## Applied Statistry -MATH 7343

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Flai Dittil

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20.5.8
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(a)

```
> data <- read.table(file="stenosis.txt", header = TRUE)</pre>
   > lr.fit <- glm(data$disease~data$smoke, data = data, family = binomial(link = "logit"))
   > summary(lr.fit)
   Call:
   glm(formula = data$disease ~ data$smoke, family = binomial(link = "logit"),
      data = data)
   Deviance Residuals:
   Min 1Q Median 3Q -1.251 -1.087 -1.087 1.188
                                 1.270
   Coefficients:
             Estimate Std. Error z value Pr(>|z|)
   (Intercept) -0.2157 0.1829 -1.180
                                          0.238
   data$smoke 0.3863
                         0.2762 1.399
                                          0.162
   (Dispersion parameter for binomial family taken to be 1)
      Null deviance: 297.94 on 214 degrees of freedom
   Residual deviance: 295.97 on 213 degrees of freedom
   AIC: 299.97
   Number of Fisher Scoring iterations: 3
                    \ln\left(\frac{\dot{p}}{\dot{n}}\right) = -0.2157 + 0.3863 \,\chi_1
            0.3863 >0 => log obds of developing
                                            aortic stenonis
                   p is higher for individuels who smoke
relative to those who don't . Further, it implies that with smooting the log odds of developing and in creeded by 0.3863. Also, log adds
 increases as "p" increases.
```

(d)

> exp(0.3863)
[1] 1.471526

estimated odds of suffering from antic stensis for individually who snoke relative to those who do not is  $\hat{OR} = 1.47$ 

(c)

> exp(0.3863 + c(-1, 1) \* 1.96 \* 0.2762)
[1] 0.8563706 2.5285651

95% CI (0.856, 2.529)

It contains 1' > there is no evidence

that probability of decloping

acroic stends is different

depending on smoking stadus.

Since "1" is in the interval,

we can conclude that 1.47

is not different from 1".

(d)

> summary(lr.fit2)

```
Call:
qlm(formula = data$disease ~ data$smoke + data$sex, family = binomial(link = "logit"),
   data = data
Deviance Residuals:
Min 1Q Median 3Q
-1.3630 -1.0555 -0.9783 1.0807
                                     Max
                                   1.3905
Coefficients:
          Estimate Std. Error z value Pr(>|z|)
                                      0.0238 *
                       0.2159 -2.261
(Intercept) -0.4882
                       0.2903 0.670
data$smoke
            0.1946
             0.7199
                       0.2881
                              2.499 0.0125 *
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
   Null deviance: 297.94 on 214 degrees of freedom
Residual deviance: 289.64 on 212 degrees of freedom
AIC: 295.64
Number of Fisher Scoring iterations: 4
> \exp(0.1946)
[1] 1.214825
                   OR = 1.215. This is little lower then unadjusted for gender.
              > \exp(0.1946 + c(-1, 1) * 1.96 * 0.2903)
              [1] 0.687710 2.145962
          95%. CI - (6.688, 2.146)
     It Contains "1". Since "1" is in the interval we can conclude that 1.21 is not different from "1".
```

> lr.fit2 <- glm(data\$disease ~ data\$smoke + data\$sex, data = data, family = binomial(link = "logit"))

In order to determine if the relationship between the presence of artic stenotic and between the presence of artic stenotic and females, and in the logistic repression we need to include in the logistic repression we need to interaction term that is the product hadel an interaction term that is the product hadel an interaction term that is the product.