## CSC 423 Homework 6

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## Flight Response of Geese.

a)

```
load("rdata/PACGEESE.Rdata")
glm_model <- glm(as.integer(RESPONSE) ~ ALTITUDE + LATERAL, data = PACGEESE)</pre>
summary(glm_model)
##
## Call:
## glm(formula = as.integer(RESPONSE) ~ ALTITUDE + LATERAL, data = PACGEESE)
## Deviance Residuals:
##
      Min
                 1Q
                      Median
                                   30
                                           Max
## -1.0006 -0.2714
                      0.1229
                                        1.3468
                               0.2318
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.840750
                           0.035457 51.915 < 2e-16 ***
## ALTITUDE
                0.026951
                           0.007765
                                      3.471 0.000568 ***
## LATERAL
               -0.028716
                           0.001615 -17.776 < 2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
  (Dispersion parameter for gaussian family taken to be 0.1395234)
##
##
##
      Null deviance: 109.95 on 463 degrees of freedom
## Residual deviance: 64.32 on 461 degrees of freedom
## AIC: 407.91
##
## Number of Fisher Scoring iterations: 2
```

As we can see both the explanatory variable has p-values less than  $\alpha$  which is 0.01 which tells they are realy significant. b) & c)

```
summary(glm(as.integer(RESPONSE) ~ ALTITUDE, data = PACGEESE))
```

```
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                                   38.250
## (Intercept) 1.52962
                           0.03999
                0.02577
                           0.01007
                                              0.0108 *
## ALTITUDE
                                      2.559
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for gaussian family taken to be 0.2346535)
##
##
       Null deviance: 109.95 on 463 degrees of freedom
## Residual deviance: 108.41 on 462 degrees of freedom
## AIC: 648.14
##
## Number of Fisher Scoring iterations: 2
summary(glm(as.integer(RESPONSE) ~ LATERAL, data = PACGEESE))
##
## Call:
## glm(formula = as.integer(RESPONSE) ~ LATERAL, data = PACGEESE)
## Deviance Residuals:
                                    3Q
##
       Min
                 1Q
                      Median
                                            Max
## -0.9241 -0.2932
                      0.1220
                                0.2143
                                         1.3782
##
## Coefficients:
##
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.928725
                           0.025088
                                      76.88
                                               <2e-16 ***
## LATERAL
               -0.028668
                           0.001635 -17.54
                                               <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1428595)
##
       Null deviance: 109.946 on 463 degrees of freedom
##
## Residual deviance: 66.001 on 462 degrees of freedom
## AIC: 417.88
##
## Number of Fisher Scoring iterations: 2
When \alpha = 0.01 flight response of the geese doesn't depends on altitude of the helicopter. But it depends on
Altitude based on our test.
 d)
pred <- predict(glm_model, newdata = data.frame("ALTITUDE" = 6, "LATERAL" = 3))</pre>
\exp(\text{pred})/(1 + \exp(\text{pred}))
##
## 0.8717264
```

## 9.25 Gender Discrimination In Hiring.

a)

```
load("rdata/DISCRIM.Rdata")
glm_model <- glm(HIRE ~ ., data = DISCRIM)</pre>
summary(glm_model)
##
## Call:
## glm(formula = HIRE ~ ., data = DISCRIM)
## Deviance Residuals:
##
        Min
                   1Q
                         Median
                                        3Q
                                                 Max
## -0.49572 -0.24532 -0.00046
                                   0.15100
                                             0.72310
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) -0.84436
                           0.28128 -3.002 0.006179 **
## EDUC
                0.10692
                           0.04230
                                      2.528 0.018482 *
                0.08863
                           0.02069
                                      4.285 0.000256 ***
## EXP
## GENDER
                0.48473
                            0.13814
                                      3.509 0.001802 **
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for gaussian family taken to be 0.1178536)
##
##
       Null deviance: 6.1071 on 27 degrees of freedom
## Residual deviance: 2.8285 on 24 degrees of freedom
## AIC: 25.272
##
## Number of Fisher Scoring iterations: 2
Since, all variables have p-values less than \alpha we can conclude that this model is adequate to make predictions.
```

b)

```
summary(glm(HIRE ~ GENDER, data = DISCRIM))
```

```
##
## Call:
## glm(formula = HIRE ~ GENDER, data = DISCRIM)
##
## Deviance Residuals:
##
                 1Q
                      Median
                                    ЗQ
                                            Max
## -0.4615 -0.2654 -0.2000
                                         0.8000
                               0.5385
##
## Coefficients:
##
               Estimate Std. Error t value Pr(>|t|)
                 0.2000
## (Intercept)
                            0.1202
                                      1.664
                                               0.108
## GENDER
                 0.2615
                            0.1763
                                      1.483
                                               0.150
##
```

```
## (Dispersion parameter for gaussian family taken to be 0.216568)
##
## Null deviance: 6.1071 on 27 degrees of freedom
## Residual deviance: 5.6308 on 26 degrees of freedom
## AIC: 40.55
##
## Number of Fisher Scoring iterations: 2
```

If yo loos at the p-values which is 0.15 which is more than alpha we can conclude that it is not an important indicator for predictions.

c)

```
ex = data.frame("EDUC" = 4,"EXP" =0,"GENDER" = 1)
preds = predict(glm_model, newdata = ex, type = "link", se.fit = TRUE)
critval = 1.96
logit.point = preds$fit
logit.lower = preds$fit - (critval * preds$se.fit)
logit.upper = preds$fit + (critval * preds$se.fit)
logit.point; logit.lower; logit.upper
```

```
## 1
## 0.06804796
## 1
## -0.1806073
## 1
## 0.3167032
```

These results can be interpreted as, 95 out of 100 samples with the following observations log odds lies between these values.