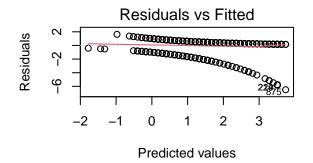
## Lab 21 R Script

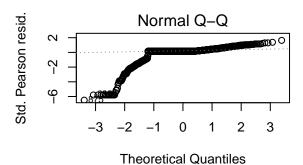
## Alexander Hernandez

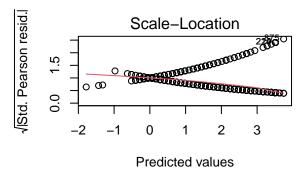
11/15/2022

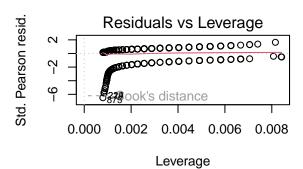
## 1) American Football

```
placekick = read.csv("C:\\repos\\STAT 50001\\Lab 21\\Placekick.csv")
attach(placekick)
model = glm(good ~ distance,
           family = binomial(logit),
                 = placekick)
summary(model)
##
## glm(formula = good ~ distance, family = binomial(logit), data = placekick)
## Deviance Residuals:
                    Median
      Min
           1Q
                                  ЗQ
                                          Max
## -2.7441 0.2425
                    0.2425
                              0.3801
                                       1.6092
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
## (Intercept) 5.812080
                          0.326277
                                    17.81 <2e-16 ***
## distance
              -0.115027
                          0.008339 -13.79 <2e-16 ***
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 1013.43 on 1424 degrees of freedom
## Residual deviance: 775.75 on 1423 degrees of freedom
## AIC: 779.75
## Number of Fisher Scoring iterations: 6
# Simple Logistic Regression Model:
\# good = [1 + exp(-5.812080 + .115027(distance))]^{-1}
par(mfrow=c(2,2))
plot(model)
```









```
##
       data = placekick)
##
##
  Deviance Residuals:
##
       Min
                 1Q
                      Median
                                   3Q
                                           Max
##
   -2.8818
             0.1781
                      0.1781
                               0.4781
                                        1.4183
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
  (Intercept)
               4.51557
                           0.45381
                                     9.950 < 2e-16 ***
               -0.08587
                           0.01100
                                    -7.808 5.83e-15 ***
##
  distance
## PAT
                1.33827
                           0.37951
                                     3.526 0.000421 ***
##
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
##
  (Dispersion parameter for binomial family taken to be 1)
##
       Null deviance: 1013.43 on 1424 degrees of freedom
## Residual deviance: 762.41 on 1422 degrees of freedom
```

```
## AIC: 768.41
##
## Number of Fisher Scoring iterations: 6
# Using all significant regressor variables with 95% confidence, the model is:
# good = [1 + exp( -5.812080 + 0.08587(distance) - 1.33827(PAT) )]^-1
```