# **Stat 207 HW6**

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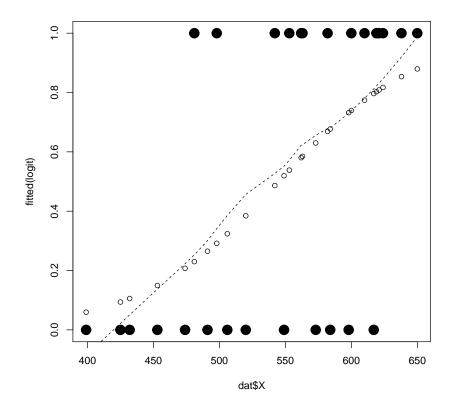


#### 1 14.9

```
dat = read.table("CH14PR09.txt")
  names(dat) = c("Y", "X")
 logit = glm(Y ~ X, data = dat, family = "binomial")
  summary(logit)
##
## Call:
## glm(formula = Y ~ X, family = "binomial", data = dat)
##
## Deviance Residuals:
     Min 1Q Median
##
                               3Q
                                         Max
## -1.7845 -0.8350 0.5065 0.8371
##
## Coefficients:
                Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -10.308925 4.376997 -2.355 0.0185 *
## X
              0.018920
                         0.007877 2.402 0.0163 *
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 37.393 on 26 degrees of freedom
## Residual deviance: 29.242 on 25 degrees of freedom
## AIC: 33.242
##
## Number of Fisher Scoring iterations: 4
```

From the summary, the maximum likelihood estimates of  $\hat{\beta}_0 = -10.308925$ ,  $\hat{\beta}_1 = 0.018920$ ,

$$\hat{\pi} = \frac{exp(\beta_0 + \beta_1 X)}{1 + exp(\beta_0 + \beta_1 X)} = \frac{exp(-10.308925 + 0.018920X)}{1 + exp(-10.308925 + 0.018920X)}$$



The fitted logistic response function appears to be well.

```
(c) exp(0.018920)
## [1] 1.0191
```

 $exp(\beta_1) = 1.0191$ , so that the odds of employee's ability increased by 1.91% with each additional employee's emotional stability.

```
(d) newdat = data.frame(X = 550)
    predict(logit, newdata = newdat, type = "response")

## 1
## 0.5242263
```

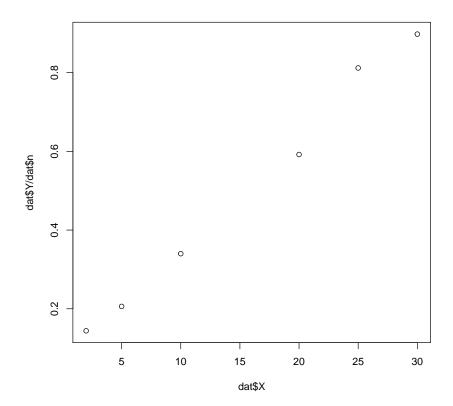
The estimated probability that employees with an emotional stability test score of 550 will be able to perform in a task group is 0.5242263.

```
(e) newpi = 0.7
   pi_2 = log(newpi/(1-newpi))
      (pi_2 - (-10.308925))/0.018920
## [1] 589.6524
```

The emotional stability test score for which 70 percents of the employees with this test score are expected to be able to perform in a task group is 598.6524.

## 2 14.11

```
(a) dat = read.table("CH14PR11.txt")
    names(dat) = c("X", "n", "Y")
    plot(dat$X, dat$Y/dat$n)
```



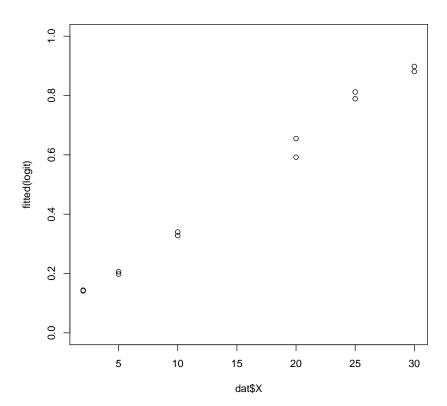
The plot support the analyst's belief that the logistic response function is appropriate.

```
(b) logit = glm(Y/n ~ X, data = dat, family = "binomial")
  ## Warning: non-integer #successes in a binomial glm!
    summary(logit)
   ##
   ## Call:
   ## glm(formula = Y/n ~ X, family = "binomial", data = dat)
  ##
  ## Deviance Residuals:
           1 2
                                3
     0.054601
   ##
   ##
   ## Coefficients:
                Estimate Std. Error z value Pr(>|z|)
   ## (Intercept) -2.0766
                           1.8970 -1.095
                                            0.274
                                   1.273
                            0.1067
                                            0.203
   ## X
                  0.1359
   ##
   ## (Dispersion parameter for binomial family taken to be 1)
   ##
         Null deviance: 2.216343 on 5 degrees of freedom
   ## Residual deviance: 0.024363 on 4 degrees of freedom
   ## AIC: 7.1154
   ##
   ## Number of Fisher Scoring iterations: 4
```

From the summary, the maximum likelihood estimates of  $\hat{\beta}_0 = -2.0766,$   $\hat{\beta}_1 = 0.1359,$ 

$$\hat{\pi} = \frac{exp(\beta_0 + \beta_1 X)}{1 + exp(\beta_0 + \beta_1 X)} = \frac{exp(-2.0766 + 0.1359X)}{1 + exp(-2.0766 + 0.1359X)}$$

```
(c) plot(dat$X, fitted(logit), ylim = c(0, 1))
points(dat$X, dat$Y/dat$n, lwd = 1)
```



The fitted logistic response function appears to be well.

```
(d) exp(0.1359)
## [1] 1.145567
```

 $exp(\beta_1)=1.145567$ , so that the odds of the bottles being returned is increased by 14.5567% with each one deposit level increased.

```
(e) newdat = data.frame(X = 15)
    predict(logit, newdata = newdat, type = "response")

## 1
## 0.4903005
```

The estimated probability that a bottle will be returned when the deposit is 15 cents is 0.4903005.

```
(f) newpi = 0.75

pi_2 = log(newpi/(1-newpi))

(pi_2 - (-2.0766))/0.1359

## [1] 23.36433
```

Estimate the amount of deposit for which 75% of the bottles are expected to be returned is 23.36433.

#### 3 14.14

```
dat = read.table("CH14PR14.txt")
  names(dat) = c("Y", "X1", "X2", "X3")
 logit = glm(Y ~ X1 + X2 + X3, data = dat, family = "binomial")
  summary(logit)
##
## Call:
## glm(formula = Y ~ X1 + X2 + X3, family = "binomial", data = dat)
##
## Deviance Residuals:
      Min
##
           1Q Median
                                  3Q
                                          Max
## -1.4037 -0.5637 -0.3352 -0.1542
                                       2.9394
##
## Coefficients:
              Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -1.17716
                          2.98242 -0.395 0.69307
                          0.03038
## X1
               0.07279
                                   2.396 0.01658 *
              -0.09899
## X2
                          0.03348 -2.957 0.00311 **
               0.43397
                          0.52179
                                   0.832 0.40558
## X3
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 134.94 on 158 degrees of freedom
## Residual deviance: 105.09 on 155 degrees of freedom
## AIC: 113.09
##
## Number of Fisher Scoring iterations: 6
```

From the summary, the maximum likelihood estimates  $\hat{\beta}_0 = -1.17716$ ,

```
\hat{\beta}_1 = 0.07279, \ \hat{\beta}_2 = -0.09899, \ \hat{\beta}_4 = 0.43397 \hat{\pi} = \frac{exp(\beta_0 + \beta_1 X1 + \beta_2 X2 + \beta_3 X3)}{1 + exp(\beta_0 + \beta_1 X1 + \beta_2 X2 + \beta_3 X3)} = \frac{exp(-1.17716 + 0.07279X1 - 0.09899X2 + 0.43397X3)}{1 + exp(-1.17716 + 0.07279X1 - 0.09899X2 + 0.43397X3)}
```

```
(b) exp(0.07279)

## [1] 1.075505

exp(-0.09899)

## [1] 0.9057518

exp(0.43397)

## [1] 1.543373
```

- $exp(\beta_1) = 1.075505$ , so that the odds of getting a flu shot is increased by 7.5% with each one age increased.
- $exp(\beta_2) = 0.9057518$ , so that the odds of getting a flu shot is decreased by 9.4% with each one health awareness index increased.
- $exp(\beta_3) = 1.543373$ , so that the odds of getting a flu shot is increased by 54.3% from woman to man.

The estimated probability with X1=55, X2=60 and X3=1 is 0.06422197

#### $4 \quad 14.19$

(a) dat = read.table("CH14PR13.txt")
 names(dat) = c("Y", "X1", "X2")
 logit = glm(Y ~ X1 + X2, data = dat, family = "binomial")
 summary(logit)

```
##
## Call:
## glm(formula = Y ~ X1 + X2, family = "binomial", data = dat)
##
## Deviance Residuals:
      Min 1Q Median
##
                                 3Q
                                         Max
## -1.6189 -0.8949 -0.5880
                             0.9653
                                      2.0846
##
## Coefficients:
##
              Estimate Std. Error z value Pr(>|z|)
## (Intercept) -4.73931
                         2.10195 -2.255 0.0242 *
             0.06773
                         0.02806
                                 2.414 0.0158 *
## X2
              0.59863
                         0.39007
                                  1.535
                                           0.1249
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## (Dispersion parameter for binomial family taken to be 1)
##
##
      Null deviance: 44.987 on 32 degrees of freedom
## Residual deviance: 36.690 on 30 degrees of freedom
## AIC: 42.69
##
## Number of Fisher Scoring iterations: 4
```

$$H_0: \beta_2 = 0$$
  
VS.  $H_1: \beta_2 \neq 0$   
 $z^* = \frac{b2}{s(b_2)} = 0.59863/0.39007 = 1.535$ 

we can reject  $H_0$  if  $|z^*| > Z(1 - 0.05/2) = 1.959964$ ,otherwise reject  $H_1$  so that reject  $H_1$  because  $|z^*| < 1.959964$ ,

therefore, X2 can be dropped from the regression model, and the P-value is  $0.1249\,$ 

```
(c) logLik(logit)
## 'log Lik.' -18.34482 (df=3)
```

```
logitR = glm(Y ~ X1, data = dat, family = "binomial")
logLik(logitR)

## 'log Lik.' -19.65227 (df=2)

qchisq(1-0.05, 3-2)

## [1] 3.841459

pchisq(2.614, 1, lower.tail = FALSE)

## [1] 0.1059243
```

$$\begin{split} H_0: &\beta_2 = 0 \\ \text{VS. } H_1: &\beta_2 \neq 0 \end{split}$$
 The full model:  $\pi = [1 + exp(-(\beta_0 + \beta_1 X 1 + \beta_2 X 2))]^{-1} \\ \text{L}(F) = -18.34482 \end{split}$  The reduced model:  $\pi = [1 + exp(-(\beta_0 + \beta_1 X 1))]^{-1} \\ \text{L}(R) = -19.65227 \\ G^2 = -2(\ln(\text{L}(R) - \ln(\text{L}(F)))) = 2.614 \end{split}$  we can reject  $H_0$  if  $G^2 > \chi^2(1 - 0.05, 3 - 2) = 3.8415, \text{otherwise reject} H_1$  so that reject  $H_1$  because  $G^2 < 3.8415$ ,

therefore, X2 can be dropped from the regression model, and the P-value is 0.1059. And the result is same as the result we get in (b).

```
(d) logLik(logit)
## 'log Lik.' -18.34482 (df=3)

logitF = glm(Y ~ X1 + X2 +I(X1^2) + I(X2^2) + I(X1*X2), data = dat, family = "binomial logLik(logitF)

## 'log Lik.' -17.12634 (df=6)

qchisq(1-0.05, 6-3)

## [1] 7.814728

pchisq(2.436953, 3, lower.tail = FALSE)
```

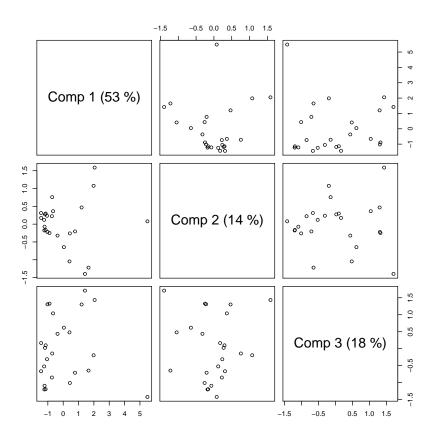
## [1] 0.4867929

```
H_0:\beta_3 = \beta_4 = \beta_5 = 0 VS. H_1:not all \beta_k = 0,k=3,4,5 The full model: \pi = [1 + exp(-(\beta_0 + \beta_1 X1 + \beta_2 X2 + \beta_3 X3 + \beta_4 X4 + \beta_5 X5))]^{-1} L(F) = -17.12634 The reduced model: \pi = [1 + exp(-(\beta_0 + \beta_1 X1 + \beta_2 X2))]^{-1} L(R) = -18.34482 G^2 = -2(\ln(L(R)-\ln(L(F)))) = 2.436953 we can reject H_0 if G^2 > \chi^2(1 - 0.05, 6 - 3) = 7.81,otherwise reject H_1 so that reject H_1 because G^2 < 7.81, therefore, X3,X4,X5 can be dropped from the regression model, and the P-value is 0.4867929.
```

### 5 Problem 5

```
(a) dat = read.table("apartment.txt", header = TRUE)
     require("pls")
   ## Loading required package: pls
   ## Warning: package 'pls' was built under R version 3.1.2
   ##
   ## Attaching package: 'pls'
   ##
   ## The following object is masked from 'package:stats':
   ##
   ##
         loadings
     dat.stan = dat
     for(j in 1:ncol(dat))
       dat.stan[,j] = (dat[,j] - mean(dat[,j]))/sd(dat[,j])
     dat = dat.stan
     fit = plsr(Y ~ X1 + X2 + X3 + X4 + X5, data = dat, 5, validation="CV")
     summary(fit)
   ## Data: X dimension: 25 5
   ## Y dimension: 25 1
   ## Fit method: kernelpls
   ## Number of components considered: 5
   ##
   ## VALIDATION: RMSEP
   ## Cross-validated using 10 random segments.
```

```
## (Intercept) 1 comps 2 comps 3 comps 4 comps 5 comps
## CV
          1.021 0.3139 0.2964 0.2419 0.1853
                                                   0.2259
            1.021 0.3094 0.2901 0.2340 0.1823
## adjCV
                                                   0.2213
##
## TRAINING: % variance explained
## 1 comps 2 comps 3 comps 4 comps 5 comps
## X 52.58 66.94 85.17 91.35
                                    100.00
## Y 92.14
            96.53
                    97.92 98.01
                                    98.05
 loadings(fit)[, 1:3]
         Comp 1
                   Comp 2
## X1 -0.07983743 0.6903977 -0.76418606
## X2 0.59805736 0.1008482 -0.08164449
## X3 0.54164949 -0.5440785 -0.25282151
## X4 0.16890957 -0.7416964 0.59414237
## X5 0.56738864 0.5744813 0.04300711
plot(fit, plottype = "scores", comps = 1:3)
```



```
(b) Radj = c(0, 95.1 , 95.21 , 96.61 , 98.05 , 98.05)/100
    n = 25
    ans = integer(5)
    for (i in 2:6)
    {
        ans[i-1] = (n-i-2)*(Radj[i]-Radj[i-1])/(1-Radj[i])
    }
```

(c)

- 6 Problem 6
- 7 Problem 7
- 8 Problem 8