# **Logistic Regression**

AC

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### **Logistic Regression - Introduction**

Logistic regression is a predictive analysis approach conducted when the dependent variable is dichotomous (binary). It is used to describe data and to explain the relationship between one dependent binary variable and one or more metric independent variables.

#### **Logistic Regression - Basic Equation**

The logistic function can be understood simply as finding the  $\beta$  parameters that best fit:

$$y = 1$$
 if  $\beta_0 + \beta_1 x + \epsilon > 0$ 

y = 0 where  $\epsilon$  is an error distributed by the standard logistic distribution. (If standard normal distribution is used instead, it is a probit regression.)

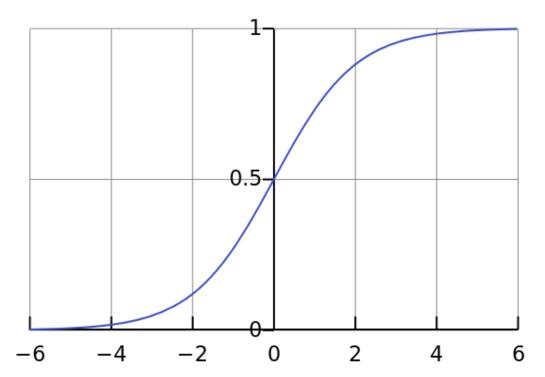


Figure 1. The standard logistic function  $\sigma(t)$  (where  $\sigma(t) \in (0,1)$ )

The logistic function  $\sigma(t)$  is defined as follows:

$$\sigma(t) = \frac{1}{1 + e^{-t}}$$

In our case,  $t = \beta_0 + \beta_1 x$ .

So the logistic function can now be written as:

$$F(x) = \frac{1}{1 + e^{-(\beta_0 + \beta_1 x)}}$$

### **Logistic Regression - Assumptions**

The major assumptions are:

- 1. The outcomes must be discrete i.e.the dependent variable should be binary in nature (eg. presence vs absence).
- 2. There should be no outliers in the data, this can be done by converting the continuous predictors to standardized z scores and remove values below -3.29 or greater than 3.29.
- 3. There should be no high intercorrelations (multicollinearity) among the predictors (Correlation coefficients should be less than 0.9).

#### Sample Example

```
rm(list = ls())
library(ggplot2)
library(aod)
## Warning: package 'aod' was built under R version 3.3.1
library(Rcpp)
dat0 <- data.frame(read.csv("http://www.ats.ucla.edu/stat/data/binary.csv"))</pre>
head(dat0)
##
    admit gre gpa rank
## 1
        0 380 3.61
## 2
        1 660 3.67
                      3
## 3
        1 800 4.00
## 4
        1 640 3.19
## 5
        0 520 2.93
                      4
        1 760 3.00
## 6
summary(dat0)
##
        admit
                                                         rank
                         gre
                                         gpa
## Min.
          :0.0000
                    Min.
                          :220.0
                                    Min.
                                          :2.260
                                                    Min.
                                                           :1.000
## 1st Qu.:0.0000
                    1st Qu.:520.0
                                    1st Qu.:3.130
                                                    1st Qu.:2.000
## Median :0.0000
                    Median :580.0
                                    Median :3.395
                                                    Median :2.000
```

```
## Mean :0.3175
                                   Mean :3.390
                    Mean
                           :587.7
                                                   Mean
                                                          :2.485
## 3rd Qu.:1.0000
                    3rd Qu.:660.0
                                   3rd Qu.:3.670
                                                   3rd Qu.:3.000
## Max.
          :1.0000
                    Max.
                           :800.0
                                   Max.
                                          :4.000
                                                   Max.
                                                          :4.000
sapply(dat0, sd)
##
        admit
                                            rank
                      gre
                                  gpa
##
    0.4660867 115.5165364
                            0.3805668
                                       0.9444602
#table(dat0$admit,dat0$rank)
#For categorical data, looking at contingency table
xtabs(~ admit + rank, data = dat0)
##
       rank
## admit 1 2 3 4
##
      0 28 97 93 55
##
      1 33 54 28 12
#Using the Logit model
dat0$rank <- as.factor(dat0$rank)</pre>
logit <- glm(admit ~ gre + gpa + rank, data = dat0, family = "binomial")</pre>
summary(logit)
##
## Call:
## glm(formula = admit ~ gre + gpa + rank, family = "binomial",
      data = dat0
##
## Deviance Residuals:
##
      Min
                10
                     Median
                                  30
                                         Max
## -1.6268 -0.8662 -0.6388
                              1.1490
                                      2.0790
##
## Coefficients:
               Estimate Std. Error z value Pr(>|z|)
##
## (Intercept) -3.989979 1.139951 -3.500 0.000465 ***
               0.002264
## gre
                          0.001094
                                   2.070 0.038465 *
## gpa
               0.804038
                          0.331819
                                    2.423 0.015388 *
              ## rank2
## rank3
              -1.340204
                          0.345306
                                   -3.881 0.000104 ***
                          0.417832 -3.713 0.000205 ***
## rank4
              -1.551464
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for binomial family taken to be 1)
##
      Null deviance: 499.98 on 399 degrees of freedom
##
## Residual deviance: 458.52 on 394 degrees of freedom
## AIC: 470.52
##
## Number of Fisher Scoring iterations: 4
```

```
#Obtaining confidence intervals using profiled log-liklihood
confint(logit)
## Waiting for profiling to be done...
##
                       2.5 %
                                   97.5 %
## (Intercept) -6.2716202334 -1.792547080
                0.0001375921 0.004435874
## gre
## gpa
                0.1602959439 1.464142727
## rank2
               -1.3008888002 -0.056745722
## rank3
               -2.0276713127 -0.670372346
## rank4
               -2.4000265384 -0.753542605
#Obtaining confidence intervals using standard errors
confint.default(logit)
##
                       2.5 %
                                   97.5 %
## (Intercept) -6.2242418514 -1.755716295
## gre
               0.0001202298 0.004408622
## gpa
                0.1536836760 1.454391423
## rank2
               -1.2957512650 -0.055134591
## rank3
               -2.0169920597 -0.663415773
## rank4
               -2.3703986294 -0.732528724
#Can test the overall effect of rank
wald.test(b = coef(logit), Sigma = vcov(logit), Terms = 4:6)
## Wald test:
## -----
##
## Chi-squared test:
## X2 = 20.9, df = 3, P(> X2) = 0.00011
#Interpreting the odds ratios
exp(coef(logit))
## (Intercept)
                                             rank2
                       gre
                                   gpa
                                                          rank3
                                                                      rank4
##
     0.0185001
                 1.0022670
                             2.2345448
                                         0.5089310
                                                     0.2617923
                                                                  0.2119375
#Interpreting the odds ratio and 95% CI
exp(cbind(OR = coef(logit), confint(logit)))
## Waiting for profiling to be done...
                               2.5 %
                                        97.5 %
##
                      OR
## (Intercept) 0.0185001 0.001889165 0.1665354
## gre
               1.0022670 1.000137602 1.0044457
               2.2345448 1.173858216 4.3238349
## gpa
## rank2
               0.5089310 0.272289674 0.9448343
## rank3
               0.2617923 0.131641717 0.5115181
               0.2119375 0.090715546 0.4706961
## rank4
```

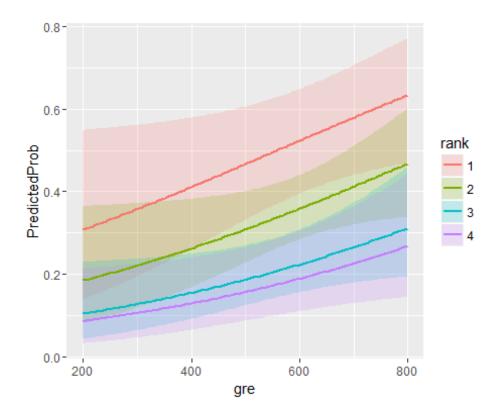
```
#Calculating the predicted probability of admission at each
#value of rank, holding gre and gpa at their means.
dat1 <- with(dat0, data.frame(gre = mean(gre), gpa = mean(gpa),</pre>
             rank = factor(1:4)))
dat1
##
       gre
              gpa rank
## 1 587.7 3.3899
## 2 587.7 3.3899
                     2
## 3 587.7 3.3899
                     3
## 4 587.7 3.3899
                     4
#Prediction
dat1$rankprob <- predict(logit, newdata = dat1, type = "response")</pre>
dat1
##
              gpa rank rankprob
       gre
## 1 587.7 3.3899
                     1 0.5166016
## 2 587.7 3.3899
                     2 0.3522846
## 3 587.7 3.3899
                     3 0.2186120
## 4 587.7 3.3899
                     4 0.1846684
#Preparing data to plot
dat2 <- with(dat0,</pre>
             data.frame(gre = rep(seq(from = 200, to = 800, length.out =
100), 4),
             gpa = mean(gpa), rank = factor(rep(1:4, each = 100))))
dat3 <- cbind(dat2, predict(logit, newdata = dat2, type = "link", se = TRUE))</pre>
dat3 <- within(dat3, {</pre>
               PredictedProb <- plogis(fit)</pre>
               LL <- plogis(fit - (1.96*se.fit))
               UL <- plogis(fit + (1.96*se.fit))
})
head(dat3)
                 gpa rank
                                 fit
                                         se.fit residual.scale
          gre
## 1 200.0000 3.3899
                                                             1 0.5492064
                        1 -0.8114870 0.5147714
## 2 206.0606 3.3899
                       1 -0.7977632 0.5090986
                                                             1 0.5498513
## 3 212.1212 3.3899
                     1 -0.7840394 0.5034491
                                                             1 0.5505074
## 4 218.1818 3.3899
                      1 -0.7703156 0.4978239
                                                             1 0.5511750
## 5 224.2424 3.3899
                        1 -0.7565919 0.4922237
                                                            1 0.5518545
                        1 -0.7428681 0.4866494
## 6 230.3030 3.3899
                                                             1 0.5525464
            LL PredictedProb
## 1 0.1393812
                   0.3075737
```

```
## 2 0.1423880
                   0.3105042
## 3 0.1454429
                   0.3134499
## 4 0.1485460
                   0.3164108
## 5 0.1516973
                   0.3193867
## 6 0.1548966
                   0.3223773
#Testing Model Fit
with(logit, null.deviance - deviance)
## [1] 41.45903
with(logit, df.null - df.residual)
## [1] 5
#Evaluating p-value
with(logit, pchisq(null.deviance - deviance, df.null - df.residual,
lower.tail = FALSE))
## [1] 7.578194e-08
#Log Liklihood
logLik(logit)
## 'log Lik.' -229.2587 (df=6)
```

#### **Plots**

The above result can be graphically shown as follows:

```
ggplot(dat3, aes(x = gre, y = PredictedProb)) +
  geom_ribbon(aes(ymin = LL, ymax = UL, fill = rank), alpha = 0.2) +
  geom_line(aes(colour = rank), size = 1)
```



# References

- 1. R Data Analysis Examples: Logit Regression from http://www.ats.ucla.edu/stat/r/dae/logit.htm
- 2. Statistics Solutions http://www.statisticssolutions.com/what-is-logistic-regression/
- 3. Wikipedia https://en.wikipedia.org/wiki/Logistic\_regression