Brauchle HW4

```
a_model <- glm(Direction ~ Lag1 + Lag2, data = Weekly, family = "binomial")
summary(a_model)</pre>
```

```
##
## Call:
## glm(formula = Direction ~ Lag1 + Lag2, family = "binomial", data = Weekly)
## Deviance Residuals:
##
     Min
              1Q Median
                              3Q
                                     Max
## -1.623 -1.261
                  1.001
                           1.083
                                   1.506
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) 0.22122 0.06147
                                    3.599 0.000319 ***
## Lag1
              -0.03872
                          0.02622 -1.477 0.139672
## Lag2
               0.06025
                          0.02655
                                   2.270 0.023232 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 1496.2 on 1088 degrees of freedom
##
## Residual deviance: 1488.2 on 1086 degrees of freedom
## AIC: 1494.2
##
## Number of Fisher Scoring iterations: 4
```

a. The model including the full dataset only had Lag2 as a significant predictor of Direction (z = 2.27, p < 0.05).

```
weekly_2 <- Weekly[-1,]
b_model <- glm(Direction ~ Lag1 + Lag2, data = weekly_2, family = "binomial")
summary(b_model)</pre>
```

2/21/2019 Brauchle_HW4

```
##
## Call:
### glm(formula = Direction ~ Lag1 + Lag2, family = "binomial", data = weekly_2)
##
## Deviance Residuals:
##
      Min
                1Q Median
                                   3Q
                                          Max
                     0.9999
##
  -1.6258 -1.2617
                              1.0819
                                       1.5071
##
## Coefficients:
##
               Estimate Std. Error z value Pr(>|z|)
                                    3.630 0.000283 ***
## (Intercept) 0.22324
                          0.06150
## Lag1
               -0.03843
                          0.02622 -1.466 0.142683
## Lag2
                0.06085
                          0.02656
                                    2.291 0.021971 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 1494.6 on 1087
                                      degrees of freedom
##
## Residual deviance: 1486.5 on 1085 degrees of freedom
## AIC: 1492.5
##
## Number of Fisher Scoring iterations: 4
```

b. Removing the first observation does not change the model very significantly, and Lag2 remains as the only significant predictor of Direction (z = 2.29, p < 0.05).

```
#c
glm.probs = predict(b_model, weekly[1,], type="response")
if (glm.probs > 0.5) {
   pred_dir = "Up"
}else {
   pred_dir = "Down"
}
pred_dir
```

```
## [1] "Up"
```

```
pred_dir == Weekly[1,]$Direction
```

```
## [1] FALSE
```

c. The first observation was predicted as "Up." However, this observation was not classified correctly, as the true direction of the first observation is "Down."

2/21/2019 Brauchle_HW4

```
#d
errors = vector()
for (i in (1:nrow(weekly))) {
  loop_model <- glm(Direction ~ Lag1 + Lag2, data = Weekly[-i,], family = "binomial")</pre>
  glm.probs = predict(loop_model, weekly[i,], type="response")
  if (glm.probs > 0.5) {
    pred_dir = "Up"
  } else {
    pred_dir = "Down"
  }
  if (pred_dir != Weekly[i,]$Direction){
    errors[i] = 1
  } else {
    errors[i] = 0
  }
}
errors
```

Brauchle HW4

```
##
             ##
           [35] 1 0 0 0 1 0 1 0 0 1 0 1 1 1 0 1 0 0 0 1 0 0 1 1 0 0 0 0 1 0 1 1 0 0
##
          [69] 1 0 1 1 0 0 0 1 0 1 1 0 0 1 1 0 1 1 0 0 1 0 0 1 1 1 0 0 0 0 0 1 0 1
##
        [103] 1 0 0 1 0 1 0 0 1 1 0 0 1 0 0 1 0 0 1 1 1 1 1 0 0 0 1 0 1 1 1 0 0 0
##
        ##
        [171] 0 0 1 1 1 0 1 0 1 0 0 0 0 0 0 0 0 1 1 0 1 0 1 0 1 0 0 0 0 1 0
##
        [205] 0 1 0 1 0 1 1 1 1 0 0 1 1 0 1 0 0 1 1 0 0 0 1 1 1 0 1 0 1 0 1 0 0 0 1
        [239] 1 0 1 0 1 0 1 0 1 0 1 1 0 1 0 1 0 0 1 0 0 1 0 0 0 0 0 0 1 0 0 0 1 0 0 1 0
##
        ##
        [307] 0 0 1 0 0 0 0 1 0 1 1 0 0 1 0 1 0 1 1 0 0 0 1 0 1 0 1 1 1 1 0 1 0
##
##
        [341] 0 1 0 0 0 1 0 1 0 1 0 0 0 0 0 1 1 0 0 1 0 0 1 1 0 1 1 1 1 1
##
        [375] 0 0 0 1 0 0 0 0 0 0 1 0 1 1 0 0 1 1 0 0 0 0 0 1 0 0 1 1 1 0 1 0 1 0
##
        [443] 1 1 0 1 1 0 1 0 1 1 0 1 0 1 0 0 0 1 0 0 0 1 1 0 0 0 0 1
##
##
        [477] 0 0 1 0 0 0 1 1 1 0 1 0 0 0 1 0 1 1 1 0 0 0 0 1 1 1 0 1 0 1 0 0 0
##
        ##
##
        [579] 1 1 1 1 0 1 0 0 1 0 0 1 0 1 0 0 1 0 0 1 0 1 1 1 1 1 1 1 0 1 0 1 0 1 0
         \begin{smallmatrix} 613 \end{smallmatrix} ] \hspace{.1cm} 0 \hspace{.1cm} 0 \hspace{.1cm} 1 \hspace{.1cm} 0 \hspace{.1cm} 1 \hspace{.1cm} 0 \hspace{.1cm} 1 \hspace{.1cm} 1 \hspace{.1cm} 1 \hspace{.1cm} 0 \hspace{.1cm} 1 \hspace{.1cm} 1 \hspace{.1cm} 1 \hspace{.1cm} 0 \hspace{.1cm} 1 \hspace{.
##
##
        [647] 1 1 0 1 0 0 0 1 1 1 1 1 1 1 0 1 0 0 1 0 0 0 1 1 0 1 0 1 1 1 1 0 0 0 1
##
        [681] 1 0 0 0 0 0 0 0 0 0 0 1 0 0 0 0 1 0 0 1 0 1 1 0 0 0 0 1 0 1 0 1 0 1
        ##
##
        [749] 1 1 1 1 1 0 1 0 0 0 0 0 0 1 0 1 1 1 0 0 0 1 0 0 1 0 0 0 1 1 1 0 0 0
##
        [783] 1 0 0 1 1 1 0 0 1 0 1 0 1 0 0 1 0 0 1 0 0 0 0 0 0 1 0 1 1 0 0 1 1 0 1
##
        [817] 1 1 0 0 0 0 0 1 1 0 0 1 0 0 1 0 1 0 0 0 1 1 0 1 1 0 1 0 1 0 1 0 0 0 1
##
        [851] 1 1 1 0 1 1 0 0 0 1 0 1 0 1 0 1 0 0 0 0 0 1 0 0 1 1 0 0 1 0 1 1 1
##
        [919] 0 0 0 1 0 0 1 0 0 0 0 1 0 1 1 1 1 0 0 1 1 0 1 1 1 1 0 1 0 1 0 1 0 1 0
##
        [953] 1 0 0 1 1 1 1 1 1 0 1 0 0 0 1 1 1 0 1 1 1 1 1 0 0 0 0 1 1 0 0 0 0 1 0 0
        [987] 1 1 1 0 0 1 1 1 0 1 0 0 0 0 1 0 0 1 0 1 0 0 1 1 1 1 0 1 0 0 1 0 1
## [1021] 0 0 1 1 0 1 1 1 0 1 1 0 0 0 1 0 1 1 1 1 0 0 1 0 0 0 0 0 0 1 0 1
## [1055] 1 0 0 0 0 0 1 1 1 0 0 0 0 1 0 1 1 1 0 0 0 0 0 0 0 0 0 0 0 0 0 0 1 0 0 0 0 0
## [1089] 0
```

```
mean(errors)
```

```
## [1] 0.4499541
```

e. Using the leave-one-out-cross-validation method produced an error estimate of 45%, so only about 55% of observations were classified correctly.