第五次作业参考答案

第一种:

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Homework5

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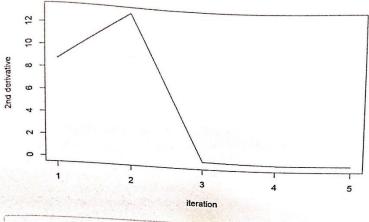
我们将把一个逻辑回归模型与人脸识别算法的测试相关。实验使用识别算法将每个人的第一张图像(称为探针)与剩余的2143张图像之一进行匹配。理想情况下,匹配同一个人的另一个图像(称为目标)。一个成功的匹配产生了yi=1的响应,而与任何其他人的索强度的绝对差异。

第一问、编写您自己的牛顿-拉斐逊算法以适应逻辑回归。初步猜测 β (0) = (β (0) 0 , β (0) 1) > = (0.96, 0)T , 并设置收敛公差 ϵ =10^5.绘制一个图 , 显示每次迭代时导数的值

```
load("C:/Users/Del1/Desktop/facerecognition.RData")
y=as.matrix(data$match)
table(y)

## y
## 0 1
## 282 760
```

```
x=matrix(c(rep(1,1042),data\$eyediff),nr=1042,nc=2)
Beta=matrix(c(0.96,0),nr=2,nc=1)
w=matrix(0,nr=1042,nc=1042)
pi=matrix(0,nr=1042,nc=1)
a=NULL
b1=NULL
b2=NULL
j=0
i=1
while(i<=1042){
pi=1/(1+exp(-x%*%Beta))
w[i,i]=(pi[i]*(1-pi[i]))
i=i+1
loga=t(x)%*%(y-pi)
logb=t(x) % * % w % * % x
Beta=Beta+solve((logb))%*%loga
h=sum(loga^2)
epsilo = 1e-5
while (sqrt(h)>epsilo){
a=c(a,sqrt(h))
b1=c(b1, Beta[1])
b2=c(b2, Beta[2])
pi=1/(1+exp(-x%*%Beta))
loga=t(x) % *% (y-pi)
logb=t(x)%*%w%*%x
Beta=Beta+solve((logb))%*%loga
while(i<=1042){
w[i,i] = (pi[i] * (1-pi[i]))
i=i+1
j=j+1
h=sum(loga^2)
plot(1:j,a,xlab = "iteration",ylab = "2nd derivative",'1')
```

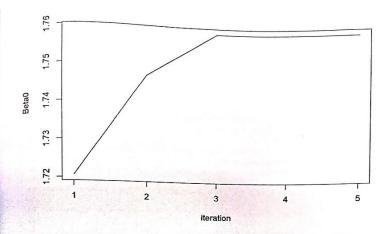


Beta

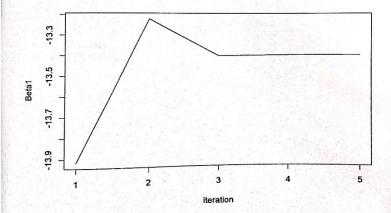
[,1]
[1,] 1.758701
[2,] -13.400040

第二问、绘制一个图,显示β(t)如何移动直到收敛。使用r中的estimate from glm()
函数检查结果(setting family="binomial")。

plot(1:j,b1,xlab = "iteration",ylab = 'Beta0','1')



plot(1:j,b2,xlab = "iteration",ylab = 'Betal','1')



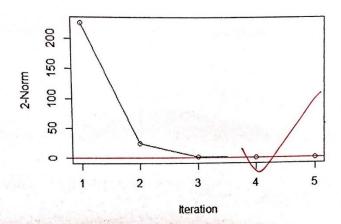
```
fit.full = glm(formula = data\$match \sim data\$eyediff, family = binomial())
   summary(fit.full)
   ##
   ## Call:
   ## glm(formula = data$match ~ data$eyediff, family = binomial())
   ## Deviance Residuals:
   ## Min 1Q Median 3Q Max
## -1.9562 -0.9227 0.6372 0.7630 1.7173
   ##
   ## Coefficients:
  ## Estimate Std. Error z value Pr(>|z|)
## (Intercept) 1.7587 0.1183 14.863 <2e-16 ***
## data$eyediff -13.4000 1.5502 -8.644 <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: 1216.8 on 1041 degrees of freedom
  ##
  ## Residual deviance: 1134.7 on 1040 degrees of freedom
  ## AIC: 1138.7
  ##
  ## Number of Fisher Scoring iterations: 4
  Beta
  ## [1,] 1.758701
  ## [2,] -13.400040
                                                                                  yt1≔1, 0/0
第三问、等高线图
 x=seq(0,10,length=50)
 x=seq(0,10,1ength=50)
y=seq(-40,0,length=50)
y=matrix(nr=50,nc=50)
 f <- function(p,q)(
   betal=c(p,q)
   pi=1/(1+exp(-x%*%betal))
    Log=y* (x%*%beta1)+log(1-Pi)
    sum (Log)
  for (i in 1:50) {
    for (j in 1:50) {
    Z[i,j]=f(X[i],Y[j])
  contour(X,Y,Z)
     9
     20
     9
                                                                                      10
                                                                        8
                                             4
                             2
             0
```

#library(AER)

第二种:

2. Homework 5 加载数据集并准备解释与被解释变量: load("D:/Sta_test.RData") facedata<-data nrow<-dim(facedata)[1]</pre> # extract data for regression y<-as.matrix(facedata\$match)</pre> x<-as.matrix(facedata\$eyediff) x0<-matrix(1,nrow,1) $x \leftarrow cbind(x,x0)$ 模型初始化和参数初值: # model input parameters niter<-100 tol<-1e-5 h2<-matrix(0,niter,1) beta<-matrix(0,2,niter+1) # initial guess for beta beta[,1]<-matrix(c(0.96,0)) 迭代过程: # Newton Raphson for (i in 1:niter){ pi<-1/(1+exp(-x%*%beta[,i])) z<-y-pi h<-t(x)%*%z h2[i]<-(h[1,]^2+h[2,]^2)^0.5 if(h2[i]>tol){ w<-pi%*%t(1-pi) w<-diag(diag(w)) dh<-(-solve(t(x)%*%w%*%x)) beta[,i+1]<-beta[,i]-dh%*%h else{ break } 每一步长的二范数: # plot 2 Norm at each iteration plot(h2[1:i],type="o",main="2-Norm at each iteration",xlab="Iteration", ylab="2-Norm") abline(h=tol,lwd=1,col="red")

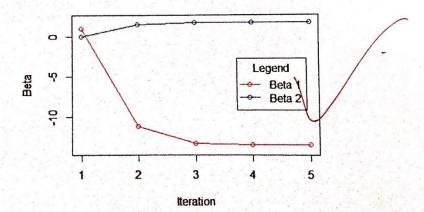
2-Norm at each iteration



每一步长的参数估计值:

plot beta value at each iteration
plot(beta[1,1:i],type="o",main="Beta at each iteration",xlab="Iteration
",ylab="Beta",ylim=c(-14,2),col="red")
par(new=TRUE)
plot(beta[2,1:i],type="o",main="Beta at each iteration",xlab="Iteration
",ylab="Beta",ylim=c(-14,2),col="blue")
legend("right", inset=.05, title="Legend",c("Beta 1","Beta 2"),lty=c(1, 1), pch=c(1, 1), col=c("red", "blue"))

Beta at each iteration



```
参数移动绘图(与 contour plot 一起显示):
# plot beta position change during optimization
plot(beta[1,1:i],beta[2,1:i],type="o",main="Beta locations",xlab="Beta
1",ylab="Beta 2",col="red",xlim=c(-13,1),ylim=c(0,2))
与广义线性回归模型比较回归参数:
# use glm function to check results
glm < -glm(match \sim eyediff, family = binomial(link = "logit"), data = data)
 ##
 ## Call: glm(formula = match ~ eyediff, family = binomial(link = "logi
 t"),
 ##
        data = data)
 ##
 ## Coefficients:
 ## (Intercept)
                     eyediff
 ##
          1.759
                     -13.400
 ## Degrees of Freedom: 1041 Total (i.e. Null); 1040 Residual
 ## Null Deviance:
                         1217
 ## Residual Deviance: 1135 AIC: 1139
  summary(glm)
  ## Call:
  ## glm(formula = match ~ eyediff, family = binomial(link = "logit"),
  ##
         data = data)
  ## Deviance Residuals:
  ##
         Min
                  1Q
                       Median
                                    3Q
  ## -1.9562 -0.9227
                        0.6372
                                 0.7630
                                         1.7173
   ##
   ## Coefficients:
   ##
                 Estimate Std. Error z value Pr(>|z|)
                                              <2e-16 ***
   ## (Intercept) 1.7587
                              0.1183 14.863
   ## eyediff
                 -13.4000
                              1.5502 -8.644
                                              <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: 1216.8 on 1041 degrees of freedom
   ## Residual deviance: 1134.7 on 1040 degrees of freedom
   ## AIC: 1138.7
   ##
   ## Number of Fisher Scoring iterations: 4
```

从上述结果可以看出,广义线性回归模型的参数估计为(1.76,-13.40),与迭代得出的结果十分相近。

绘制 log@ (L(β))的密度图作为底图:

```
# underlying contour plot for Log(L)
beta1<-matrix(seq(-13,1,0.1))
beta2<-matrix(seq(0,2,0.1))
L<-matrix(0,dim(beta1)[1],dim(beta2)[1])
for (k in 1:dim(beta1)[1]){
  for (l in 1:dim(beta2)[1]){
    beta0<-rbind(beta1[k],beta2[1])
    pi<-1/(1+exp(-x%*%beta0))
    L[k,1]<-sum(y*x%*%beta0+log(1-pi))
  }
  }
  par(new=TRUE)
  contour(beta1,beta2,L,nlevels=20,xlim=c(-13,1),ylim=c(0,2))</pre>
```

Beta locations

