3.78
            591
                  admit
## 13 3.44 692
                  admit
## 14 3.48
            528
                  admit
## 15 3.47
            552
                  admit
## 16 3.35 520
                  admit
## 17 2.89
            543
                  admit
## 18 2.28 523
                  admit
## 19 3.21
            530
                  admit
## 20 3.58 564
                  admit
## 21 3.33
            565
                  admit
## 22 2.80
           444
                 border
## 23 3.13
           416
                 border
## 24 2.89 431
                 border
## 25 3.01 471
                 border
## 26 2.91 446
                 border
## 27 2.75
            546
                 border
## 28 2.73 467
                 border
## 29 3.12 463
                 border
## 30 3.08 440 notadmit
## 31 3.01 453 notadmit
## 32 3.03 414 notadmit
## 33 3.04 446 notadmit
## 34 2.89 485 notadmit
## 35 2.79 490 notadmit
## 36 2.54 446 notadmit
## 37 2.43 425 notadmit
## 38 2.20 474 notadmit
            531 notadmit
## 39 3.36
## 40 2.57 542 notadmit
```

```
## 41 2.36 482 notadmit
## 42 3.66 420 notadmit
#Now we the predictor GMAT and fit the model
GMAT rescaled<- Admission$GMAT/100</pre>
summary(fitted.model<- clm(as.factor(status) ~ GPA + GMAT rescaled, data=Admi</pre>
ssion,
link="logit"))
## formula: as.factor(status) ~ GPA + GMAT rescaled
## data:
            Admission
##
## link threshold nobs logLik AIC niter max.grad cond.H
## logit flexible 42 -27.54 63.09 5(0) 2.17e-07 1.6e+04
##
## Coefficients:
##
                 Estimate Std. Error z value Pr(>|z|)
## GPA
                  -3.1194
                              1.1913 -2.618 0.008833 **
## GMAT_rescaled -2.7755
                              0.8406 -3.302 0.000961 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Threshold coefficients:
                   Estimate Std. Error z value
## admit|border
                    -23.390
                                 5.857 -3.994
## border | notadmit -21.853
                                 5.670 -3.854
#Computations AICC and BIC
p<- 4
n<- 42
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 64.16976 (df=4)
BIC(fitted.model)
## [1] 70.03935
#Here we check to see if the model fits using a deviance test
null.model<- clm(as.factor(status) ~ 1, data=Admission, link="logit")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 31.04588 (df=2)
print(p.value<- pchisq(deviance, df=2, lower.tail = FALSE))</pre>
## 'log Lik.' 1.813312e-07 (df=2)
#Now we will use the fitted model to predict the probabilities of each admiss
ion status for someone with 3.1 GPA and GMAT score of 550
#print(predict(fitted.model, data.frame(GPA=3.1, GMAT_rescaled=5.50), type="p
rob"))
```

Cumulative Logistic: 4.2

```
#First we import the data and fit the cumlogit model
TVService <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise4.2Data.csv", header = TRUE, sep =",")
TVService
##
      subscribed magazine resolved satisf
## 1
                5
                                            5
                        yes
                                   no
## 2
               49
                                            5
                        yes
                                   no
                                            3
## 3
               56
                         no
                                   no
                                            5
## 4
               13
                        yes
                                  yes
## 5
               27
                         no
                                  yes
                                            4
               41
                                            5
## 6
                        yes
                                  yes
                2
                                            5
## 7
                        yes
                                  yes
## 8
               64
                                            4
                        yes
                                  yes
## 9
               88
                                            4
                        yes
                                  yes
## 10
               43
                                            4
                        yes
                                  yes
## 11
               94
                        yes
                                   no
                                            4
## 12
                8
                         no
                                   no
                                            1
                9
                                            2
## 13
                        yes
                                   no
## 14
               68
                                            4
                        yes
                                   no
## 15
                5
                                            2
                         no
                                  yes
                                            3
## 16
              108
                         no
                                  yes
## 17
               21
                        yes
                                            4
                                  yes
               25
                                            3
## 18
                        yes
                                   no
## 19
                2
                         no
                                  yes
                                            4
## 20
               11
                                            2
                         no
                                   no
                                            5
## 21
               98
                        yes
                                  yes
                                            5
## 22
               11
                         no
                                  yes
## 23
                                            4
               46
                         no
                                   no
## 24
                7
                                            3
                         no
                                   no
                7
                                            5
## 25
                         no
                                  yes
                9
                                            5
## 26
                        yes
                                  yes
                                            2
## 27
               17
                         no
                                   no
## 28
                8
                         no
                                  yes
                                            2
## 29
                9
                                            1
                         no
                                  yes
               95
## 30
                                            4
                         no
                                   no
                                            3
## 31
               60
                         no
                                  yes
                                            4
## 32
               80
                         no
                                  yes
## 33
                2
                                            3
                        yes
                                   no
## 34
               33
                                            4
                        yes
                                  yes
                5
                                            3
## 35
                        yes
                                   no
## 36
                7
                                            1
                         no
                                   no
#Making the response satisf a categorical variable
satisf_categorical<- as.factor(TVService$satisf)</pre>
#Specifying reference categories and making them categorical variables
magazine releveled<- relevel(as.factor(TVService$magazine), ref="yes")</pre>
resolved_releveled<- relevel(as.factor(TVService$resolved), ref="yes")</pre>
```

```
#Fitting the model
summary(fitted.model<- clm(satisf categorical ~ subscribed + magazine relevel</pre>
+ resolved_releveled, data=TVService, link="logit"))
## formula:
## satisf categorical ~ subscribed + magazine releveled + resolved releveled
## data:
            TVService
##
## link threshold nobs logLik AIC
                                       niter max.grad cond.H
## logit flexible 36 -47.05 108.10 5(0) 8.00e-10 6.2e+04
##
## Coefficients:
##
                         Estimate Std. Error z value Pr(>|z|)
## subscribed
                                    0.009686
                                               1.086 0.27760
                         0.010516
## magazine_releveledno -1.917509
                                    0.677141 -2.832 0.00463 **
## resolved_releveledno -1.428832   0.655864   -2.179   0.02937 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Threshold coefficients:
       Estimate Std. Error z value
##
## 1 2 -4.2359
                    0.9678 -4.377
## 2|3
       -2.8298
                    0.8126 -3.483
## 3 4 -1.5740
                    0.7442 -2.115
## 4 | 5
        0.3350
                    0.6708
                             0.499
#Computations for AICC and BIC
p<-7
n<-36
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 112.0974 (df=7)
BIC(fitted.model)
## [1] 119.182
#Checking to see how well the model fits using a deviance test
null.model<- clm(satisf_categorical ~ 1, data=TVService, link="logit")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 14.7994 (df=4)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.001996357 (df=4)
#Predicting P probabilities of each of the five levels of the satisfaction sc
ore for a caller who had been subscribed for 3 months, doesn't receive the ma
gazine, and whose issue was resolved over the phone
```

Cumulative Probit: 4.1

```
#We start by importing the data
Admission <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise4.1Data.csv", header = TRUE, sep =",")
#Here we rescale the predictor GMAT and fit the cumprobit model
GMAT_rescaled<- Admission$GMAT/100</pre>
summary(fitted.model<- clm(as.factor(status) ~ GPA + GMAT_rescaled, data=Admi</pre>
ssion,
link="probit"))
## formula: as.factor(status) ~ GPA + GMAT_rescaled
## data:
           Admission
##
          threshold nobs logLik AIC
                                      niter max.grad cond.H
   probit flexible 42 -27.59 63.19 6(0) 1.04e-13 1.3e+04
##
##
## Coefficients:
##
                 Estimate Std. Error z value Pr(>|z|)
## GPA
                  -1.7356
                             0.6222 -2.790 0.005276 **
                             0.4573 -3.618 0.000297 ***
## GMAT_rescaled -1.6546
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Threshold coefficients:
                   Estimate Std. Error z value
##
## admit|border
                    -13.603
                                 3.078
                                         -4.42
## border notadmit -12.724
                                         -4.24
                                 3.001
Admission
##
      GPA GMAT
                  status
## 1 2.96 596
                  admit
## 2 3.14 473
                  admit
## 3 3.22 482
                  admit
## 4 3.29 527
                  admit
## 5 3.69 505
                  admit
## 6 2.46 693
                  admit
## 7 3.03 626
                  admit
## 8 3.19 663
                  admit
## 9 3.63 447
                  admit
## 10 3.59 588
                  admit
```

```
## 11 3.30
            563
                   admit
## 12 3.78
            591
                   admit
## 13 3.44 692
                   admit
## 14 3.48
            528
                   admit
## 15 3.47
            552
                   admit
## 16 3.35
            520
                   admit
## 17 2.89
            543
                   admit
## 18 2.28
            523
                   admit
## 19 3.21
            530
                   admit
## 20 3.58
            564
                   admit
## 21 3.33
            565
                   admit
## 22 2.80 444
                  border
## 23 3.13
            416
                  border
                  border
## 24 2.89
            431
## 25 3.01
            471
                  border
## 26 2.91
            446
                  border
## 27 2.75
            546
                  border
## 28 2.73
            467
                  border
## 29 3.12
            463
                  border
## 30 3.08 440 notadmit
## 31 3.01 453 notadmit
## 32 3.03
           414 notadmit
## 33 3.04 446 notadmit
## 34 2.89
            485 notadmit
## 35 2.79
            490 notadmit
## 36 2.54 446 notadmit
## 37 2.43
            425 notadmit
## 38 2.20 474 notadmit
## 39 3.36 531 notadmit
## 40 2.57
            542 notadmit
## 41 2.36 482 notadmit
## 42 3.66 420 notadmit
#Computations for AICC and BIC
p<- 4
n<- 42
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 64.26707 (df=4)
BIC(fitted.model)
## [1] 70.13666
#Now we check to see if the model fits using a deviance test
null.model<- clm(as.factor(status) ~ 1, data=Admission, link="probit")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 30.94857 (df=2)
print(p.value<- pchisq(deviance, df=2, lower.tail = FALSE))</pre>
```

```
## 'log Lik.' 1.903718e-07 (df=2)

#Now we will use the fitted model to predict the probabilities of each admiss
ion status for someone with 3.1 GPA and GMAT score of 550

#print(predict(fitted.model, data.frame(GPA=3.1, GMAT_rescaled=5.50), type="p
rob"))
```

Cumulative Probit 4.2

```
#First we import the data and fit the cumprobit model
TVService <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise4.2Data.csv", header = TRUE, sep =",")
#Making the response satisf a categorical variable
satisf_categorical<- as.factor(TVService$satisf)</pre>
#Specifying reference categories and making them categorical variables
magazine releveled<- relevel(as.factor(TVService$magazine), ref="yes")</pre>
resolved releveled<- relevel(as.factor(TVService$resolved), ref="yes")</pre>
#Fitting the model
summary(fitted.model<- clm(satisf categorical ~ subscribed + magazine relevel</pre>
ed + resolved_releveled,
data=TVService, link="probit"))
## formula:
## satisf categorical ~ subscribed + magazine releveled + resolved releveled
## data:
         TVService
##
## link
         threshold nobs logLik AIC
                                    niter max.grad cond.H
## probit flexible 36 -46.69 107.39 4(0) 1.70e-09 5.5e+04
##
## Coefficients:
##
                       Estimate Std. Error z value Pr(>|z|)
                      0.006061 0.005718 1.060 0.28919
## subscribed
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Threshold coefficients:
      Estimate Std. Error z value
##
## 1 2 -2.6005
                  0.5384 -4.830
## 2 3 -1.7878
                  0.4709 -3.796
                  0.4411 -2.334
## 3 4 -1.0295
## 4 | 5
       0.1114
                  0.3933
                          0.283
#Computations for AICC and BIC
p<-7
```

```
n<-36
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 111.3854 (df=7)
BIC(fitted.model)
## [1] 118.47
#Checking to see how well the model fits using a deviance test
null.model<- clm(satisf_categorical ~ 1, data=TVService, link="probit")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 15.51138 (df=4)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.001427906 (df=4)
#Predicting P probabilities of each of the five levels of the satisfaction sc
ore for a caller who had been subscribed for 3 months, doesn't receive the ma
gazine, and whose issue was resolved over the phone
print(predict(fitted.model, data.frame(subscribed=3, magazine_releveled="no",
resolved_releveled="yes"), type="prob"))
## $fit
##
## 1 0.0783845 0.1947673 0.2884392 0.3408959 0.09751309
```

Cumulative Complementary Log Log: 4.1

```
#We start by importing the data
Admission <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise4.1Data.csv", header = TRUE, sep =",")
#Here we rescale the predictor GMAT and fit the cumulative cloglog model
GMAT rescaled<- Admission$GMAT/100
summary(fitted.model<- clm(as.factor(status) ~ GPA + GMAT rescaled, data=Admi</pre>
ssion,
link="cloglog"))
## formula: as.factor(status) ~ GPA + GMAT rescaled
## data:
           Admission
##
## link
           threshold nobs logLik AIC niter max.grad cond.H
## cloglog flexible 42 -29.50 66.99 5(0) 5.38e-07 1.3e+04
##
## Coefficients:
##
                 Estimate Std. Error z value Pr(>|z|)
                  -1.6479
## GPA
                            0.6316 -2.609 0.009081 **
                              0.4303 -3.530 0.000415 ***
## GMAT_rescaled -1.5190
## ---
```

```
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Threshold coefficients:
                   Estimate Std. Error z value
## admit|border
                    -13.137
                                  3.059 -4.294
## border notadmit -12.265
                                  2.989 -4.104
#Computations for AICC and BIC
p<- 4
n<- 42
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 68.07119 (df=4)
BIC(fitted.model)
## [1] 73.94079
#Now we check to see if the model fits using a deviance test
null.model<- clm(as.factor(status) ~ 1, data=Admission, link="cloglog")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 27.14445 (df=2)
print(p.value<- pchisq(deviance, df=2, lower.tail = FALSE))</pre>
## 'log Lik.' 1.275435e-06 (df=2)
#Now we will use the fitted model to predict the probabilities of each admiss
ion status for someone with 3.1 GPA and GMAT score of 550
#print(predict(fitted.model, data.frame(GPA=3.1, GMAT_rescaled=5.50), type="p
rob"))
```

Cumulative Complementary Log Log: 4.2

```
#First we import the data and fit the cum CLogLog model
TVService <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise4.2Data.csv", header = TRUE, sep =",")
#Making the response satisf a categorical variable and fitting the model
satisf categorical<- as.factor(TVService$satisf)</pre>
summary(fitted.model<- clm(satisf_categorical ~ subscribed + magazine + resol</pre>
ved,
data=TVService, link="cloglog"))
## formula: satisf_categorical ~ subscribed + magazine + resolved
## data:
           TVService
##
           threshold nobs logLik AIC
                                         niter max.grad cond.H
  cloglog flexible 36 -48.37 110.73 6(0) 4.42e-12 5.2e+04
##
## Coefficients:
```

```
Estimate Std. Error z value Pr(>|z|)
## subscribed 0.002997
                          0.006339
                                     0.473 0.63641
## magazineyes 1.142173
                          0.422566
                                     2.703
                                            0.00687 **
## resolvedyes 1.009893
                          0.430080
                                     2.348 0.01887 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Threshold coefficients:
       Estimate Std. Error z value
## 1 2 -1.4523
                    0.6499 -2.235
## 2|3 -0.3188
                    0.4857 -0.656
## 3 4
       0.5528
                    0.4616
                             1.198
## 4 5
        1.7226
                    0.4941
                             3,487
#Computations for AICC and BIC
p<-7
n<-36
print(AICC<- -2*logLik(fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 114.7302 (df=7)
BIC(fitted.model)
## [1] 121.8149
#Checking to see how well the model fits using a deviance test
null.model<- clm(satisf_categorical ~ 1, data=TVService, link="cloglog")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 12.16653 (df=4)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.006833923 (df=4)
#Predicting P probabilities of each of the five levels of the satisfaction sc
ore for a caller who had been subscribed for 3 months, doesn't receive the ma
gazine, and whose issue was resolved over the phone
print(predict(fitted.model, data.frame(subscribed=3, magazine="no",
resolved="yes"), type="prob"))
## $fit
                                 3
## 1 0.0810142 0.1498225 0.2352053 0.4014798 0.1324782
```

Generalized Logit Model (for nominal response) 4.4

```
#First we import the data and fitting the generalized probit model
Forecast <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/E
xercise Files/Exercise4.4Data.csv", header = TRUE, sep =",")
#Specifying reference categories and changing them into categorical variables</pre>
```

```
outcome releveled <- relevel(as.factor(Forecast$outcome), ref="C")</pre>
water_releveled <- relevel(as.factor(Forecast$water), ref="yes")</pre>
#Running the model and checking to see how well it fits using a deviance test
summary(fitted.model<- multinom(outcome_releveled ~ elevation + winddir + wi</pre>
ndspeed +
water_releveled , data=Forecast))
## # weights: 18 (10 variable)
## initial value 32.958369
## iter 10 value 17.771391
## iter 20 value 17.356931
## final value 17.356923
## converged
## Call:
## multinom(formula = outcome_releveled ~ elevation + winddir +
       windspeed + water releveled, data = Forecast)
##
## Coefficients:
      (Intercept)
                    elevation
                                  winddir windspeed water releveledno
##
## FA
        -7.764088 0.001089404 0.025033087 0.1116808
                                                            0.22092498
        -5.218338 0.002986672 0.007571138 0.1421470
## FD
                                                            0.04887474
##
## Std. Errors:
      (Intercept)
                    elevation
                                  winddir windspeed water_releveledno
##
         3.582384 0.001953411 0.012024957 0.1756380
## FA
         3.049625 0.001807177 0.008640188 0.1962268
## FD
                                                              1.585012
##
## Residual Deviance: 34.71385
## AIC: 54.71385
#Deviance test
null.model<- multinom(outcome_releveled ~ 1, data=Forecast)</pre>
## # weights: 6 (2 variable)
## initial value 32.958369
## final value 30.288413
## converged
print(deviance<- deviance(null.model)-deviance(fitted.model))</pre>
## [1] 25.86298
print(p.value<- pchisq(deviance, df=8, lower.tail = FALSE))</pre>
## [1] 0.001108523
#Using the fitted model to predict the probabilities for an airport at 2000 f
eet, away from water, winf at 5 knots eastward.
print(predict(fitted.model, data.frame(elevation=2000, winddir=90, windspeed=
```

```
5,
water releveled ="no"), type="prob"))
            C
                      FΑ
                                  FD
## 0.09933447 0.00773318 0.89293235
```

Generalized Logit Model (for nominal response) 4.5

```
#First we import the data and fitting the generalized probit model
AnkleCondition <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT
410/Exercise Files/Exercise4.5Data.csv", header = TRUE, sep =",")
#Specifying reference categories and changing them into categorical variables
condition releveled<- relevel(as.factor(AnkleCondition$condition), ref="sprai
ned")
gender_releveled<- relevel(as.factor(AnkleCondition$gender), ref="female")</pre>
#Running the model and checking to see how well it fits using a deviance test
summary(fitted.model<- multinom(condition releveled ~ age + gender releveled,</pre>
data=AnkleCondition))
## # weights: 12 (6 variable)
## initial value 27.465307
## iter 10 value 21.077309
## final value 21.066688
## converged
## Call:
## multinom(formula = condition releveled ~ age + gender releveled,
       data = AnkleCondition)
##
##
## Coefficients:
            (Intercept)
                                age gender_releveledmale
##
## broken
            -4.6658881 0.21077974
                                                  4.195560
              0.3352406 -0.04544743
## torn
                                                   1.556776
##
## Std. Errors:
            (Intercept)
                              age gender_releveledmale
## broken
               3.162765 0.2023320
                                                1.655647
               1.679667 0.1379255
## torn
                                                1.273358
##
## Residual Deviance: 42.13338
## AIC: 54.13338
#Deviance test
summary(null.model<- multinom(condition_releveled ~ 1, data=AnkleCondition))</pre>
## # weights: 6 (2 variable)
## initial value 27.465307
```

```
## final value 27.300482
## converged
## Call:
## multinom(formula = condition releveled ~ 1, data = AnkleCondition)
## Coefficients:
            (Intercept)
##
              0.2513144
## broken
              0.2513144
## torn
##
## Std. Errors:
            (Intercept)
## broken
              0.5039526
## torn
              0.5039526
## Residual Deviance: 54.60096
## AIC: 58.60096
print(deviance<- deviance(null.model)-deviance(fitted.model))</pre>
## [1] 12.46759
print(p.value<- pchisq(deviance, df=4, lower.tail = FALSE))</pre>
## [1] 0.01419266
#We will now use the fitted model to predict the probability of each type of
injury for a 9 year old girl.
print(predict(fitted.model, data.frame(age=9, gender releveled="female"), typ
e="prob"))
     sprained
                broken
##
                           torn
## 0.50210809 0.03149866 0.46639325
```

Poisson 5.1

```
#First we begin by importing the data
DefectiveData <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT
410/Exercise Files/Exercise5.1Data.csv", header = TRUE, sep =",")

#Releveling shift and setting day as the reference
shift_releveled<- relevel(as.factor(DefectiveData$shift), ref="day")

#Fitting the model
summary(fitted.model<- glm(ndefectives ~ experience + shift_releveled,
data=DefectiveData, family=poisson(link=log)))

##
## Call:
## glm(formula = ndefectives ~ experience + shift_releveled, family = poisson</pre>
```

```
(link = log),
       data = DefectiveData)
##
##
## Deviance Residuals:
##
       Min
                   10
                        Median
                                       3Q
                                                Max
## -2.05694 -0.56800
                        0.05144
                                  0.47579
                                            2.04325
## Coefficients:
                          Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                           0.35714
                                      0.33734
                                                1.059 0.28973
## experience
                           0.03552
                                      0.04709
                                                0.754 0.45066
                                                1.276 0.20195
## shift releveledevening 0.40813
                                      0.31985
## shift releveledmorning 0.10090
                                      0.33419
                                                0.302 0.76270
## shift relevelednight
                           0.90671
                                      0.30630 2.960 0.00307 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for poisson family taken to be 1)
##
##
       Null deviance: 39.019 on 35 degrees of freedom
## Residual deviance: 27.731 on 31 degrees of freedom
## AIC: 128.62
##
## Number of Fisher Scoring iterations: 5
#Checking to see how well the model fits with a deviance test
null.model<- glm(ndefectives ~ 1, data=DefectiveData, family=poisson(link=log</pre>
))
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 11.28837 (df=1)
print(p.value<- pchisq(deviance, df=4, lower.tail = FALSE))</pre>
## 'log Lik.' 0.02350729 (df=1)
#Using the model to predict the number of defective items produced during a n
ight shift by an operator with six months of experience
print(predict(fitted.model, data.frame(experience=6, shift_releveled="night")
type="response"))
## 4.379627
```

Poisson 5.2

```
#First we begin by importing the data
CarInsuranceData <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/ST
AT 410/Exercise Files/Exercise5.2Data.csv", header = TRUE, sep =",")</pre>
```

```
#Releveling gender and setting day as the reference
gender releveled<- relevel(as.factor(CarInsuranceData$gender), ref="F")</pre>
#Fitting the model
summary(fitted.model<- glm(naccidents ~ gender_releveled + age + miles,</pre>
data=CarInsuranceData, family=poisson(link=log)))
##
## Call:
## glm(formula = naccidents ~ gender_releveled + age + miles, family = poisso
n(link = log),
       data = CarInsuranceData)
##
##
## Deviance Residuals:
##
        Min
                   10
                         Median
                                        3Q
                                                 Max
## -2.63470 -0.83300
                        0.00054
                                  0.67182
                                             1.70119
##
## Coefficients:
                      Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                      0.449155
                                 0.370804
                                             1.211
                                                      0.226
## gender_releveledM 0.218899
                                 0.160938
                                             1.360
                                                      0.174
                                             2.543
                                                      0.011 *
## age
                      0.017103
                                 0.006726
## miles
                     -0.001283
                                 0.001781 -0.721
                                                      0.471
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '* 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for poisson family taken to be 1)
##
##
       Null deviance: 70.348 on 47 degrees of freedom
## Residual deviance: 61.761 on 44 degrees of freedom
## AIC: 202.64
##
## Number of Fisher Scoring iterations: 5
#Checking to see how well the model fits with a deviance test
null.model<- glm(naccidents ~ 1, data=CarInsuranceData, family=poisson(link=1
og))
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 8.587217 (df=1)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.03531361 (df=1)
#Using the model to predict the value of the total number of auto accidents c
aused by a 35-year-old woman who has driven a total of one hundred thousand m
print(predict(fitted.model, data.frame(gender_releveled="F", age=35, miles=10
0),
type="response"))
```

```
## 1
## 2.507913
```

Zero-Truncated Poisson 5.4

```
#First we begin by importing the data which is taken from 5.1
DefectiveData <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT</pre>
410/Exercise Files/Exercise5.1Data.csv", header = TRUE, sep =",")
#Getting rid of zeros in data set and fitting the model
DefectiveData<- DefectiveData[which(DefectiveData$ndefectives != 0),]</pre>
#Fitting the model
summary(fitted.model<- vglm(ndefectives ~ experience + shift,</pre>
data=DefectiveData, family = pospoisson()))
##
## Call:
## vglm(formula = ndefectives ~ experience + shift, family = pospoisson(),
       data = DefectiveData)
##
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                 0.19203
                            0.44108
                                    0.435 0.66330
                 0.03355
                            0.05606
                                      0.598 0.54951
## experience
## shiftevening 0.47641
                                      1.173 0.24080
                            0.40615
## shiftmorning 0.36479
                            0.42558
                                    0.857 0.39135
## shiftnight
                 1.06314
                            0.38115 2.789 0.00528 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Name of linear predictor: loglink(lambda)
## Log-likelihood: -49.6747 on 28 degrees of freedom
##
## Number of Fisher scoring iterations: 4
## No Hauck-Donner effect found in any of the estimates
#Using a deviance test to check the fit of the model
null.model<- vglm(ndefectives ~ 1, data=DefectiveData, family = pospoisson())</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## [1] 9.904659
print((p.value<- pchisq(deviance, df=4, lower.tail = FALSE)))</pre>
## [1] 0.04206469
#Using the model to predict the number of defective items produced during a n
ight shift by an operator with six months of experience.
```

```
print(predict(fitted.model, data.frame(experience=6, shift="night"),
type="response"))
##
            [,1]
## [1,] 4.350423
```

Zero-Truncated Poisson 5.5

```
#First we begin by importing the data which is taken from 5.2
CarInsuranceData <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/ST</pre>
AT 410/Exercise Files/Exercise5.2Data.csv", header = TRUE, sep =",")
#Getting rid of zeros in data set and fitting the model
CarInsuranceData<- CarInsuranceData[which(CarInsuranceData$naccidents !=0),]</pre>
#Fitting the model
summary(fitted.model<- vglm(naccidents ~ gender + age + miles,</pre>
data=CarInsuranceData, family = pospoisson()))
##
## Call:
## vglm(formula = naccidents ~ gender + age + miles, family = pospoisson(),
       data = CarInsuranceData)
##
##
## Coefficients:
                 Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 0.4869342 0.3981630
                                       1.223
                                                0.2213
                                       0.485
## genderM
                0.0826963 0.1705286
                                                0.6277
                0.0157404 0.0072558 2.169
                                                0.0301 *
## age
               -0.0002013 0.0019055 -0.106
## miles
                                                0.9159
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Name of linear predictor: loglink(lambda)
##
## Log-likelihood: -84.0799 on 40 degrees of freedom
##
## Number of Fisher scoring iterations: 4
## No Hauck-Donner effect found in any of the estimates
#Using a deviance test to check the fit of the model
null.model<- vglm(naccidents ~ 1, data=CarInsuranceData, family = pospoisson(</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## [1] 5.087076
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## [1] 0.1655309
```

```
#Using the model to predict the total number of auto accidents caused by a 35
-year-old woman who has driven a total of one hundred thousand miles
print(predict(fitted.model, data.frame(gender="F", age=35, miles=100),
type="response"))
## [,1]
## [1,] 2.952453
```

Zero-Inflated Poisson 5.8

```
#First we begin by importing the data
Races<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exerc</pre>
ise Files/Exercise5.8Data.csv", header = TRUE, sep =",")
#Setting reference levels and turning run and gender into categorical variabl
run releveled<- relevel(as.factor(Races$run), ref="5K")</pre>
gender relevled<- relevel(as.factor(Races$gender), ref="M")</pre>
#Fitting the model
summary(fitted.model<- zeroinfl(nraces ~ gender_relevled + age + run_relevele</pre>
d | pace,
data=Races))
##
## Call:
## zeroinfl(formula = nraces ~ gender_relevled + age + run_releveled | pace,
       data = Races)
##
##
## Pearson residuals:
                10 Median
       Min
                                3Q
                                        Max
## -1.4548 -0.7087 -0.2621 0.4742 2.1409
##
## Count model coefficients (poisson with log link):
##
                      Estimate Std. Error z value Pr(>|z|)
                                             2.897 0.003769 **
## (Intercept)
                       1.63134
                                  0.56313
                                  0.30360
                                             3.369 0.000753 ***
## gender_relevledF
                       1.02296
## age
                      -0.04431
                                  0.01478 -2.998 0.002719 **
## run releveled10K
                                  0.36080
                                             0.514 0.607359
                       0.18539
## run releveledFull
                       0.75468
                                  0.32369 2.331 0.019727 *
##
## Zero-inflation model coefficients (binomial with logit link):
               Estimate Std. Error z value Pr(>|z|)
                            3.9941
## (Intercept) -8.7064
                                      -2.18
                                              0.0293 *
                 0.7255
                            0.3628
                                      2.00
                                              0.0455 *
## pace
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Number of iterations in BFGS optimization: 15
## Log-likelihood: -51.02 on 7 Df
```

```
#Using a deviance test to check the fit of the model
null.model<- zeroinfl(nraces ~ 1, data=Races)
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))
## 'log Lik.' 19.54518 (df=2)
print(p.value<- pchisq(deviance, df=5, lower.tail = FALSE))
## 'log Lik.' 0.001520639 (df=2)
#Using the fitted model to predict the number of races in the past four month
s for a female runner, aged 45, who ran at an average pace of 10 minutes per
mile, if she ran 10K
print(predict(fitted.model, data.frame(gender_relevled="F", run_releveled="10
K", age=45,
pace=10)))
## 1
## 1.88771</pre>
```

Zero-Inflated Poisson 5.9

```
#First we begin by importing the data
Reading <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Ex
ercise Files/Exercise5.9Data.csv", header = TRUE, sep =",")
#Setting reference levels and turning hw and gender into categorical variable
homework releveled<- relevel(as.factor(Reading$hw), ref="no")</pre>
gender_relevled<- relevel(as.factor(Reading$gender), ref="M")</pre>
#Fitting the model
summary(fitted.model<- zeroinfl(nbooks ~ homework_releveled + gender_relevled</pre>
grade,
data=Reading))
##
## Call:
## zeroinfl(formula = nbooks ~ homework releveled + gender relevled | grade,
##
       data = Reading)
##
## Pearson residuals:
        Min
                       Median
                                             Max
                  10
                                     30
## -1.22436 -0.71216 0.01381 0.81085
                                         1.68542
## Count model coefficients (poisson with log link):
                         Estimate Std. Error z value Pr(>|z|)
##
## (Intercept)
                          0.63085
                                      0.25643
                                                2.460
                                                        0.0139 *
## homework_releveledyes 0.07149
                                      0.22003
                                                0.325
                                                        0.7453
## gender relevledF
                          0.49764
                                      0.24547
                                                2.027
                                                        0.0426 *
##
```

```
## Zero-inflation model coefficients (binomial with logit link):
##
               Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                 1.4649
                            1.0651
                                     1.375
                                              0.1690
                -1.2981
                            0.5616 -2.312
                                              0.0208 *
## grade
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Number of iterations in BFGS optimization: 10
## Log-likelihood: -82.49 on 5 Df
#Using a deviance test to check the fit of the model
null.model<- zeroinfl(nbooks ~ 1, data=Reading)</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 11.00896 (df=2)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.01167754 (df=2)
#Using the model to predict the number of books read by a second-grade girl f
or whom the reading is part of the homework
print(predict(fitted.model, data.frame(grade=2, gender relevled="F", homework
releveled="yes")))
##
## 2.51027
```

Hurdle Poisson 5.12

```
#First we begin by importing the data
Librarianship <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT
410/Exercise Files/Exercise5.12Data.csv", header = TRUE, sep =",")
#Fitting the model
summary(fitted.model<- hurdle(ncomps ~ nbooks + njrnls | budget,</pre>
data=Librarianship, dist="poisson", zero.dist="binomial", link="logit"))
##
## Call:
## hurdle(formula = ncomps ~ nbooks + njrnls | budget, data = Librarianship,
##
       dist = "poisson", zero.dist = "binomial", link = "logit")
##
## Pearson residuals:
      Min
                10 Median
                                       Max
                                3Q
## -3.3994 -1.2434 -0.7700 0.9434 6.9410
##
## Count model coefficients (truncated poisson with log link):
                Estimate Std. Error z value Pr(>|z|)
## (Intercept) 2.733815
                           0.140691 19.431
                                              <2e-16 ***
## nbooks -0.025741 0.010592 -2.430
                                              0.0151 *
```

```
## njrnls
               0.001832 0.002425
                                      0.755
                                              0.4500
## Zero hurdle model coefficients (binomial with logit link):
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) -0.4086
                            0.6625 -0.617
                                             0.5374
                                             0.0389 *
## budget
                0.4490
                            0.2174
                                   2.065
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Number of iterations in BFGS optimization: 9
## Log-likelihood: -155.5 on 5 Df
#Using devieance test to see how well the model fits
null.model<- hurdle(ncomps ~ 1, data=Librarianship, dist="poisson",</pre>
zero.dist="binomial", link="logit")
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 19.21102 (df=2)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.0002472598 (df=2)
#Using the fitted model to predict the number of computers in a library with
10,000 books, 25 periodicals, and an annual budget of $15 per student
print(predict(fitted.model, data.frame(nbooks=10, njrnls=25, budget=15)))
## 12,43384
```

Hurdle Poisson 5.13

```
#First we begin by importing the data
Medication<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/</pre>
Exercise Files/Exercise5.13Data.csv", header = TRUE, sep =",")
#Releveling gender and making it a categorical variable
gender_releveled<- relevel(as.factor(Medication$gender), ref="M")</pre>
#Fitting the model
summary(fitted.model<- hurdle(ndaysnomeds ~ gender_releveled + age | notherme</pre>
data=Medication, dist="poisson", zero.dist = "binomial", link="logit"))
##
## Call:
## hurdle(formula = ndaysnomeds ~ gender releveled + age | nothermeds, data =
Medication,
##
       dist = "poisson", zero.dist = "binomial", link = "logit")
##
## Pearson residuals:
      Min 10 Median
                                3Q
                                       Max
```

```
## -1.5558 -0.7444 0.1306 0.6857 2.0317
##
## Count model coefficients (truncated poisson with log link):
                     Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                     -1.02120
                                 1.48329 -0.688
                                                    0.4912
                                                    0.0168 *
## gender_releveledF 0.72882
                                 0.30489
                                            2.390
                      0.02155
                                 0.01992
                                           1.082
                                                   0.2794
## age
## Zero hurdle model coefficients (binomial with logit link):
               Estimate Std. Error z value Pr(>|z|)
                                            0.00365 **
## (Intercept)
                 3.0554
                            1.0511
                                     2.907
## nothermeds
                -0.2288
                            0.1122 -2.039 0.04144 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Number of iterations in BFGS optimization: 9
## Log-likelihood: -52.23 on 5 Df
#Using devieance test to see how well the model fits
null.model<- hurdle(ndaysnomeds ~ 1, data=Medication, dist="poisson",</pre>
zero.dist="binomial", link="logit")
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 12.7902 (df=2)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.005113002 (df=2)
#Using the fitted model to predict the number of computers in a library with
10,000 books, 25 periodicals, and an annual budget of $15 per student
print(predict(fitted.model, data.frame(gender_releveled="M", age=78, notherme
ds=0)))
##
## 2.158935
#Negative Binomial 6.2
#First we begin by importing the data
BioMussels<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise6.2Data.csv", header = TRUE, sep =",")
#Specifying the reference level and making feeding level a categorical variab
FeedingLevel releveled<- relevel(as.factor(BioMussels$feeding.level), ref="hi
gh")
#Fitting the model
summary(fitted.model<- glm.nb(ndead.mussels ~ max.temp + min.temp + FeedingLe</pre>
vel_releveled, data=BioMussels))
```

```
##
## Call:
## glm.nb(formula = ndead.mussels ~ max.temp + min.temp + FeedingLevel_releve
       data = BioMussels, init.theta = 10.63638151, link = log)
##
##
## Deviance Residuals:
        Min
                   10
                         Median
                                        30
                                                 Max
## -1.78262
             -0.93015
                       -0.05483
                                   0.43714
                                             1.68131
##
## Coefficients:
##
                               Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                              -11.45070
                                           3.98601 -2.873 0.00407 **
## max.temp
                                0.03026
                                           0.02222
                                                     1.361
                                                            0.17340
                                                     2.048
## min.temp
                                0.14185
                                           0.06925
                                                            0.04052 *
## FeedingLevel releveledlow
                                1.77858
                                           0.48825
                                                     3.643
                                                            0.00027 ***
## FeedingLevel_releveledmed
                                1.41255
                                           0.48347
                                                     2.922 0.00348 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for Negative Binomial(10.6364) family taken to be 1)
##
##
       Null deviance: 75.013 on 23
                                      degrees of freedom
## Residual deviance: 23.855 on 19 degrees of freedom
## AIC: 104.34
## Number of Fisher Scoring iterations: 1
##
##
##
                 Theta:
                         10.6
##
             Std. Err.:
                         12.0
##
    2 x log-likelihood:
                         -92.338
#Using deviance test to see how well the model fits
null.model<- glm.nb(ndead.mussels ~ 1, data = BioMussels)</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 27.27491 (df=2)
print(p.value<- pchisq(deviance, df=4, lower.tail = FALSE))</pre>
## 'log Lik.' 1.749015e-05 (df=2)
#Using the model to predict the number of dead mussels that were fed a high l
evel of food, and were located in an area with a maximum temperature of 75 de
grees and minimum temperature of 60 degrees.
print(predict(fitted.model, data.frame(max.temp=75, min.temp=60,
FeedingLevel releveled="high"), type="response"))
```

```
## 1
## 0.5113223
```

#Negative Binomial 6.3

```
#First we begin by importing the data
HSAllowance<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410
/Exercise Files/Exercise6.3Data.csv", header = TRUE, sep =",")
#Specifying the reference level and making gender and job categorical variabl
gender releveled<- relevel(as.factor(HSAllowance$gender), ref="M")</pre>
job_releveled<- relevel(as.factor(HSAllowance$job), ref="yes")</pre>
#Fitting the model
summary(fitted.model<- glm.nb(allowance ~ age + gender_releveled + job_releve</pre>
led,
data=HSAllowance))
##
## Call:
## glm.nb(formula = allowance ~ age + gender_releveled + job_releveled,
       data = HSAllowance, init.theta = 3.29253741, link = log)
##
## Deviance Residuals:
       Min
                      Median
##
                 10
                                   3Q
                                            Max
                    -0.2815
## -2.5167 -0.8701
                               0.1905
                                         2.5725
##
## Coefficients:
##
                     Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                      -0.1297
                                   2.3982
                                          -0.054
                                                   0.95688
## age
                       0.0359
                                  0.1387
                                            0.259
                                                   0.79570
                       0.4523
                                  0.3067
                                            1.475
## gender releveledF
                                                   0.14033
                                  0.3937
                                            3.208 0.00134 **
## job_releveledno
                       1.2630
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for Negative Binomial(3.2925) family taken to be 1)
##
##
       Null deviance: 48.685 on 29
                                     degrees of freedom
## Residual deviance: 34.148 on 26 degrees of freedom
## AIC: 150.68
##
## Number of Fisher Scoring iterations: 1
##
##
##
                 Theta:
                         3.29
##
             Std. Err.:
                         1.67
##
## 2 x log-likelihood:
                         -140.68
```

```
#Using deviance test to see how well the model fits
null.model<- glm.nb(allowance ~ 1, data=HSAllowance)</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 11.87264 (df=2)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.00783244 (df=2)
#Using the model to predict the amount of weekly allowance for a male student
, age 16, who hasn't held a summer job.
print(predict(fitted.model, data.frame(age=16, gender releveled="M", job rele
veled="no"),
type="response"))
##
## 5.516843
#Zero-Truncated Negative Binomial 6.4
#First we begin by importing the data
StateParks<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise6.4Data.csv", header = TRUE, sep =",")
#Specifying the reference level and making camped a categorical variable
camped_releveled<- relevel(as.factor(StateParks$camped), ref="yes")</pre>
#Fitting the model
summary(fitted.model<- vglm(nkayaks ~ partysize + routelength + camped_releve</pre>
led,
data=StateParks, family=posnegbinomial()))
##
## Call:
## vglm(formula = nkayaks ~ partysize + routelength + camped releveled,
       family = posnegbinomial(), data = StateParks)
##
##
## Coefficients:
##
                      Estimate Std. Error z value Pr(>|z|)
## (Intercept):1
                      -0.85770
                                   0.55260 -1.552
                                                      0.121
## (Intercept):2
                       2.19308
                                   1.15141
                                                NA
                                                         NΔ
## partysize
                       0.18331
                                  0.03440
                                             5.329 9.89e-08 ***
## routelength
                       0.02644
                                  0.03582
                                             0.738
                                                      0.460
## camped releveledno 0.27304
                                  0.29855
                                             0.915
                                                      0.360
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '* 0.05 '.' 0.1 ' ' 1
## Names of linear predictors: loglink(munb), loglink(size)
## Log-likelihood: -48.2332 on 55 degrees of freedom
```

```
##
## Number of Fisher scoring iterations: 13
## Warning: Hauck-Donner effect detected in the following estimate(s):
## '(Intercept):2'
#Using deviance test to see how well the model fits
null.model<- vglm(nkayaks ~ 1, data=StateParks, family=posnegbinomial())</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## [1] 24.63644
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## [1] 1.839315e-05
#Using the model to predict the number of rented kayaks for a party of 5 peop
le who plan to take a 6-hour route and to
#camp overnight.
print(predict(fitted.model, data.frame(partysize=5, routelength=6,
camped_releveled="yes"), type="response"))
##
         [,1]
## 1 1.807214
#Zero-Truncated Negative Binomial 6.5
#First we begin by importing the data
Youtube<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exe
rcise Files/Exercise6.5Data.csv", header = TRUE, sep =",")
#Specifying the reference level and making type a categorical variable
type_releveled<- relevel(as.factor(Youtube$type), ref="fashion")</pre>
#Fitting the model
summary(fitted.model<- vglm(nnewvideos ~ nvideos + nsubscr + nviews + type re</pre>
leveled,
data=Youtube, family = posnegbinomial()))
##
## Call:
## vglm(formula = nnewvideos ~ nvideos + nsubscr + nviews + type releveled,
       family = posnegbinomial(), data = Youtube)
##
```

Estimate Std. Error z value Pr(>|z|)

0.0049484 0.0016068 3.080 0.00207 **

0.930 0.35246

1.657 0.09753 .

0.3625437 0.3899071

0.7660984 0.4623516

type_releveledcomedy 0.3336973 0.5483930 0.609 0.54286

Coefficients:

(Intercept):1

(Intercept):2

nvideos

nsubscr

nviews

##

```
## type_releveledlifeadvice 0.5772579 0.4939778
                                                   1.169 0.24257
## type relevelednews
                            1.5310712 0.5050049
                                                   3.032 0.00243 **
## type_releveledproducts
                            0.3359591 0.4596850
                                                   0.731 0.46487
## type releveledscience
                                                   0.015 0.98789
                            0.0101134 0.6663842
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Names of linear predictors: loglink(munb), loglink(size)
## Log-likelihood: -95.9302 on 70 degrees of freedom
##
## Number of Fisher scoring iterations: 8
##
## No Hauck-Donner effect found in any of the estimates
#Using deviance test to see how well the model fits
null.model<- vglm(nnewvideos ~ 1, data = Youtube, family=posnegbinomial())</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## [1] 20.89688
print(p.value<- pchisq(deviance, df=8, lower.tail = FALSE))</pre>
## [1] 0.007426463
#Using the model to predict the number of new videos for a vlogger who posted
a total
#of 87 videos on popular science, has 50,000 subscribers, and 254,000 views.
print(predict(fitted.model, data.frame(nvideos=87, nsubscr=50, nviews=254,
type_releveled="science"), type="response"))
##
         [,1]
## 1 1.793339
```

#Zero-Inflated Negative Binomial 6.6

```
#First we begin by importing the data
NegBiInsurance<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT
410/Exercise Files/Exercise6.6Data.csv", header = TRUE, sep =",")

#Specifying the reference level and making gender a categorical variable
gender_releveled<- relevel(as.factor(NegBiInsurance$gender), ref="M")

#Fitting the model
summary(fitted.model<- zeroinfl(nclaimspast5ys ~ age + gender_releveled |
nclaimsprev5ys, data=NegBiInsurance, dist="negbin"))

##
## Call:
## zeroinfl(formula = nclaimspast5ys ~ age + gender_releveled | nclaimsprev5y
s,
## data = NegBiInsurance, dist = "negbin")</pre>
```

```
##
## Pearson residuals:
##
       Min
                10 Median
                                3Q
                                       Max
## -1.6473 -0.8728 -0.1159 0.5660 2.2521
##
## Count model coefficients (negbin with log link):
                      Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                     -0.108008
                                 0.536028 -0.201 0.84031
                                            3.221 0.00128 **
## age
                      0.029678
                                 0.009213
## gender_releveledF 0.127863
                                 0.184205
                                            0.694 0.48760
## Log(theta)
                      3.475375
                                 2.344593
                                            1.482 0.13826
##
## Zero-inflation model coefficients (binomial with logit link):
##
                  Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                    0.8826
                               0.7204
                                        1.225
                                                0.2205
## nclaimsprev5ys -1.3297
                               0.5397 -2.464
                                                0.0138 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Theta = 32.3099
## Number of iterations in BFGS optimization: 10
## Log-likelihood: -82.02 on 6 Df
#Using deviance test to see how well the model fits
null.model<- zeroinfl(nclaimspast5ys ~ 1, data=NegBiInsurance, dist="negbin")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 20.2445 (df=3)
#Using the model to predict the number number of claims made in the past five
years by a 55-year-old female
#policyholder who has made no claims in the previous five years.
print(predict(fitted.model, data.frame(nclaimsprev5ys=0, age=55,
gender_releveled="F")))
##
## 1.526963
#Zero-Inflated Negative Binomial 6.7
#First we begin by importing the data
Dentist<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exe</pre>
```

```
#First we begin by importing the data
Dentist<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exe
rcise Files/Exercise6.7Data.csv", header = TRUE, sep =",")

#Specifying the reference level and making gender and oralhygiene categorical
variables
gender_releveled<- relevel(as.factor(Dentist$gender), ref="F")
oralhygiene_releveled<- relevel(as.factor(Dentist$oralhygiene), ref="med")

#Fitting the model
summary(fitted.model<- zeroinfl(DMFTindex ~ gender_releveled + oralhygiene_re</pre>
```

```
leveled | age,
data=Dentist, dist="negbin"))
##
## Call:
## zeroinfl(formula = DMFTindex ~ gender releveled + oralhygiene releveled |
##
       age, data = Dentist, dist = "negbin")
##
## Pearson residuals:
                10 Median
##
       Min
                                3Q
                                        Max
## -1.0646 -0.7071 -0.1889 0.4202 1.9276
##
## Count model coefficients (negbin with log link):
##
                             Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                               1.2352
                                          0.2762 4.472 7.75e-06 ***
## gender releveledM
                               0.6655
                                           0.3130
                                                    2.126
                                                            0.0335 *
## oralhygiene releveledhigh
                               0.2468
                                           0.3623
                                                    0.681
                                                            0.4957
## oralhygiene releveledlow
                                                            0.7022
                               0.1577
                                           0.4125
                                                    0.382
                               0.8514
                                           0.4703
                                                            0.0702 .
## Log(theta)
                                                    1.810
##
## Zero-inflation model coefficients (binomial with logit link):
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) 2.10729
                           1.42547
                                     1.478
                                              0.1393
## age
               -0.10126
                           0.05073 -1.996
                                              0.0459 *
## ---
                   0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
## Theta = 2.3429
## Number of iterations in BFGS optimization: 13
## Log-likelihood: -88.86 on 7 Df
#Using deviance test to see how well the model fits
null.model<- zeroinfl(DMFTindex ~ 1, data=Dentist, dist="negbin")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 12.76369 (df=3)
print(p.value<- pchisq(deviance, df=4, lower.tail = FALSE))</pre>
## 'log Lik.' 0.01249009 (df=3)
#Using the model to find the predicted value of the DMFT index for a man, age
d 28, with a high level of oral hygiene.
print(predict(fitted.model, data.frame(age=28, gender_releveled="M",
oralhygiene releveled="high")))
##
## 5.77492
```

```
#First we begin by importing the data which is taken from 6.6
HNBInsurance<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 41
0/Exercise Files/Exercise6.6Data.csv", header = TRUE, sep =",")
#Specifying the reference level and making gender a categorical variable
gender releveled<- relevel(as.factor(HNBInsurance$gender), ref="M")</pre>
#Fitting the model
summary(fitted.model<- hurdle(nclaimspast5ys ~ age + gender_releveled | nclai</pre>
msprev5ys,
data=HNBInsurance, dist="negbin", zero.dist = "binomial", link="logit"))
##
## Call:
## hurdle(formula = nclaimspast5ys ~ age + gender_releveled | nclaimsprev5ys,
       data = HNBInsurance, dist = "negbin", zero.dist = "binomial", link = "
logit")
##
## Pearson residuals:
        Min
                  10
                       Median
                                    3Q
                                            Max
## -1.58030 -0.89629 -0.09889 0.56517 2.24100
## Count model coefficients (truncated negbin with log link):
##
                      Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                     -0.187878
                                 0.573901 -0.327 0.74339
## age
                      0.030928
                                 0.009815
                                            3.151 0.00163 **
## gender releveledF 0.125341
                                 0.193554
                                            0.648 0.51726
## Log(theta)
                      3.111167
                                 1.872968
                                            1.661 0.09670 .
## Zero hurdle model coefficients (binomial with logit link):
                  Estimate Std. Error z value Pr(>|z|)
## (Intercept)
                   -0.8814
                               0.7050 -1.250
                                                0.2112
                                                0.0115 *
## nclaimsprev5ys
                   1.2816
                               0.5071
                                        2.527
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Theta: count = 22.4472
## Number of iterations in BFGS optimization: 18
## Log-likelihood: -81.69 on 6 Df
#Using deviance test to see how well the model fits
null.model<- hurdle(nclaimspast5ys ~ 1, data=HNBInsurance, dist="negbin",</pre>
zero.dist="binomial", link="logit")
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 20.90545 (df=3)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.0001101446 (df=3)
```

```
#Using the model to find the predicted number of claims made in the past five
years by a 55-year-old female policyholder who has made no claims in the prev
ious five years
print(predict(fitted.model, data.frame(nclaimsprev5ys=0, age=55,
gender releveled="F")))
##
          1
## 1.522423
#Hurdle Negative Binomial 6.9
#First we begin by importing the data
SportsMed<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/E</pre>
xercise Files/Exercise6.9Data.csv", header = TRUE, sep =",")
#Specifying the reference level and making gender a categorical variable
gender_releveled<- relevel(as.factor(SportsMed$gender), ref="F")</pre>
#Fitting the model
summary(fitted.model<- hurdle(ngameinjuries ~ gender releveled +</pre>
nsports|npracticeinjuries, data=SportsMed, dist="negbin",
zero.dist="binomial", link="logit"))
##
## Call:
## hurdle(formula = ngameinjuries ~ gender_releveled + nsports | npracticeinj
uries,
       data = SportsMed, dist = "negbin", zero.dist = "binomial", link = "log
##
it")
##
## Pearson residuals:
                10 Median
       Min
                                 3Q
                                        Max
##
## -1.2907 -0.8023 -0.2172 0.4246 2.4498
## Count model coefficients (truncated negbin with log link):
##
                     Estimate Std. Error z value Pr(>|z|)
```

0.47606

0.19854

0.89322

0.5401

Estimate Std. Error z value Pr(>|z|)

0.8030 -1.441

2.499

0.32510

1.522 0.12796

0.405 0.68525

1.975 0.04825 *

0.1495

0.0124 *

2.729 0.00635 **

(Intercept)

Log(theta)

(Intercept)

npracticeinjuries

Theta: count = 5.8374

Log-likelihood: -59.46 on 6 Df

nsports

##

gender releveledM 0.88730

0.72465

0.08047

1.76429

-1.1574

Number of iterations in BFGS optimization: 17

1.3498

Zero hurdle model coefficients (binomial with logit link):

Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1

```
#Using deviance test to see how well the model fits
null.model<- hurdle(ngameinjuries ~ 1, data=SportsMed, dist="negbin",</pre>
zero.dist="binomial", link="logit")
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 21.23368 (df=3)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 9.414351e-05 (df=3)
#Using the model to find the predicted number of injuries for a male athlete
who throughout his college years has participated in two sports, and who has
received one minor injury during practice games.
print(predict(fitted.model, data.frame(gender releveled="M", nsports=2,
npracticeinjuries=1)))
##
          1
## 3.282217
#Beta 7.2
Birds <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exer</pre>
cise Files/Exercise7.2Data.csv", header = TRUE, sep =",")
#Rescaling predictiors according to the size of their observations and creati
ng response
mass<- Birds$mass/1000
wingspan<- Birds$wingspan/100
distance<- Birds$distance/1000</pre>
propsuccess<- Birds$nmigrated/Birds$nringed</pre>
#Fitting the model
summary(fitted.model<- betareg(propsuccess ~ mass + wingspan + distance,</pre>
link="logit"))
##
## Call:
## betareg(formula = propsuccess ~ mass + wingspan + distance, link = "logit"
)
##
## Standardized weighted residuals 2:
                10 Median
                                30
                                        Max
## -2.4465 -0.7549 0.0988 0.5438 2.3560
## Coefficients (mean model with logit link):
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) 2.78304
                           0.78201 3.559 0.000373 ***
                           0.41088 -0.037 0.970848
## mass
               -0.01502
## wingspan
               0.04487
                           0.93935 0.048 0.961901
## distance -1.31857 0.42237 -3.122 0.001797 **
```

```
##
## Phi coefficients (precision model with identity link):
         Estimate Std. Error z value Pr(>|z|)
                                3.293 0.000991 ***
## (phi)
            4.173
                       1.267
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Type of estimator: ML (maximum likelihood)
## Log-likelihood: 6.406 on 5 Df
## Pseudo R-squared: 0.3677
## Number of iterations: 12 (BFGS) + 5 (Fisher scoring)
#Checking to see how well the model fits
null.model<- betareg(propsuccess ~ 1, link="logit")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 8.55842 (df=2)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.03577627 (df=2)
#Using the model to predict the number of birds that successfully reach the w
inter grounds for a flock of 70 birds with an average mass of 600 grams, an a
verage wingspan of 65 centimeters, that travel a distance of 1650 kilometers.
print(70*predict(fitted.model, data.frame(mass=.6, wingspan=.65, distance=1.6
5,
nringed=70)))
##
          1
## 45.63551
#Beta 7.3
Hospitals <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/</pre>
Exercise Files/Exercise7.3Data.csv", header = TRUE, sep =",")
#Setting reference levels, changing to numeric, and creating response
prophospitalized<- Hospitals$perchospitalized/100</pre>
location.rel<- relevel(as.factor(Hospitals$location), ref="urban")</pre>
type.rel<- relevel(as.factor(Hospitals$type), ref="private")</pre>
#Fitting the model
summary(fitted.model<- betareg(prophospitalized ~ location.rel</pre>
+ type.rel + nbeds, data=Hospitals, link="logit"))
##
## Call:
## betareg(formula = prophospitalized ~ location.rel + type.rel + nbeds,
       data = Hospitals, link = "logit")
##
## Standardized weighted residuals 2:
```

```
Min 10 Median
                                30
                                       Max
## -2.9504 -0.7758 0.1150 0.9524 1.6070
##
## Coefficients (mean model with logit link):
##
                       Estimate Std. Error z value Pr(>|z|)
                     -1.6735254 0.2787233 -6.004 1.92e-09 ***
## (Intercept)
## location.relrural 0.5632707 0.2494671
                                             2.258 0.02395 *
                      0.0116530 0.2511991
                                             0.046 0.96300
## type.relpublic
                      0.0021170 0.0007219
                                             2.933 0.00336 **
## nbeds
##
## Phi coefficients (precision model with identity link):
         Estimate Std. Error z value Pr(>|z|)
                               3.984 6.78e-05 ***
## (phi)
           9.808
                       2.462
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Type of estimator: ML (maximum likelihood)
## Log-likelihood: 20.79 on 5 Df
## Pseudo R-squared: 0.2884
## Number of iterations: 19 (BFGS) + 2 (Fisher scoring)
#Checking to see how well the model fits
null.model<- betareg(prophospitalized ~ 1, data=Hospitals, link="logit")</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 11.00244 (df=2)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.01171267 (df=2)
#Using the model to predict the predicted proportion of hospitalized ER patie
nts for a rural public hospital with 50 beds.
print(predict(fitted.model, data.frame(location.rel="rural", type.rel="public")
nbeds=50)))
##
## 0.270369
#Zero Inflated Beta 7.5
RealEstate <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410
/Exercise Files/Exercise7.5Data.csv", header = TRUE, sep =",")
#Recaling predictors
RealEstate$propsold<- RealEstate$percsold/100
RealEstate$avgprice.res<- RealEstate$avgprice/100
RealEstate$nhouses.res<- RealEstate$nhouses/100
#Fitting the model
```

```
summary(fitted.model<- gamlss(propsold ~ avgprice.res + nhouses.res,</pre>
mu.link="logit", nu.formula = ~ age, nu.link="logit", data=RealEstate,
family = BEZI))
## GAMLSS-RS iteration 1: Global Deviance = 12.5187
## GAMLSS-RS iteration 2: Global Deviance = 11.0988
## GAMLSS-RS iteration 3: Global Deviance = 11.0973
## GAMLSS-RS iteration 4: Global Deviance = 11.0973
## Family: c("BEZI", "Zero Inflated Beta")
##
## Call: gamlss(formula = propsold ~ avgprice.res + nhouses.res,
      nu.formula = ~age, family = BEZI, data = RealEstate,
      mu.link = "logit", nu.link = "logit")
##
## Fitting method: RS()
##
## Mu link function: logit
## Mu Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.16393 0.55373
                                  2.102
                                           0.0462 *
## avgprice.res -0.20024
                       0.09321
                                  -2.148
                                           0.0420 *
## nhouses.res -0.16791
                        0.24421 -0.688
                                          0.4983
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Sigma link function: log
## Sigma Coefficients:
              Estimate Std. Error t value Pr(>|t|)
                                  6.238 1.9e-06 ***
## (Intercept)
                1.703
                           0.273
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Nu link function: logit
## Nu Coefficients:
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -4.51609 1.55696 -2.901 0.00785 **
                                 2.448 0.02205 *
## age
              0.12468
                         0.05093
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## -----
## No. of observations in the fit: 30
## Degrees of Freedom for the fit: 6
        Residual Deg. of Freedom:
##
                                  24
##
                       at cycle:
##
```

```
## Global Deviance:
                       11.09732
##
              AIC:
                       23.09732
##
              SBC:
                       31.50451
## *************************
#Checking to see how well the model fits
null.model<- gamlss(propsold ~ 1, mu.link="logit", nu.formula= ~ 1,</pre>
nu.link="logit", data=RealEstate, family=BEZI)
## GAMLSS-RS iteration 1: Global Deviance = 25.2804
## GAMLSS-RS iteration 2: Global Deviance = 25.2182
## GAMLSS-RS iteration 3: Global Deviance = 25.2182
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 14.12085 (df=3)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 0.002745187 (df=3)
#Using the model to predict the percent houses sold for a subdivision with 30
0 houses, built 50 years ago, and where houses are sold, on average, for $450
,000
param.pred<- predictAll(fitted.model, newdata=data.frame(avgprice.res=4.5,</pre>
nhouses.res=3, age=50), type="response")
print((1-param.pred$nu)*param.pred$mu)
## [1] 0.06694997
#Zero Inflated Beta 7.6
MartialArts <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 41</pre>
0/Exercise Files/Exercise7.6Data.csv", header = TRUE, sep =",")
#Creating response variable
MartialArts$propfirst<- MartialArts$nfirstplaces/MartialArts$ntrophies
#Fitting the model
summary(fitted.model<- gamlss(propfirst ~ nyears + nblackbelts, mu.link="logi</pre>
t",
nu.formula = ~ npupils, nu.link="logit", data=MartialArts, family=BEZI))
## GAMLSS-RS iteration 1: Global Deviance = -0.7884
## GAMLSS-RS iteration 2: Global Deviance = -11.4115
## GAMLSS-RS iteration 3: Global Deviance = -11.578
## GAMLSS-RS iteration 4: Global Deviance = -11.5783
## Family: c("BEZI", "Zero Inflated Beta")
##
## Call: gamlss(formula = propfirst ~ nyears + nblackbelts,
## nu.formula = ~npupils, family = BEZI, data = MartialArts,
```

```
mu.link = "logit", nu.link = "logit")
##
## Fitting method: RS()
## -----
## Mu link function: logit
## Mu Coefficients:
           Estimate Std. Error t value Pr(>|t|)
## nyears 0.12740 0.03938 3.235 0.00415 **
## nblackbelts 0.22669 0.10602 2.138 0.04502 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Sigma link function: log
## Sigma Coefficients:
            Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.7057 0.3077 8.795 2.62e-08 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Nu link function: logit
## Nu Coefficients:
           Estimate Std. Error t value Pr(>|t|)
## (Intercept) 2.99552 1.79441 1.669 0.111
## npupils -0.07133 0.03452 -2.066
                                     0.052 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## No. of observations in the fit: 26
## Degrees of Freedom for the fit: 6
##
      Residual Deg. of Freedom:
                             20
##
                   at cycle: 4
##
## Global Deviance: -11.57825
##
            AIC:
                   0.4217455
            SBC:
                   7,970325
#Checking to see how well the model fits
null.model<- gamlss(propfirst ~ 1, mu.link="logit", nu.formula = ~ 1,</pre>
nu.link="logit", data=MartialArts, family=BEZI)
## GAMLSS-RS iteration 1: Global Deviance = 17.6774
## GAMLSS-RS iteration 2: Global Deviance = 15.8111
## GAMLSS-RS iteration 3: Global Deviance = 15.8073
## GAMLSS-RS iteration 4: Global Deviance = 15.8073
```

```
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 27.38556 (df=3)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 4.887314e-06 (df=3)
#Using the model to predict the proportion of first-place trophies won by a s
tudio that #has been around for 10 years, has 85 students and three black-bel
t instructors.
param.pred<- predictAll(fitted.model, newdata=data.frame(nyears=10,</pre>
nblackbelts=3, npupils=85), type="response")
print((1-param.pred$nu)*param.pred$mu)
## [1] 0.4800583
#One Inflated Beta 7.7
ParksAndRec <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 41
0/Exercise Files/Exercise7.7Data.csv", header = TRUE, sep =",")
#Creating response variable
ParksAndRec$propsurvived<- ParksAndRec$nsurvived/ParksAndRec$nplanted
#Fitting the model
summary(fitted.model<- gamlss(propsurvived ~ pestcontrol + fertilization,</pre>
mu.link="logit", nu.formula = ~ precipitation + windspeed, nu.link="logit",
data=ParksAndRec, family=BEOI))
## GAMLSS-RS iteration 1: Global Deviance = 8.5599
## GAMLSS-RS iteration 2: Global Deviance = 3.9565
## GAMLSS-RS iteration 3: Global Deviance = 3.7543
## GAMLSS-RS iteration 4: Global Deviance = 3.7504
## GAMLSS-RS iteration 5: Global Deviance = 3.7503
## Family: c("BEOI", "One Inflated Beta")
##
## Call: gamlss(formula = propsurvived ~ pestcontrol + fertilization,
##
      nu.formula = ~precipitation + windspeed, family = BEOI,
##
      data = ParksAndRec, mu.link = "logit", nu.link = "logit")
##
## Fitting method: RS()
##
## Mu link function: logit
## Mu Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
                             0.5642 -1.964
                                             0.0643
## (Intercept)
                 -1.1080
## pestcontrol
                  0.3233
                             0.1364
                                      2.371
                                             0.0285 *
## fertilization 0.7327
                         0.2821
                                     2.597
                                             0.0177 *
```

```
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Sigma link function: log
## Sigma Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.6048 0.3015 5.323 3.88e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## -----
## Nu link function: logit
## Nu Coefficients:
               Estimate Std. Error t value Pr(>|t|)
##
## (Intercept)
              6.91004 4.29740 1.608
## precipitation -0.04206
                         0.09075 -0.463
                                          0.6483
## windspeed -0.78224 0.37748 -2.072
                                          0.0521 .
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## ------
## No. of observations in the fit: 26
## Degrees of Freedom for the fit: 7
       Residual Deg. of Freedom: 19
##
                      at cycle:
##
## Global Deviance:
                    3.750299
##
                     17.7503
             AIC:
##
             SBC:
                     26.55697
## ***********************
#Checking to see how well the model fits
null.model<- gamlss(propsurvived ~ 1, mu.link="logit", nu.formula=~1,</pre>
nu.link="logit", data=ParksAndRec, family=BEOI)
## GAMLSS-RS iteration 1: Global Deviance = 23.3027
## GAMLSS-RS iteration 2: Global Deviance = 22.2044
## GAMLSS-RS iteration 3: Global Deviance = 22.1958
## GAMLSS-RS iteration 4: Global Deviance = 22.1958
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 18.4455 (df=3)
print(p.value<- pchisq(deviance, df=4, lower.tail = FALSE))</pre>
## 'log Lik.' 0.001009668 (df=3)
#Using the model to predict the predicted proportion of trees that would surv
ive for two years
```

```
param1<- predictAll(fitted.model, newdata = data.frame(pestcontrol=0,</pre>
fertilization=0, precipitation=2, windspeed=12.5), type="response")
param2<- predictAll(fitted.model, newdata = data.frame(pestcontrol=0,</pre>
fertilization=0, precipitation=25, windspeed=6), type="response")
print(param1$nu+(1-param1$nu)*param1$mu)
## [1] 0.2855535
print(param2$nu+(1-param2$nu)*param2$mu)
## [1] 0.8212778
#one Inflated Beta 7.8
LuggageSales <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 4</pre>
10/Exercise Files/Exercise7.8Data.csv", header = TRUE, sep =",")
#Setting reference level and making gender numeric
LuggageSales$gender.rel<- relevel(as.factor(LuggageSales$gender), ref="M")</pre>
#Fitting the model
summary(fitted.model<- gamlss(propsales ~ gender.rel + bonus, mu.link="logit"</pre>
nu.formula = ~ expyr, nu.link="logit", data=LuggageSales, family=BEOI))
## GAMLSS-RS iteration 1: Global Deviance = -1.75
## GAMLSS-RS iteration 2: Global Deviance = -19.2634
## GAMLSS-RS iteration 3: Global Deviance = -22.4945
## GAMLSS-RS iteration 4: Global Deviance = -22.5962
## GAMLSS-RS iteration 5: Global Deviance = -22.5971
## ***********************************
## Family: c("BEOI", "One Inflated Beta")
## Call: gamlss(formula = propsales ~ gender.rel + bonus, nu.formula = ~expy
r,
       family = BEOI, data = LuggageSales, mu.link = "logit",
##
       nu.link = "logit")
##
## Fitting method: RS()
##
## -
## Mu link function: logit
## Mu Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.3532
                            0.4533 -0.779 0.445044
                            0.2599 -2.702 0.013712 *
## gender.relF -0.7023
                            0.4038 4.492 0.000223 ***
## bonus
                1.8137
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Sigma link function: log
## Sigma Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
                                 9.151 1.38e-08 ***
## (Intercept) 2.9540
                        0.3228
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Nu link function: logit
## Nu Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -3.7949 1.2964 -2.927 0.00833 **
## expyr
               0.4619
                         0.1933
                                 2.389 0.02684 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## No. of observations in the fit: 26
## Degrees of Freedom for the fit:
##
        Residual Deg. of Freedom: 20
##
                      at cycle:
##
## Global Deviance:
                     -22.59715
##
             AIC:
                     -10.59715
##
             SBC:
                     -3.048571
#Checking to see how well the model fits
null.model<- gamlss(propsales ~ 1, mu.link="logit", nu.formula = ~ 1,</pre>
nu.link="logit", data=LuggageSales, family=BEOI)
## GAMLSS-RS iteration 1: Global Deviance = 17.0857
## GAMLSS-RS iteration 2: Global Deviance = 10.0505
## GAMLSS-RS iteration 3: Global Deviance = 9.5774
## GAMLSS-RS iteration 4: Global Deviance = 9.5682
## GAMLSS-RS iteration 5: Global Deviance = 9.568
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 32.16519 (df=3)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 4.830298e-07 (df=3)
#Using the model to predict the proportion of completed sales for a salesman
with 3 years of work experience and who received $1,500 in bonuses the previo
us year.
param<- predictAll(fitted.model, newdata = data.frame(expyr=3, gender.rel="M"</pre>
```

```
bonus=1.5), type="response")
print(param$nu+(1-param$nu)*param$mu)
## [1] 0.9213704
#Zero-One Inflated Beta 7.9
AgLab <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/Exer
cise Files/Exercise7.9Data.csv", header = TRUE, sep =",")
#Rescaling
AgLab$altitudeK<- AgLab$altitude/1000
#Fitting the model
summary(fitted.model<- gamlss(germrate ~ altitudeK, mu.link="logit", nu.formu</pre>
= ~ EC, nu.link="log", tau.formula = ~ soiltemp, tau.link="log", data=AgLab,
family=BEINF))
## GAMLSS-RS iteration 1: Global Deviance = 18.173
## GAMLSS-RS iteration 2: Global Deviance = 17.6438
## GAMLSS-RS iteration 3: Global Deviance = 17.639
## GAMLSS-RS iteration 4: Global Deviance = 17.639
## ***********************************
## Family: c("BEINF", "Beta Inflated")
##
## Call: gamlss(formula = germrate ~ altitudeK, nu.formula = ~EC,
      tau.formula = ~soiltemp, family = BEINF, data = AgLab,
##
      mu.link = "logit", nu.link = "log", tau.link = "log")
##
## Fitting method: RS()
##
## Mu link function: logit
## Mu Coefficients:
              Estimate Std. Error t value Pr(>|t|)
                1.6631
                           0.4713
                                    3.529 0.00224 **
## (Intercept)
## altitudeK
               -0.4790
                           0.1262 -3.796 0.00122 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## -
## Sigma link function:
                        logit
## Sigma Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) -0.5594
                           0.2370 -2.361 0.0291 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## ---
## Nu link function: log
```

```
## Nu Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -13.716 6.161 -2.226
                                        0.0383 *
                                        0.0348 *
## EC
                          2.579
                5.863
                                 2.273
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Tau link function: log
## Tau Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -24.8615 12.9583 -1.919
                                       0.0702 .
                                        0.0780 .
## soiltemp 0.3599
                        0.1932 1.863
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## No. of observations in the fit: 26
## Degrees of Freedom for the fit:
        Residual Deg. of Freedom: 19
##
                      at cycle: 4
##
## Global Deviance:
                    17.63899
##
             AIC:
                     31.63899
##
             SBC:
                     40.44567
#Checking to see how well the model fits
null.model<- gamlss(germrate ~ 1, mu.link="logit", nu.formula = ~ 1,</pre>
nu.link="log", tau.formula = ~ 1, tau.link="log", data=AgLab, family=BEINF)
## GAMLSS-RS iteration 1: Global Deviance = 49.2481
## GAMLSS-RS iteration 2: Global Deviance = 49.2279
## GAMLSS-RS iteration 3: Global Deviance = 49.2278
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 31.5888 (df=4)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 6.389023e-07 (df=4)
#Using the fitted model to predict the germination rate for a plot with EC of
1.5 mS/cm2, soil temperature of 68F, and altitude of 950 feet.
param<- predictAll(fitted.model, newdata = data.frame(EC=1.5, soiltemp=68,</pre>
altitudeK=0.95), type="response")
print((param$tau+param$mu)/(1+param$nu+param$tau))
## [1] 0.8593549
```

```
CollegeFootball <- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STA
T 410/Exercise Files/Exercise7.10Data.csv", header = TRUE, sep =",")
#Fitting the model
summary(fitted.model<- gamlss(propgames ~ vertical + bench, mu.link="logit",</pre>
nu.formula = ~ broad, nu.link="log",tau.formula = ~ BMI + fortyyd, tau.link
="log", data=CollegeFootball, family=BEINF))
## GAMLSS-RS iteration 1: Global Deviance = 15.291
## GAMLSS-RS iteration 2: Global Deviance = 13.6346
## GAMLSS-RS iteration 3: Global Deviance = 13.6244
## GAMLSS-RS iteration 4: Global Deviance = 13.6244
## **********************************
## Family: c("BEINF", "Beta Inflated")
##
## Call: gamlss(formula = propgames ~ vertical + bench, nu.formula = ~broad,
      tau.formula = ~BMI + fortyyd, family = BEINF, data = CollegeFootball,
##
      mu.link = "logit", nu.link = "log", tau.link = "log")
##
## Fitting method: RS()
## -----
## Mu link function: logit
## Mu Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -6.33600 2.44008 -2.597 0.01395 *
## vertical 0.21407
                       0.06938
                               3.086 0.00409 **
## bench
             -0.01399 0.03066 -0.456 0.65117
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## -----
## Sigma link function: logit
## Sigma Coefficients:
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.7429 0.1665 -4.463 8.9e-05 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## -----
## Nu link function: log
## Nu Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) 45.5488 18.5894
                                2.450
## broad
              -0.4148
                        0.1652 -2.511
                                        0.0171 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Tau link function: log
```

```
## Tau Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) -27.0077
                          26.3530 -1.025
                                            0.3129
                                            0.0336 *
## BMI
               -1.0664
                           0.4808 -2.218
## fortyyd
              12.4225
                           5.7375
                                  2.165
                                            0.0377 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## --
## No. of observations in the fit: 42
## Degrees of Freedom for the fit: 9
        Residual Deg. of Freedom: 33
##
##
                        at cycle: 4
##
## Global Deviance:
                       13.62437
##
              AIC:
                       31.62437
##
              SBC:
                       47.26339
#Checking to see how well the model fits
null.model<- gamlss(propgames ~ 1, mu.link="logit", nu.formula = ~ 1,</pre>
nu.link="log", tau.formula = ~ 1, tau.link="log", data=CollegeFootball,
family= BEINF)
## GAMLSS-RS iteration 1: Global Deviance = 47.9109
## GAMLSS-RS iteration 2: Global Deviance = 47.2882
## GAMLSS-RS iteration 3: Global Deviance = 47.2861
## GAMLSS-RS iteration 4: Global Deviance = 47.2861
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 33.66176 (df=4)
print(p.value<- pchisq(deviance, df=5, lower.tail = FALSE))</pre>
## 'log Lik.' 2.780153e-06 (df=4)
#sing the fitted model to predict the proportion of games that a new player w
ill play, if his BMI 27.8 kg/m2, forty-dash run is 4.67 seconds, the vertical
jump is 32 inches, broad jump is 117 inches, and bench press is 16 repetition
5.
param<- predictAll(fitted.model, newdata = data.frame(BMI=27.8, fortyyd=4.67,</pre>
vertical=32, broad=117, bench=16), type="response")
print((param$tau+param$mu)/(1+param$nu+param$tau))
## [1] 0.9032484
#Random Slope and Intercept for Normal Response 8.2
```

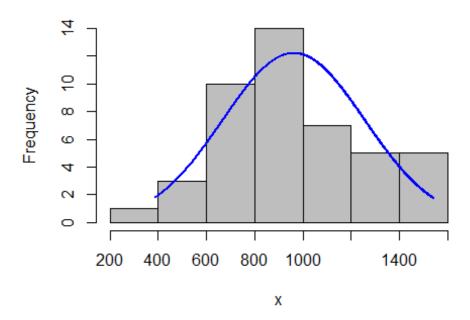
DepartmentStore<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT

410/Exercise Files/Exercise8.2Data.csv", header = TRUE, sep =",")

```
#Creating a long form data set
LongForm<- melt(DepartmentStore, id.vars=c("id","totalyears","status"),
variable.name="bonus.year", value.name="bonus")

#Creating response variable
year<- ifelse(LongForm$bonus.year=="bonus18",1.8,
ifelse(LongForm$bonus.year=="bonus19",1.9,2.0))

#Checking normality of reponse variable
plotNormalHistogram(LongForm$bonus)</pre>
```



```
shapiro.test(LongForm$bonus)

##

## Shapiro-Wilk normality test

##

## data: LongForm$bonus

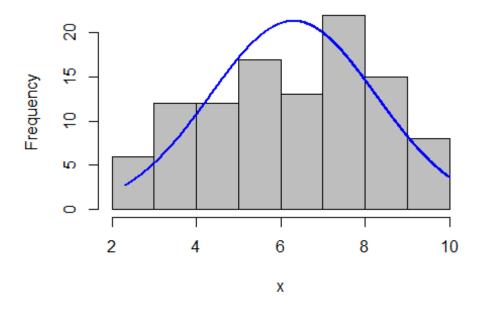
## W = 0.96686, p-value = 0.222

#Creating reference Level for status
status.rel<- relevel(as.factor(LongForm$status), ref="part")

#fitting the model
summary(fitted.model<- lme(bonus ~ totalyears + status.rel + year, random = ~ 1 + year | id, data=LongForm))</pre>
```

```
## Linear mixed-effects model fit by REML
##
     Data: LongForm
##
          AIC
                   BIC
                          logLik
##
     529.3133 543.0219 -256.6567
##
## Random effects:
## Formula: ~1 + year | id
## Structure: General positive-definite, Log-Cholesky parametrization
               StdDev
## (Intercept) 1499.85877 (Intr)
## year
               797.95782 -0.998
## Residual
                67.88526
##
## Fixed effects:
                   bonus ~ totalyears + status.rel + year
                       Value Std.Error DF
                                            t-value p-value
##
## (Intercept)
                  -2246.1138 457.2023 29 -4.912735
## totalyears
                     58.8982
                                8.2368 12 7.150631
                                                      0.000
## status.relfull
                     53.9643
                               63.3473 12 0.851879
                                                      0.411
## year
                   1394.6667 240.4381 29 5.800523
                                                      0.000
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.111
## status.relfull 0.013 -0.600
                  -0.990 0.000 0.000
## year
##
## Standardized Within-Group Residuals:
           Min
                        Q1
                                   Med
                                                Q3
                                                           Max
## -1.55954603 -0.32012837 -0.04131555 0.31523299 1.89915223
##
## Number of Observations: 45
## Number of Groups: 15
intervals(fitted.model)
## Approximate 95% confidence intervals
##
  Fixed effects:
##
                                     est.
##
                        lower
                                                upper
## (Intercept)
                  -3181.19760 -2246.11380 -1311.03001
## totalyears
                     40.95181
                                 58.89822
                                             76.84464
## status.relfull
                    -84.05771
                                 53.96428
                                            191.98628
                    902.91556 1394.66667 1886.41777
## year
## attr(,"label")
## [1] "Fixed effects:"
##
  Random Effects:
##
     Level: id
##
##
                               lower
                                             est.
                                                         upper
## sd((Intercept))
                         891.8860755 1499.8587688 2522.2686935
## sd(year)
                         476.3219559 797.9578238 1336.7779518
```

```
## cor((Intercept), year) -0.9995787 -0.9983042 -0.9931881
##
## Within-group standard error:
##
      lower est.
                        upper
## 47.49499 67.88526 97.02937
#Checking to see how well the model fits
null.model<- glm(bonus ~ totalyears + status.rel + year, data=LongForm)</pre>
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 45.81206 (df=5)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 6.218073e-10 (df=5)
#Using fitted model to predict the predicted bonus in 2021 for a full-time em
ployee who has been with the company for 7 years
#print(predict(fitted.model, data.frame(status.rel="full", totalyears=7, year
=2.1), level=0))
#Random Slope and Intercept for Normal Response 8.3
Orthopedic<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise8.3Data.csv", header = TRUE, sep =",")
#Creating a long form data set
data1<- melt(Orthopedic[,c("id","gender","age","doctor","length1",</pre>
"length2", "length3")], id. vars=c("id", "gender", "age", "doctor"),
variable.name="length.visit",value.name="length")
data2<- melt(Orthopedic[,c("score1", "score2", "score3")],</pre>
variable.name="score.visit", value.name="score")
## No id variables; using all as measure variables
Longform<- cbind(data1,data2)</pre>
#Creating response variable
visit<- ifelse(Longform$score.visit=="score1",1,</pre>
ifelse(Longform$score.visit=="score2",2,3))
#Checking normality of reponse variable
plotNormalHistogram(Longform$score)
```



```
shapiro.test(Longform$score)
##
##
    Shapiro-Wilk normality test
##
## data: Longform$score
## W = 0.97598, p-value = 0.05332
#Creating reference levels
gender.rel<- relevel(as.factor(Longform$gender), ref="M")</pre>
doctor.rel<- relevel(as.factor(Longform$doctor), ref="B")</pre>
#fitting the model
summary(fitted.model<- lme(score ~ gender.rel + age + doctor.rel</pre>
+ length + visit, random = ~ 1 + visit | id, data=Longform))
## Linear mixed-effects model fit by REML
     Data: Longform
##
##
          AIC
                   BIC
                           logLik
##
     271.0585 297.0097 -125.5293
##
## Random effects:
    Formula: ~1 + visit | id
    Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                          Corr
## (Intercept) 1.5090310 (Intr)
## visit
               0.3709390 -0.245
```

```
## Residual 0.2852501
##
## Fixed effects:
                  score ~ gender.rel + age + doctor.rel + length + visit
                  Value Std.Error DF t-value p-value
## (Intercept) 5.221580 2.5707920 68 2.031117 0.0462
## gender.relF 2.084700 0.5198854 31 4.009921
                                                0.0004
               0.002997 0.0385434 31 0.077755 0.9385
## age
## doctor.relA 0.158469 0.5084557 31 0.311668 0.7574
## length
              -0.010510 0.0047844 68 -2.196820 0.0314
## visit
               -0.046100 0.0714364 68 -0.645336 0.5209
## Correlation:
##
               (Intr) gndr.F age
                                 dctr.A length
## gender.relF 0.033
## age
               -0.979 -0.166
## doctor.relA -0.213 0.086 0.095
## length
              -0.083 0.114 0.014 0.042
## visit
              -0.030 -0.005 -0.001 -0.002 -0.043
##
## Standardized Within-Group Residuals:
          Min
                       Q1
                                  Med
                                                          Max
                                               Q3
## -1.89038162 -0.46995687 0.01261292 0.40384503 1.67790854
## Number of Observations: 105
## Number of Groups: 35
intervals(fitted.model)
## Approximate 95% confidence intervals
##
## Fixed effects:
##
                    lower
                                  est.
                                               upper
## (Intercept) 0.09164411 5.221579728 10.3515153426
## gender.relF 1.02438617 2.084699512 3.1450128591
## age
              -0.07561292 0.002996939 0.0816068002
## doctor.relA -0.87853313 0.158469127 1.1954713812
              -0.02005743 -0.010510384 -0.0009633373
## length
## visit
              -0.18864960 -0.046100468 0.0964486669
## attr(,"label")
## [1] "Fixed effects:"
##
##
  Random Effects:
    Level: id
##
##
                              lower
                                          est.
                                                   upper
## sd((Intercept))
                          1.1473569 1.5090310 1.9847134
## sd(visit)
                          0.2705064 0.3709390 0.5086597
## cor((Intercept), visit) -0.5845323 -0.2446523 0.1682708
##
##
   Within-group standard error:
      lower
                 est.
                          upper
## 0.2246113 0.2852501 0.3622595
```

```
#Checking to see how well the model fits
null.model<- glm(score ~ gender.rel + age + doctor.rel + length + visit,</pre>
data=Longform)
print(deviance<- -2*(logLik(null.model)-logLik(fitted.model)))</pre>
## 'log Lik.' 72.63075 (df=7)
print(p.value<- pchisq(deviance, df=3, lower.tail = FALSE))</pre>
## 'log Lik.' 1.166255e-15 (df=7)
#Using fitted model to predict the quality of service score that would be giv
en by a 55-year-old male patient on his fourth visit to Dr. A with a 30-minut
e appointment.
#print(predict(fitted.model, data.frame(id=136,gender.rel="M",age=55,doctor.r
el="A", length=30, visit=4), level=0))
#Random Slope and Intercept with Variance Structure for Error 8.6
#Uses data from 8.2
DepartmentStore<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT
410/Exercise Files/Exercise8.2Data.csv", header = TRUE, sep =",")
#Creating Longform data
Longform<- melt(DepartmentStore, id.vars=c("id", "totalyears", "status"),</pre>
variable.name="bonus.year", value.name="bonus")
#Creating variables
year<- ifelse(Longform$bonus.year=="bonus18",1.8,</pre>
ifelse(Longform$bonus.year=="bonus19",1.9,2.0))
#Rescaling and reference
status.rel<- relevel(as.factor(Longform$status), ref="part")</pre>
#Fitting random slope and intercept model with unstructured covariance matrix
of error terms
summary(un.fitted.model<- lme(bonus ~ totalyears + status.rel + year,</pre>
random = ~ 1 + year | id, data=Longform, correlation=corSymm(),
weights=varIdent(form = ~ id | year)))
## Linear mixed-effects model fit by REML
## Data: Longform
##
         AIC
                  BIC
                         logLik
##
     529.393 551.6694 -251.6965
##
## Random effects:
## Formula: ~1 + year | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                         Corr
## (Intercept) 127.86136 (Intr)
         72.16169 -0.815
## year
```

```
## Residual
               108.15388
##
## Correlation Structure: General
## Formula: ~1 | id
## Parameter estimate(s):
## Correlation:
##
   1
## 2 -0.247
## 3 -0.616 0.735
## Variance function:
## Structure: Different standard deviations per stratum
## Formula: ~id | year
## Parameter estimates:
         1.8
                   1.9
## 1.0000000 0.8826214 0.8955668
## Fixed effects: bonus ~ totalyears + status.rel + year
##
                       Value Std.Error DF
                                           t-value p-value
## (Intercept)
                  -2032.3403 338.3799 29 -6.006090 0.0000
## totalyears
                                7.9297 12 7.717549 0.0000
                     61.1977
## status.relfull
                     40.7302
                               60.9854 12 0.667868
                                                     0.5169
                   1272.6088 174.2924 29 7.301573 0.0000
## year
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.145
## status.relfull 0.017 -0.600
## year
                  -0.983 0.000 0.000
##
## Standardized Within-Group Residuals:
           Min
                                   Med
                        Q1
                                                Q3
                                                           Max
## -2.02149053 -0.42338681 -0.05195106 0.66991469 2.11525768
## Number of Observations: 45
## Number of Groups: 15
#AICC
n<- 45
p<- 14
print(AICC<- -2*logLik(un.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 545.393 (df=13)
#Fitting random slope and intercept model with Toeplitz covariance matrix of
error terms
summary(toep.fitted.model<- lme(bonus ~ totalyears + status.rel + year,</pre>
random = ~ 1 + year | id, data=Longform,
correlation=corARMA(form = ~ 1 | id, p=1, q=1)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
                   BIC
##
          AIC
                          logLik
##
     532.5445 549.6802 -256.2722
```

```
##
## Random effects:
## Formula: ~1 + year | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                            Corr
## (Intercept) 4.092560e-02 (Intr)
               2.105377e-05 0
## year
## Residual
               1.299681e+02
##
## Correlation Structure: ARMA(1,1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi1
                  Theta1
## 0.09194164 0.99999473
## Fixed effects: bonus ~ totalyears + status.rel + year
                       Value Std.Error DF
                                            t-value p-value
## (Intercept)
                  -2263.5843 443.8512 29 -5.099872 0.0000
## totalyears
                     60.9362
                                8.0566 12 7.563516
                                                      0.0000
## status.relfull
                     42.5230
                               61.9615 12
                                           0.686280
                                                      0.5056
## year
                   1394.6667 231.2558 29
                                           6.030839
                                                      0.0000
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.112
## status.relfull 0.013 -0.600
## year
                  -0.990 0.000 0.000
## Standardized Within-Group Residuals:
##
          Min
                      Q1
                                Med
                                             Q3
                                                       Max
## -1.7345232 -0.7151969 -0.0438371 0.8018594 1.9381448
## Number of Observations: 45
## Number of Groups: 15
#AICC
p < -11
print(AICC<- -2*logLik(toep.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 542.5445 (df=10)
#Fitting random slope and intercept model with spatial power covariance matri
x of error terms
summary(sppow.fitted.model<- lme(bonus ~ totalyears + status.rel + year,</pre>
random = ~ 1 + year | id, data=Longform,
correlation=corCAR1(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
##
     532.8405 548.2627 -257.4203
##
## Random effects:
```

```
## Formula: ~1 + year | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                            Corr
## (Intercept) 1.682603e-02 (Intr)
## year
               3.010918e-03 0
## Residual
               1.309026e+02
##
## Correlation Structure: Continuous AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi
## 0.5467992
## Fixed effects: bonus ~ totalyears + status.rel + year
                       Value Std.Error DF
                                            t-value p-value
## (Intercept)
                  -2253.4538 386.4705 29 -5.830856 0.0000
## totalyears
                     59.8379
                                8.9004 12 6.723065 0.0000
## status.relfull
                     48.2277
                               68.4510 12 0.704558 0.4945
                   1394.6667 200.1013 29 6.969804 0.0000
## year
## Correlation:
##
                  (Intr) ttlyrs stts.r
                  -0.142
## totalyears
## status.relfull 0.017 -0.600
## year
                  -0.984 0.000 0.000
##
## Standardized Within-Group Residuals:
          Min
                      Q1
                                Med
                                            Q3
                                                       Max
## -1.7491915 -0.7551994 -0.0889853 0.7951945 1.8956271
##
## Number of Observations: 45
## Number of Groups: 15
#AICC
p<- 10
print(AICC<- -2*logLik(sppow.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 541.3111 (df=9)
#Fitting random intercept-only model with spatial power covariance matrix of
error terms
summary(sppowint.fitted.model<- lme(bonus ~ totalyears + status.rel + year,</pre>
random = ~ 1 | id, data=Longform,
correlation=corCAR1(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
     Data: Longform
##
##
          AIC
                   BIC
                          logLik
##
     528.8405 540.8355 -257.4203
##
## Random effects:
## Formula: ~1 | id
           (Intercept) Residual
```

```
## StdDev: 0.01464535 130.9026
##
## Correlation Structure: Continuous AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi
## 0.5467992
## Fixed effects: bonus ~ totalyears + status.rel + year
                       Value Std.Error DF
                                          t-value p-value
                  -2253.4538 386.4705 29 -5.830856 0.0000
## (Intercept)
## totalyears
                     59.8379
                                8.9004 12 6.723065 0.0000
## status.relfull
                               68.4510 12 0.704558 0.4945
                     48.2277
## year
                   1394.6667 200.1013 29 6.969804 0.0000
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.142
## status.relfull 0.017 -0.600
## year
                  -0.984 0.000 0.000
##
## Standardized Within-Group Residuals:
                        Q1
          Min
                                   Med
                                                Q3
                                                           Max
## -1.74919151 -0.75519942 -0.08898529 0.79519451 1.89562705
##
## Number of Observations: 45
## Number of Groups: 15
#AICC
p<- 8
print(AICC<- -2*logLik(sppowint.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 534.8405 (df=7)
#Fitting random slope and intercept model with autoregressive covariance matr
ix of error terms
summary(ar.fitted.model<- lme(bonus ~ totalyears + status.rel + year,</pre>
random = ~ 1 + year | id, data=Longform,
correlation=corAR1(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
     532.8405 548.2627 -257.4203
##
##
## Random effects:
## Formula: ~1 + year | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                            Corr
## (Intercept)
                 0.02036143 (Intr)
## year
                 0.00073145 0
## Residual
              130.90256777
##
```

```
## Correlation Structure: AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi
## 0.5467992
## Fixed effects: bonus ~ totalyears + status.rel + year
                       Value Std.Error DF
                                           t-value p-value
                  -2253.4538 386.4705 29 -5.830856 0.0000
## (Intercept)
## totalyears
                     59.8379
                                8.9004 12 6.723065 0.0000
## status.relfull
                     48.2277 68.4510 12 0.704558 0.4945
## year
                   1394.6667 200.1013 29 6.969804 0.0000
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.142
## status.relfull 0.017 -0.600
## year
                  -0.984 0.000 0.000
##
## Standardized Within-Group Residuals:
##
          Min
                      01
                                            03
                                Med
                                                      Max
## -1.7491915 -0.7551994 -0.0889853 0.7951945 1.8956271
##
## Number of Observations: 45
## Number of Groups: 15
#AICC
p<- 10
print(AICC<- -2*logLik(ar.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 541.3111 (df=9)
#Fitting random slope and intercept model with compound symmetric covariance
matrix of error terms
summary(cs.fitted.model<- lme(bonus ~ totalyears + status.rel + year,</pre>
random = ~ 1 + year | id, data=Longform,
correlation=corCompSymm(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
##
     531.3133 546.7355 -256.6567
##
## Random effects:
## Formula: ~1 + year | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                          Corr
## (Intercept) 1499.84438 (Intr)
## year
                797.95695 -0.998
## Residual
                 68.15637
##
## Correlation Structure: Compound symmetry
## Formula: ~1 | id
```

```
## Parameter estimate(s):
##
           Rho
## 0.007941571
## Fixed effects:
                   bonus ~ totalyears + status.rel + year
                       Value Std.Error DF
##
                                            t-value p-value
## (Intercept)
                  -2246.1138 457.2018 29 -4.912741
                                                       0.000
## totalyears
                                8.2368 12 7.150613
                                                       0.000
                     58.8982
                               63.3475 12 0.851878
## status.relfull
                     53.9643
                                                       0.411
                   1394.6667 240.4378 29 5.800529
## year
                                                      0.000
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.111
## status.relfull 0.013 -0.600
## year
                  -0.990 0.000 0.000
##
## Standardized Within-Group Residuals:
           Min
                        Q1
                                   Med
                                                Q3
                                                            Max
## -1.55566258 -0.31195060 -0.04880253 0.30786515 1.89838030
##
## Number of Observations: 45
## Number of Groups: 15
#AICC
p<- 10
print(AICC<- -2*logLik(cs.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 539.7839 (df=9)
#Fitting random slope and intercept model with independent covariance matrix
of error terms
summary(ind.fitted.model<- lme(bonus ~ totalyears + status.rel+ year,</pre>
random = ~ 1 + year | id, data=Longform))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
##
     529.3133 543.0219 -256.6567
##
## Random effects:
## Formula: ~1 + year | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                          Corr
## (Intercept) 1499.85877 (Intr)
## year
                797.95782 -0.998
## Residual
                 67.88526
## Fixed effects:
                   bonus ~ totalyears + status.rel + year
##
                                            t-value p-value
                       Value Std.Error DF
                  -2246.1138 457.2023 29 -4.912735
## (Intercept)
                                                       0.000
## totalyears
                     58.8982
                                8.2368 12 7.150631
                                                       0.000
## status.relfull
                     53.9643
                               63.3473 12 0.851879
                                                      0.411
```

```
1394.6667 240.4381 29 5.800523
## year
                                                      0.000
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.111
## status.relfull 0.013 -0.600
## year
                  -0.990 0.000 0.000
##
## Standardized Within-Group Residuals:
           Min
                        Q1
                                                Q3
                                                           Max
## -1.55954603 -0.32012837 -0.04131555 0.31523299 1.89915223
## Number of Observations: 45
## Number of Groups: 15
#AICC
p<- 8
print(AICC<- -2*logLik(ind.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 533.3133 (df=8)
#Fitting random intercept-only model with spatial power covariance matrix of
error terms
summary(sppowint.fitted.model<- lme(bonus ~ totalyears + status.rel + year, r</pre>
andom = ~ 1 | id, data=Longform, correlation=corCAR1(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          ATC
                   BIC
                          logLik
##
     528.8405 540.8355 -257.4203
##
## Random effects:
## Formula: ~1 | id
           (Intercept) Residual
##
## StdDev: 0.01464535 130.9026
##
## Correlation Structure: Continuous AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi
## 0.5467992
## Fixed effects: bonus ~ totalyears + status.rel + year
                       Value Std.Error DF
##
                                          t-value p-value
## (Intercept)
                  -2253.4538 386.4705 29 -5.830856 0.0000
## totalyears
                     59.8379
                                8.9004 12 6.723065 0.0000
## status.relfull
                     48.2277
                               68.4510 12
                                          0.704558
                                                     0.4945
                   1394.6667 200.1013 29 6.969804 0.0000
## year
## Correlation:
##
                  (Intr) ttlyrs stts.r
## totalyears
                  -0.142
## status.relfull 0.017 -0.600
                  -0.984 0.000 0.000
## year
```

```
##
## Standardized Within-Group Residuals:
           Min
                        Q1
                                   Med
                                                Q3
                                                           Max
## -1.74919151 -0.75519942 -0.08898529 0.79519451 1.89562705
##
## Number of Observations: 45
## Number of Groups: 15
intervals(sppowint.fitted.model)
## Approximate 95% confidence intervals
##
## Fixed effects:
##
                        lower
                                     est.
                                                upper
## (Intercept)
                  -3043.87467 -2253.45377 -1463.03287
## totalyears
                     40.44562
                                 59.83791
                                             79.23019
## status.relfull -100.91419
                                 48.22769
                                            197.36956
## year
                    985.41361 1394.66667 1803.91972
## attr(,"label")
## [1] "Fixed effects:"
##
##
  Random Effects:
##
    Level: id
##
                          lower
                                      est.
                                                  upper
## sd((Intercept)) 1.224958e-49 0.01464535 1.750969e+45
##
## Correlation structure:
##
           lower
                      est.
## Phi 0.2728708 0.5467992 0.7950434
## attr(,"label")
## [1] "Correlation structure:"
##
## Within-group standard error:
##
       lower
                  est.
  99.34883 130.90257 172.47795
##
getVarCov(sppowint.fitted.model, type="conditional")
## id 1
## Conditional variance covariance matrix
           1
                   2
## 1 17135.0 9369.7 5123.3
## 2 9369.7 17135.0 9369.7
## 3 5123.3 9369.7 17135.0
     Standard Deviations: 130.9 130.9 130.9
##
#Using the fitted model to predict the predicted bonus in 2021 for a full-tim
e employee who has been with the company for 7 years
#print(predict(sppowint.fitted.model, data.frame(totalyears=7, status.rel="fu
ll",year=2.1), level=0))
```

```
#Uses data from 8.3
Orthopedic<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise8.3Data.csv", header = TRUE, sep =",")
#Creating Longform data
data1<- melt(Orthopedic[,c("id","gender","age","doctor", "length1",</pre>
"length2", "length3")], id.vars=c("id", "gender", "age", "doctor"), variable.nam
="length.visit", value.name="length")
data2<- melt(Orthopedic[,c("score1","score2","score3")], variable.name</pre>
="score.visit", value.name="score")
## No id variables; using all as measure variables
Longform<- cbind(data1,data2)</pre>
#Creating variable for time
visit<- ifelse(Longform$score.visit=="score1",1,</pre>
ifelse(Longform$score.visit=="score2",2,3))
#Refernce Levels
gender.rel<- relevel(as.factor(Longform$gender), ref="M")</pre>
doctor.rel<- relevel(as.factor(Longform$doctor), ref="B")</pre>
#Fitting the model with untructured covariance matrix of error terms
#summary(un.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + length
#visit, random = ~ 1 + visit | id, control= lmeControl(opt="optim"),
#data=Longform, correlation=corSymm(), weights=varIdent(form=~id|Length)))
#DOESN'T CONVERGE
#fitting random intercept-only model with
#untructured covariance matrix of error terms
#summary(un.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + length
#visit, random = ~ 1 | id, control= lmeControl(opt="optim"),
#data=Longform,correlation=corSymm(), weights=varIdent(form=~id|length)))
#DOESN'T CONVERGE
#Fitting random slope and intercept model with Toeplitz covariance matrix of
error terms
summary(toep.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + lengt</pre>
visit, random = ~ 1 + visit | id, data=Longform,
correlation = corARMA(form = ~ 1 | id, p=1, q=1)))
```

```
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
         AIC
                  BIC
                          logLik
##
     274.9041 306.0456 -125.4521
##
## Random effects:
## Formula: ~1 + visit | id
## Structure: General positive-definite, Log-Cholesky parametrization
              StdDev
                         Corr
## (Intercept) 1.5680940 (Intr)
## visit
              0.4221922 -0.33
## Residual
              0.1746268
##
## Correlation Structure: ARMA(1,1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi1
                 Theta1
## -0.9999999 0.7063602
## Fixed effects:
                  score ~ gender.rel + age + doctor.rel + length + visit
##
                   Value Std.Error DF t-value p-value
## (Intercept) 5.032941 2.5636381 68 1.963203 0.0537
## gender.relF 2.065154 0.5186892 31 3.981486 0.0004
## age
               0.005701 0.0384599 31 0.148241 0.8831
## doctor.relA 0.195455 0.5074278 31
                                      0.385188 0.7027
## length
               -0.010430 0.0047788 68 -2.182628 0.0325
## visit
               -0.046152 0.0714296 68 -0.646117 0.5204
## Correlation:
##
               (Intr) gndr.F age dctr.A length
## gender.relF 0.034
              -0.979 -0.167
## age
## doctor.relA -0.213 0.086 0.095
## length
              -0.075 0.112 0.005 0.043
## visit
               -0.031 -0.005 0.000 -0.002 -0.043
##
## Standardized Within-Group Residuals:
          Min
                        01
                                   Med
                                                03
                                                           Max
## -2.27827542 -0.43564923 -0.01232409 0.47950177 2.27827475
##
## Number of Observations: 105
## Number of Groups: 35
#AICC
n<-105
p<- 12
print(AICC<- -2*logLik(toep.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 278.2954 (df=12)
#fitting random slope and intercept model with spatial power covariance matri
x of error terms
```

```
summary(sppow.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + leng</pre>
th +
visit, random = ~ 1 + visit| id, data=Longform,
correlation=corCAR1(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
##
     273.0585 301.6048 -125.5293
##
## Random effects:
## Formula: ~1 + visit | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                        Corr
## (Intercept) 1.509031 (Intr)
## visit
               0.370939 -0.245
## Residual
               0.285250
##
## Correlation Structure: Continuous AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##
           Phi
## 1.33668e-07
## Fixed effects:
                  score ~ gender.rel + age + doctor.rel + length + visit
##
                   Value Std.Error DF
                                      t-value p-value
## (Intercept) 5.221582 2.5707924 68 2.031118 0.0462
## gender.relF 2.084699 0.5198855 31 4.009920 0.0004
                0.002997 0.0385434 31 0.077754 0.9385
## age
## doctor.relA 0.158469 0.5084558 31 0.311667 0.7574
## length
               -0.010510 0.0047844 68 -2.196820 0.0314
               -0.046100 0.0714364 68 -0.645336 0.5209
## visit
## Correlation:
               (Intr) gndr.F age dctr.A length
## gender.relF 0.033
## age
               -0.979 -0.166
## doctor.relA -0.213 0.086 0.095
## length
               -0.083 0.114 0.014 0.042
               -0.030 -0.005 -0.001 -0.002 -0.043
## visit
##
## Standardized Within-Group Residuals:
           Min
                        Q1
                                   Med
                                                03
                                                           Max
## -1.89038180 -0.46995700 0.01261281 0.40384486 1.67790864
##
## Number of Observations: 105
## Number of Groups: 35
#AICC
p<- 11
print(AICC<- -2*logLik(sppow.fitted.model)+2*p*n/(n-p-1))</pre>
```

```
## 'log Lik.' 275.8972 (df=11)
#Fitting random slope and intercept model with autoregressive covariance matr
ix of error terms
summary(ar.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + length</pre>
visit, random = ~ 1 + visit| id, data=Longform,
correlation=corAR1(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
##
     272.9041 301.4504 -125.4521
##
## Random effects:
## Formula: ~1 + visit | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                         Corr
## (Intercept) 1.5680951 (Intr)
## visit
               0.4221926 -0.33
## Residual
               0.1746265
##
## Correlation Structure: AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
## Phi
## -1
## Fixed effects: score ~ gender.rel + age + doctor.rel + length + visit
                   Value Std.Error DF
                                      t-value p-value
## (Intercept) 5.032941 2.5636400 68 1.963201 0.0537
## gender.relF 2.065154 0.5186896 31 3.981483
                                                 0.0004
## age
                0.005701 0.0384599 31 0.148241 0.8831
## doctor.relA 0.195455 0.5074282 31
                                      0.385187
                                                 0.7027
## length
               -0.010430 0.0047788 68 -2.182628 0.0325
               -0.046152 0.0714297 68 -0.646116 0.5204
## visit
## Correlation:
               (Intr) gndr.F age
                                    dctr.A length
## gender.relF 0.034
               -0.979 -0.167
## age
## doctor.relA -0.213 0.086 0.095
## length
               -0.075 0.112 0.005 0.043
               -0.031 -0.005 0.000 -0.002 -0.043
## visit
##
## Standardized Within-Group Residuals:
##
           Min
                        Q1
                                   Med
                                                Q3
                                                           Max
## -2.27827948 -0.43564986 -0.01232358 0.47950317 2.27827946
## Number of Observations: 105
## Number of Groups: 35
```

```
#AICC
p<- 11
print(AICC<- -2*logLik(ar.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 275.7428 (df=11)
#Fitting random intercept-only model with autoregressive covariance matrix of
summary(arint.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + leng</pre>
visit, random = ~ 1 | id, data=Longform, correlation=corAR1(form = ~ 1 |
id)))
## Linear mixed-effects model fit by REML
     Data: Longform
##
         AIC
                  BIC
                        logLik
     272.612 295.9681 -127.306
##
##
## Random effects:
## Formula: ~1 | id
            (Intercept) Residual
## StdDev: 0.0005791566 1.57967
##
## Correlation Structure: AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi
## 0.9396084
## Fixed effects: score ~ gender.rel + age + doctor.rel + length + visit
##
                   Value Std.Error DF
                                       t-value p-value
## (Intercept) 4.686268 2.6627508 68 1.759935 0.0829
## gender.relF 2.229715 0.5388793 31 4.137689
                                                 0.0002
## age
                0.010122 0.0398445 31 0.254043 0.8011
## doctor.relA 0.167042 0.5250427 31 0.318150 0.7525
               -0.011298 0.0051494 68 -2.194026 0.0317
## length
## visit
               -0.045594 0.0647041 68 -0.704658 0.4834
## Correlation:
               (Intr) gndr.F age dctr.A length
## gender.relF 0.028
## age
               -0.978 -0.163
## doctor.relA -0.211 0.083 0.095
## length
               -0.101 0.138 0.030 0.016
## visit
               -0.043 -0.007 -0.002 -0.001 -0.051
##
## Standardized Within-Group Residuals:
##
                        Q1
                                   Med
                                                Q3
## -2.23484469 -0.65331876 0.02324913 0.52261910 2.69537024
## Number of Observations: 105
## Number of Groups: 35
```

```
getVarCov(arint.fitted.model, type="conditional")
## id 101
## Conditional variance covariance matrix
                 2
## 1 2.4954 2.3447 2.2031
## 2 2.3447 2.4954 2.3447
## 3 2.2031 2.3447 2.4954
     Standard Deviations: 1.5797 1.5797 1.5797
##
#AICC
p<- 9
print(AICC<- -2*logLik(arint.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 274.5067 (df=9)
#Fitting random slope and intercept model with compound symmetric covariance
matrix of error terms
summary(cs.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + length</pre>
visit, random = ~ 1 + visit | id, data=Longform,
correlation=corCompSymm(form = ~ 1 | id)))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
##
     273.0585 301.6048 -125.5293
##
## Random effects:
## Formula: ~1 + visit | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                         Corr
## (Intercept) 1.5089867 (Intr)
## visit
              0.3709369 -0.245
## Residual
              0.2854340
## Correlation Structure: Compound symmetry
## Formula: ~1 | id
## Parameter estimate(s):
           Rho
##
## 0.001279486
## Fixed effects: score ~ gender.rel + age + doctor.rel + length + visit
                   Value Std.Error DF
                                      t-value p-value
## (Intercept) 5.221633 2.5707883 68 2.031141 0.0462
## gender.relF 2.084693 0.5198847 31 4.009914 0.0004
## age
               0.002996 0.0385434 31 0.077737 0.9385
## doctor.relA 0.158465 0.5084550 31 0.311659 0.7574
## length
               -0.010510 0.0047844 68 -2.196814 0.0314
## visit
               -0.046100 0.0714362 68 -0.645338 0.5209
## Correlation:
##
               (Intr) gndr.F age dctr.A length
```

```
## gender.relF 0.033
## age
               -0.979 -0.166
## doctor.relA -0.213 0.086 0.095
               -0.083 0.114 0.014 0.042
## length
               -0.030 -0.005 -0.001 -0.002 -0.043
## visit
##
## Standardized Within-Group Residuals:
           Min
                        Q1
                                   Med
                                                03
## -1.88928894 -0.46957901 0.01260654 0.40373339 1.67698899
##
## Number of Observations: 105
## Number of Groups: 35
#AICC
p<- 11
print(AICC<- -2*logLik(cs.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 275.8972 (df=11)
#Fitting random slope and intercept model with independent covariance matrix
of error terms
summary(ind.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + length</pre>
visit, random = ~ 1 + visit | id, data=Longform))
## Linear mixed-effects model fit by REML
##
     Data: Longform
##
          AIC
                   BIC
                          logLik
##
     271.0585 297.0097 -125.5293
##
## Random effects:
## Formula: ~1 + visit | id
## Structure: General positive-definite, Log-Cholesky parametrization
##
               StdDev
                         Corr
## (Intercept) 1.5090310 (Intr)
## visit
               0.3709390 -0.245
## Residual
               0.2852501
##
## Fixed effects:
                  score ~ gender.rel + age + doctor.rel + length + visit
                   Value Std.Error DF
                                       t-value p-value
## (Intercept) 5.221580 2.5707920 68 2.031117 0.0462
## gender.relF 2.084700 0.5198854 31 4.009921 0.0004
## age
                0.002997 0.0385434 31 0.077755 0.9385
## doctor.relA 0.158469 0.5084557 31 0.311668 0.7574
               -0.010510 0.0047844 68 -2.196820 0.0314
## length
## visit
               -0.046100 0.0714364 68 -0.645336 0.5209
## Correlation:
               (Intr) gndr.F age
                                    dctr.A length
## gender.relF 0.033
               -0.979 -0.166
## age
## doctor.relA -0.213 0.086 0.095
```

```
-0.083 0.114 0.014 0.042
## length
## visit
               -0.030 -0.005 -0.001 -0.002 -0.043
##
## Standardized Within-Group Residuals:
##
           Min
                        Q1
                                   Med
                                                Q3
                                                            Max
## -1.89038162 -0.46995687 0.01261292 0.40384503 1.67790854
## Number of Observations: 105
## Number of Groups: 35
#AICC
p<- 10
print(AICC<- -2*logLik(ind.fitted.model)+2*p*n/(n-p-1))</pre>
## 'log Lik.' 273.3989 (df=10)
#Fitting random intercept-only model with autoregressive covariance matrix of
error terms
summary(arint.fitted.model<- lme(score ~ gender.rel + age + doctor.rel + leng</pre>
th + visit, random = \sim 1 \mid id,
                                 data=Longform, correlation=corAR1(form = ~ 1
|id)))
## Linear mixed-effects model fit by REML
     Data: Longform
##
##
         AIC
                  BIC
                        logLik
##
     272.612 295.9681 -127.306
##
## Random effects:
## Formula: ~1 | id
            (Intercept) Residual
## StdDev: 0.0005791566 1.57967
##
## Correlation Structure: AR(1)
## Formula: ~1 | id
## Parameter estimate(s):
##
         Phi
## 0.9396084
## Fixed effects:
                   score ~ gender.rel + age + doctor.rel + length + visit
                   Value Std.Error DF
                                        t-value p-value
## (Intercept)
               4.686268 2.6627508 68
                                      1.759935 0.0829
## gender.relF
                2.229715 0.5388793 31 4.137689
                                                 0.0002
## age
                0.010122 0.0398445 31
                                       0.254043
                                                 0.8011
## doctor.relA 0.167042 0.5250427 31
                                       0.318150 0.7525
               -0.011298 0.0051494 68 -2.194026 0.0317
## length
## visit
               -0.045594 0.0647041 68 -0.704658 0.4834
## Correlation:
##
               (Intr) gndr.F age
                                    dctr.A length
## gender.relF 0.028
               -0.978 -0.163
## age
## doctor.relA -0.211 0.083 0.095
```

```
## length -0.101 0.138 0.030 0.016
## visit
              -0.043 -0.007 -0.002 -0.001 -0.051
##
## Standardized Within-Group Residuals:
          Min
                        Q1
                                  Med
                                                Q3
                                                           Max
## -2.23484469 -0.65331876 0.02324913 0.52261910 2.69537024
## Number of Observations: 105
## Number of Groups: 35
intervals(arint.fitted.model)
## Approximate 95% confidence intervals
##
## Fixed effects:
##
                     lower
                                  est.
                                              upper
## (Intercept) -0.62716841 4.68626801 9.999704438
## gender.relF 1.13066337 2.22971487 3.328766360
              -0.07114118 0.01012220 0.091385587
## age
## doctor.relA -0.90378924 0.16704250 1.237874234
## length
              -0.02157322 -0.01129783 -0.001022444
## visit
               -0.17470925 -0.04559425 0.083520749
## attr(,"label")
## [1] "Fixed effects:"
##
##
  Random Effects:
   Level: id
##
##
                          lower
                                       est.
## sd((Intercept)) 7.847905e-63 0.0005791566 4.274037e+55
##
##
  Correlation structure:
##
          lower
                      est.
                               upper
## Phi 0.8917995 0.9396084 0.9666649
## attr(,"label")
## [1] "Correlation structure:"
##
##
  Within-group standard error:
##
     lower
               est.
                        upper
## 1.241087 1.579670 2.010622
getVarCov(arint.fitted.model, type="conditional")
## id 101
## Conditional variance covariance matrix
##
          1
                 2
## 1 2.4954 2.3447 2.2031
## 2 2.3447 2.4954 2.3447
## 3 2.2031 2.3447 2.4954
    Standard Deviations: 1.5797 1.5797 1.5797
```

#Using the fitted model to predict the quality of service score that would be given by a 55-year-old male patient on his fourth visit to Dr. A with a 30-mi nute appointment.

#print(predict(arint.fitted.model, data.frame(id=136, gender.rel="M", age=55,
doctor.rel="A", length=30, visit=4), level=0))

#Generalized Estimating Equations with Correlation Matrix 8.10

```
#Uses data from 8.2
DepartmentStore<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT</pre>
410/Exercise Files/Exercise8.2Data.csv", header = TRUE, sep =",")
#Creating Longform dataset
Longform<- melt(DepartmentStore, id.vars=c("id","totalyears", "status"),</pre>
variable.name="bonus.year", value.name="bonus")
#Sorting data by id
Longform<- Longform[order(Longform$id),]</pre>
#Creating variable for time
year<- ifelse(Longform$bonus.year=="bonus18",1.8,</pre>
ifelse(Longform$bonus.year=="bonus19",1.9,2.0))
#Creating reference level
status.rel<- relevel(as.factor(Longform$status), ref="part")</pre>
#Fitting GEE model with unstructured working correlation matrix
summary(un.fitted.model<- geeglm(bonus ~ totalyears + status.rel + year,</pre>
data=Longform, id=id, family=gaussian(link="identity"),
corstr="unstructured"))
##
## Call:
## geeglm(formula = bonus ~ totalyears + status.rel + year, family = gaussian
(link = "identity"),
       data = Longform, id = id, corstr = "unstructured")
##
## Coefficients:
##
                   Estimate Std.err Wald Pr(>|W|)
## (Intercept)
                  -2063.298 309.562 44.425 2.64e-11 ***
                               7.806 61.146 5.33e-15 ***
## totalyears
                     61.041
## status.relfull 41.799
                               51.598 0.656
                                                 0.418
## year
                  1289.096 169.764 57.661 3.12e-14 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '* 0.05 '.' 0.1 ' ' 1
##
## Correlation structure = unstructured
## Estimated Scale Parameters:
##
```

```
##
               Estimate Std.err
## (Intercept)
                  14719
                           2717
##
     Link = identity
##
## Estimated Correlation Parameters:
             Estimate Std.err
## alpha.1:2 0.13954 0.2692
## alpha.1:3 -0.08909 0.2694
## alpha.2:3 0.80758
## Number of clusters:
                         15 Maximum cluster size: 3
QIC(un.fitted.model)
##
          OIC
                    QICu Quasi Lik
                                           CIC
                                                   params
                                                                QICC
## 6.624e+05 6.624e+05 -3.312e+05 5.188e+00 4.000e+00 6.624e+05
#Fitting GEE model with autoregressive working correlation matrix
summary(ar.fitted.model<- geeglm(bonus ~ totalyears + status.rel + year,</pre>
data=Longform, id=id, family=gaussian(link="identity"), corstr="ar1"))
##
## Call:
## geeglm(formula = bonus ~ totalyears + status.rel + year, family = gaussian
(link = "identity"),
       data = Longform, id = id, corstr = "ar1")
##
##
## Coefficients:
##
                  Estimate Std.err Wald Pr(>|W|)
## (Intercept)
                  -2251.24
                             436.52 26.60 2.5e-07 ***
## totalyears
                     59.60
                               8.15 53.41 2.7e-13 ***
## status.relfull
                              48.30 1.05
                     49.47
                                              0.31
                   1394.67 232.29 36.05 1.9e-09 ***
## year
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Correlation structure = ar1
## Estimated Scale Parameters:
##
##
               Estimate Std.err
## (Intercept)
                  14580
                           2523
    Link = identity
##
##
## Estimated Correlation Parameters:
         Estimate Std.err
            0.383
## alpha
                    0.163
## Number of clusters:
                        15 Maximum cluster size: 3
QIC(ar.fitted.model)
##
                  QICu Quasi Lik
                                       CIC
         QIC
                                              params
                                                          QICC
## 6.56e+05 6.56e+05 -3.28e+05 5.89e+00 4.00e+00 6.56e+05
```

```
#Fitting GEE model with compound symmetric working correlation matrix
summary(cs.fitted.model<- geeglm(bonus ~ totalyears + status.rel + year,</pre>
data=Longform, id=id, family=gaussian(link="identity"),
corstr="exchangeable"))
##
## Call:
## geeglm(formula = bonus ~ totalyears + status.rel + year, family = gaussian
(link = "identity"),
##
       data = Longform, id = id, corstr = "exchangeable")
##
## Coefficients:
                  Estimate Std.err Wald Pr(>|W|)
##
## (Intercept)
                  -2246.99
                             434.47 26.75
                                          2.3e-07 ***
## totalyears
                     59.14
                               8.27 51.14 8.6e-13 ***
## status.relfull
                     51.87
                              48.01 1.17
                                              0.28
                   1394.67 232.29 36.05 1.9e-09 ***
## year
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation structure = exchangeable
## Estimated Scale Parameters:
##
               Estimate Std.err
##
                  14576
                           2518
## (Intercept)
     Link = identity
##
##
## Estimated Correlation Parameters:
##
         Estimate Std.err
## alpha
            0.287
                    0.159
## Number of clusters:
                       15 Maximum cluster size: 3
QIC(cs.fitted.model)
##
         OIC
                  QICu Quasi Lik
                                                          OICC
                                       CIC
                                              params
## 6.56e+05 6.56e+05 -3.28e+05 6.05e+00 4.00e+00 6.56e+05
#Fitting GEE model with independent working correlation matrix
summary(ind.fitted.model<- geeglm(bonus ~ totalyears + status.rel + year,</pre>
data=Longform, id=id, family=gaussian(link="identity"),
corstr="independence"))
##
## Call:
## geeglm(formula = bonus ~ totalyears + status.rel + year, family = gaussian
(link = "identity"),
       data = Longform, id = id, corstr = "independence")
##
##
## Coefficients:
##
                  Estimate Std.err Wald Pr(>|W|)
## (Intercept) -2246.99 434.47 26.75 2.3e-07 ***
```

```
## totalvears
                     59.14 8.27 51.14 8.6e-13 ***
## status.relfull
                    51.87
                             48.01 1.17
                                              0.28
                            232.29 36.05 1.9e-09 ***
## year
                  1394.67
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation structure = independence
## Estimated Scale Parameters:
##
##
               Estimate Std.err
                           2518
## (Intercept)
                  14576
## Number of clusters:
                         15 Maximum cluster size: 3
QIC(ind.fitted.model)
##
                 OICu Ouasi Lik
         OIC
                                       CIC
                                              params
                                                          OICC
## 6.56e+05 6.56e+05 -3.28e+05 6.05e+00 4.00e+00 6.56e+05
summary(un.fitted.model<- geeglm(bonus ~ totalyears+status.rel + year,</pre>
data=Longform, id=id, family=gaussian(link="identity"),
corstr="unstructured"))
##
## Call:
## geeglm(formula = bonus ~ totalyears + status.rel + year, family = gaussian
(link = "identity"),
      data = Longform, id = id, corstr = "unstructured")
##
##
##
   Coefficients:
##
                  Estimate Std.err Wald Pr(>|W|)
                            309.56 44.43 2.6e-11 ***
## (Intercept)
                  -2063.30
                              7.81 61.15 5.3e-15 ***
## totalyears
                     61.04
## status.relfull
                    41.80
                             51.60 0.66
                                              0.42
                  1289.10 169.76 57.66 3.1e-14 ***
## year
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation structure = unstructured
## Estimated Scale Parameters:
##
##
               Estimate Std.err
## (Intercept)
                  14719
                          2717
##
    Link = identity
##
## Estimated Correlation Parameters:
             Estimate Std.err
## alpha.1:2
              0.1395
                        0.269
## alpha.1:3
             -0.0891
                        0.269
## alpha.2:3
              0.8076
                        0.151
## Number of clusters: 15 Maximum cluster size: 3
```

```
##Using the fitted model to predict the predicted bonus in 2021 for a full-ti
me employee who has been with the company for 7 years
print(predict(un.fitted.model, data.frame(totalyears=7, status.rel="full",
    year=2.1)))
## 1
## 1113
```

#Generalized Estimating Equations with Correlation Matrix 8.11

```
#Uses data from 8.3
Orthopedic<- read.csv(file="C:/Users/alex0/Desktop/School/Fall 2021/STAT 410/
Exercise Files/Exercise8.3Data.csv", header = TRUE, sep =",")
#creating longform dataset
data1<- melt(Orthopedic[,c("id","gender","age","doctor","length1",</pre>
"length2", "length3")], id.vars=c("id", "gender", "age", "doctor"),
variable.name="length.visit",value.name="length")
data2<- melt(Orthopedic[,c("score1","score2","score3")],</pre>
variable.name="score.visit", value.name="score")
## No id variables; using all as measure variables
Longform<- cbind(data1,data2)</pre>
#sorting data by id
Longform<- Longform[order(Longform$id),]</pre>
#creating variable for time
visit<- ifelse(Longform$score.visit=="score1",1,</pre>
ifelse(Longform$score.visit=="score2",2,3))
#specifying reference levels
gender.rel<- relevel(as.factor(Longform$gender), ref="M")</pre>
doctor.rel<- relevel(as.factor(Longform$doctor), ref="B")</pre>
#fitting GEE model with unstructured working correlation matrix
summary(un.fitted.model<- geeglm(score ~ gender.rel + age + doctor.rel</pre>
+ length + visit, data=Longform, id=id, family=gaussian(link="identity"),
corstr="unstructured"))
##
## Call:
## geeglm(formula = score ~ gender.rel + age + doctor.rel + length +
##
       visit, family = gaussian(link = "identity"), data = Longform,
       id = id, corstr = "unstructured")
##
##
## Coefficients:
               Estimate Std.err Wald Pr(>|W|)
                                           0.013 *
## (Intercept) 4.68494 1.88506 6.18
## gender.relF 2.12954 0.52062 16.73 4.3e-05 ***
```

```
## age
               0.01032 0.03105 0.11
                                         0.740
## doctor.relA 0.21578 0.48814 0.20
                                          0.658
## length
              -0.01046 0.00558 3.52
                                          0.061 .
## visit
               -0.04797 0.07809 0.38
                                          0.539
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation structure = unstructured
## Estimated Scale Parameters:
##
               Estimate Std.err
##
                  2.18
                         0.525
## (Intercept)
##
    Link = identity
##
## Estimated Correlation Parameters:
             Estimate Std.err
## alpha.1:2
               0.858 0.0522
## alpha.1:3
                0.874
                      0.0429
## alpha.2:3
                0.977
                      0.0476
## Number of clusters:
                        35 Maximum cluster size: 3
QIC(un.fitted.model)
##
         QIC
                 QICu Quasi Lik
                                      CIC
                                             params
                                                          QICC
       266.4
##
                 241.3
                         -114.7
                                      18.5
                                                 6.0
                                                         273.6
#fitting GEE model with autoregressive working correlation matrix
summary(ar.fitted.model<- geeglm(score ~ gender.rel + age + doctor.rel</pre>
+ length + visit, data=Longform, id=id, family=gaussian(link="identity"),
corstr="ar1"))
##
## Call:
## geeglm(formula = score ~ gender.rel + age + doctor.rel + length +
##
      visit, family = gaussian(link = "identity"), data = Longform,
       id = id, corstr = "ar1")
##
##
## Coefficients:
              Estimate Std.err Wald Pr(>|W|)
##
## (Intercept) 4.73590 1.90238 6.20
                                       0.0128 *
## gender.relF 2.21330 0.54843 16.29 5.4e-05 ***
               0.00998 0.03141 0.10
## age
                                        0.7507
## doctor.relA 0.16670 0.49337 0.11
                                        0.7355
                                        0.0098 **
## length
             -0.01238 0.00480 6.67
## visit
              -0.04490 0.07126 0.40
                                         0.5287
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Correlation structure = ar1
## Estimated Scale Parameters:
##
```

```
##
               Estimate Std.err
## (Intercept)
                   2.14
                          0.477
##
    Link = identity
##
## Estimated Correlation Parameters:
         Estimate Std.err
## alpha
            0.926 0.0201
## Number of clusters: 35 Maximum cluster size: 3
QIC(ar.fitted.model)
##
                                       CIC
                                                          QICC
         QIC
                  QICu Quasi Lik
                                              params
##
       262.5
                 236.9
                          -112.4
                                      18.8
                                                 6.0
                                                          266.6
#fitting GEE model with compound symmetric working correlation matrix
summary(cs.fitted.model<- geeglm(score ~ gender.rel + age + doctor.rel</pre>
+ length + visit, data=Longform, id=id, family=gaussian(link="identity"),
corstr="exchangeable"))
##
## Call:
## geeglm(formula = score ~ gender.rel + age + doctor.rel + length +
       visit, family = gaussian(link = "identity"), data = Longform,
       id = id, corstr = "exchangeable")
##
##
## Coefficients:
               Estimate Std.err Wald Pr(>|W|)
##
## (Intercept) 4.46350 1.90511 5.49
                                          0.019 *
## gender.relF 2.15436 0.53112 16.45
                                          5e-05 ***
## age
                0.01480 0.03135 0.22
                                          0.637
## doctor.relA 0.22814 0.48751 0.22
                                          0.640
## length
              -0.01398 0.00624 5.02
                                          0.025 *
## visit
               -0.04387 0.07187 0.37
                                          0.542
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Correlation structure = exchangeable
## Estimated Scale Parameters:
##
##
               Estimate Std.err
## (Intercept)
                          0.473
                   2.11
##
     Link = identity
##
## Estimated Correlation Parameters:
         Estimate Std.err
##
## alpha
            0.898 0.0253
## Number of clusters:
                         35 Maximum cluster size: 3
QIC(cs.fitted.model)
```

```
##
         OIC
                  OICu Ouasi Lik
                                       CIC
                                              params
                                                          OICC
##
       258.1
                 233.1
                          -110.6
                                      18.5
                                                 6.0
                                                         262.2
#fitting GEE model with independent working correlation matrix
summary(ind.fitted.model<- geeglm(score ~ gender.rel + age + doctor.rel</pre>
+ length + visit, data=Longform, id=id, family=gaussian(link="identity"),
corstr="independence"))
##
## Call:
## geeglm(formula = score ~ gender.rel + age + doctor.rel + length +
       visit, family = gaussian(link = "identity"), data = Longform,
##
       id = id, corstr = "independence")
##
##
##
   Coefficients:
##
               Estimate Std.err Wald Pr(>|W|)
## (Intercept) 7.92624 1.39542 32.26 1.3e-08 ***
## gender.relF 1.04380 0.44479 5.51
                                          0.019 *
                                          0.736
## age
                0.00670 0.01984 0.11
## doctor.relA 0.07969 0.31080 0.07
                                          0.798
               -0.09331 0.01397 44.62 2.4e-11 ***
## length
## visit
                0.00713 0.11386 0.00
                                          0.950
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Correlation structure = independence
## Estimated Scale Parameters:
##
##
               Estimate Std.err
## (Intercept)
                          0.189
                   1.28
## Number of clusters:
                         35 Maximum cluster size: 3
QIC(ind.fitted.model)
##
         QIC
                  QICu Quasi Lik
                                       CIC
                                                          QICC
                                              params
##
      153.30
                146.13
                          -67.07
                                      9.58
                                                6.00
                                                        156.30
#fitting GEE model with independent working correlation matrix
summary(un.fitted.model<- geeglm(score ~ gender.rel + age + doctor.rel</pre>
+ length + visit, data=Longform, id=id, family=gaussian(link="identity"),
corstr="unstructured"))
##
## Call:
## geeglm(formula = score ~ gender.rel + age + doctor.rel + length +
       visit, family = gaussian(link = "identity"), data = Longform,
       id = id, corstr = "unstructured")
##
##
## Coefficients:
##
               Estimate Std.err Wald Pr(>|W|)
## (Intercept) 4.68494 1.88506 6.18 0.013 *
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```
## gender.relF 2.12954 0.52062 16.73 4.3e-05 ***
## age
               0.01032 0.03105 0.11
                                        0.740
## doctor.relA 0.21578 0.48814 0.20
                                        0.658
                                        0.061 .
## length -0.01046 0.00558 3.52
## visit
              -0.04797 0.07809 0.38
                                        0.539
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Correlation structure = unstructured
## Estimated Scale Parameters:
##
##
              Estimate Std.err
## (Intercept)
                  2.18
                         0.525
    Link = identity
##
## Estimated Correlation Parameters:
            Estimate Std.err
## alpha.1:2
               0.858 0.0522
## alpha.1:3
               0.874 0.0429
## alpha.2:3
               0.977 0.0476
## Number of clusters: 35 Maximum cluster size: 3
#Using the fitted model to predict the quality of service score that would be
given by a 55-year-old male patient on his fourth visit to Dr. A with a 30-mi
nute appointment.
print(predict(un.fitted.model, data.frame(id=136, gender.rel="M", age=55,
doctor.rel="A",length=30, visit=4)))
##
## 4.96
```