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Course: RMSC4002
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Code:
+ The script for RMSC4002 Asg2.
+ Dataset: credit.csv
[1] "\nThe script for RMSC4002 Asg2.\nDataset: credit.csv\n"
> d <- read.csv('credit.csv') #read dataset "credit.csv" as d
                            #use the last 5 digits of student id as seed
> set.seed(63766)
> id <- sample(1:690, size=600) #generate random row index for d1
> d1 <- d[id,]
                         #training dataset
> d2 <- d[-id,]
                         #testing dataset
> lreg <- glm(Result~Age+Address+Employ+Bank+House+Save, data=d1,
binomial(link="logit"))
Warning message:
glm.fit: fitted probabilities numerically 0 or 1 occurred
> summary(lreg)
                         #summary
Call:
glm(formula = Result ~ Age + Address + Employ + Bank + House +
  Save, family = binomial(link = "logit"), data = d1)
Deviance Residuals:
                          3Q
         1Q Median
  Min
                                 Max
-3.2089 -0.7704 -0.6258 0.7919 2.0257
Coefficients:
        Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.1542051 0.3258497 -3.542 0.000397 ***
         -0.0071715 0.0092172 -0.778 0.436533
          0.0292526 0.0225961 1.295 0.195463 0.2182379 0.0439829 4.962 6.98e-07 *** 0.3170500 0.0442968 7.157 8.22e-13 ***
Address
Employ
Bank
          -0.0010539 0.0006640 -1.587 0.112462
House
          0.0004290 0.0001184 3.622 0.000292 ***
Save
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
  Null deviance: 823.59 on 599 degrees of freedom
Residual deviance: 606.30 on 593 degrees of freedom
AIC: 620.3
Number of Fisher Scoring iterations: 7
> anova(Ireg, test="Chisq") #to check sig. by chisq test
Analysis of Deviance Table
Model: binomial, link: logit
Response: Result
Terms added sequentially (first to last)
    Df Deviance Resid. Df Resid. Dev Pr(>Chi)
NULL
                  599 823.59
       1 11.139
                    598 812.45 0.0008451 ***
Age
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Address 1 20.685
                     597
                            791.77 5.415e-06 ***
                      596
                            747.23 2.489e-11 ***
Employ 1 44.542
Bank 1 110.798
                     595
                            636.43 < 2.2e-16 ***
                     594
                           635.05 0.2410979
House 1 1.374
Save 1 28.754
                    593
                           606.30 8.219e-08 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> lreg <- glm(Result~Address+Employ+Bank+House+Save, data=d1,
binomial(link="logit"))
Warning message:
glm.fit: fitted probabilities numerically 0 or 1 occurred
> summary(lreg)
                       #summary
Call:
glm(formula = Result ~ Address + Employ + Bank + House + Save,
  family = binomial(link = "logit"), data = d1)
Deviance Residuals:
         1Q Median
                        3Q
-3.2613 -0.7662 -0.6347 0.7903 2.0045
Coefficients:
        Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.3509156 0.2079861 -6.495 8.29e-11 ***
          0.0282123 0.0225509 1.251 0.210916
Address
          0.2077795  0.0417832  4.973  6.60e-07 ***
Employ
          Bank
         House
         0.0004256 0.0001178 3.612 0.000304 ***
Save
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
  Null deviance: 823.59 on 599 degrees of freedom
Residual deviance: 606.91 on 594 degrees of freedom
AIC: 618.91
Number of Fisher Scoring iterations: 7
> anova(Ireg, test="Chisq") #to check sig. by chisq test
Analysis of Deviance Table
Model: binomial, link: logit
Response: Result
Terms added sequentially (first to last)
     Df Deviance Resid. Df Resid. Dev Pr(>Chi)
                 599
NULL
                      823.59
                      598
                            798.29 4.896e-07 ***
Address 1
           25.304
Employ 1 51.061
Bank 1 110.269
                      597
                            747.23 8.954e-13 ***
                     596
                            636.96 < 2.2e-16 ***
House 1 1.324
                     595
                           635.63 0.2499
      1 28.722
                    594
                           606.91 8.354e-08 ***
Save
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> lreg <- glm(Result~Employ+Bank+House+Save, data=d1, binomial(link="logit"))
Warning message:
glm.fit: fitted probabilities numerically 0 or 1 occurred
> summary(lreg)
                       #summary
Call:
glm(formula = Result ~ Employ + Bank + House + Save, family = binomial(link = "logit"),
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data = d1
Deviance Residuals:
          1Q Median
                         3Q
                                Max
-3.1820 -0.7634 -0.6521 0.7793 1.9966
Coefficients:
        Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.2197693 0.1783248 -6.840 7.91e-12 ***
           Employ
          0.3198575  0.0442700  7.225  5.01e-13 ***
Bank
          -0.0011678 0.0006575 -1.776 0.075696 .
House
          0.0004335 0.0001183 3.664 0.000249 ***
Save
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
(Dispersion parameter for binomial family taken to be 1)
  Null deviance: 823.59 on 599 degrees of freedom
Residual deviance: 608.46 on 595 degrees of freedom
AIC: 618.46
Number of Fisher Scoring iterations: 7
> anova(Ireg, test="Chisq") #to check sig. by chisq test
Analysis of Deviance Table
Model: binomial, link: logit
Response: Result
Terms added sequentially (first to last)
    Df Deviance Resid. Df Resid. Dev Pr(>Chi)
NULL
                 599
                        823.59
Employ 1 64.949
                      598
                             758.64 7.685e-16 ***
Bank 1 117.574
                      597
                            641.07 < 2.2e-16 ***
House 1 2.105
                     596
                            638.96 0.1468
Save 1 30.502
                     595
                            608.46 3.336e-08 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
> lreg <- glm(Result~Employ+Bank+Save, data=d1, binomial(link="logit"))
Warning message:
glm.fit: fitted probabilities numerically 0 or 1 occurred
> names(Ireg) #display the items in Ireg
                                    "fitted.values"
                                                                    "R"
[1] "coefficients"
                    "residuals"
                                                     "effects"
[6] "rank"
                  "qr"
                                "family"
                                               "linear.predictors"
                                                                 "deviance"
[11] "aic"
                  "null.deviance"
                                    "iter"
                                                 "weights"
                                                                 "prior.weights"
[16] "df.residual"
                    "df.null"
                                                "converged"
                                                                  "boundary"
                    "call"
                                  "formula"
[21] "model"
                                                                 "data"
                                                  "terms"
[26] "offset"
                   "control"
                                  "method"
                                                   "contrasts"
                                                                   "xlevels"
> pr1 <- (Ireg$fitted.values>0.5) #set pr1=True if fitted > 0.5 or otherwise
> t1 <- table(pr1, d1$Result)
                               #classification table of d1
> p_d1 <- t1[1,1]/sum(t1[1,])
                                 \#precision = TP/(TP+FP)
> r d1 <- t1[1,1]/sum(t1[,1])
                                \#recall = TP/(TP+FN)
> f\bar{1} d1 < -2*p d1*r d1/(p d1+r d1) #F1 score
> m d1 < (t1[1,2]+t1[2,1])/sum(t1)#misclassification rate of d1
> pv d2 <- predict.glm(lreg, newdata=d2) #save predicted values of d2 with lreg.
> pr2 < -(pv_d2 > 0.5)
                                 #set pr2=True if predicted > 0.5 or otherwise
> t2 <- table(pr2, d2$Result)
                                   #classification table of d2
> p d2 <- t2[1,1]/sum(t2[1,])
                                    \#precision = TP/(TP+FP)
                                   \#recall = TP/(TP+FN)
> r d2 <- t2[1,1]/sum(t2[,1])
> f1_d2 <- 2*p_d2*r_d2/(p_d2+r_d2)
                                        #F1 score
> m d2 <- (t2[1,2]+t2[2,1])/sum(t2)
                                       #misclassification rate of d2
> ysort_d1 <- d1$Result[order(lreg$fit, decreasing=T)] #sort y according to lreg$fit
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> n_d1 <- length(ysort_d1)
                                                #get length of ysort
> percl_d1 <- cumsum(ysort_d1)/(1:n_d1)
                                                    #compute cumlative percentage
> plot(percl_d1, type="l", col="blue")
                                                 #plot perc with line type
> abline(h=sum(d1$Result)/n_d1)
                                                  #add the baseline
> yideal_d1 <- c(rep(1, sum(d1$Result)), rep(0, length(d1$Result)-sum(d1$Result))) #the
ideal case
> perc ideal d1 <- cumsum(yideal d1)/(1:n d1)</pre>
                                                       #compute cumulative percentage
of ideal case
> lines(perc_ideal_d1, type="l", col="red")
                                                  #plot the ideal case in red line
> ysort d2 <- d2$Result[order(pv d2, decreasing=T)] #sort y according to predict values
of Ireg as pr2
> n d2 <- length(ysort d2)
                                                #get length of ysort
> percl_d2 <- cumsum(ysort_d2)/(1:n_d2)
                                                    #compute cumlative percentage
> lines(percl_d2, type="l", col="green")
                                                   #plot perc with line type
> abline(h=sum(d2$Result)/n d2)
                                                  #add the baseline
> yideal_d2 <- c(rep(1, sum(d2$Result)), rep(0, length(d2$Result)-sum(d2$Result))) #the
ideal case
> perc_ideal_d2 <- cumsum(yideal_d2)/(1:n_d2)</pre>
                                                      #compute cumulative percentage
of ideal case
> lines(perc_ideal_d2, type="l", col="brown")
                                                     #plot the ideal case in red line
```

