

Breast Cancer Diagnosis

Hongjie Liu, Xicheng Xie, Jiajun Tao, Shaohan Chen, Yujia Li

2023-02-27

Outline

- ▶ Background
- ▶ Task 1
- ▶ Task 2
- ▶ Task 3
- ▶ Task 4
- ▶ Discussions
- ▶ Limitations and Future Work
- ▶ Reference
- ▶ Q&A

Background

- ▶ The data is the breast cancer medical data retrieved from “breast-cancer.csv”, which has 569 rows and 32 columns.
- ▶ The first column ID labels individual breast tissue images; The second column Diagnosis identifies if the image is coming from cancer tissue or benign cases (M = malignant, B = benign). There are 357 benign and 212 malignant cases.
- ▶ The other 30 columns correspond to mean, standard deviation and the largest values (points on the tails) of the distributions of the following 10 features computed for the cellnuclei;

Objectives

- ▶ The goal of the exercise is to build a predictive model based on logistic regression to facilitate cancer diagnosis.
- ▶ We will move towards the goal with the steps of task 1, 2, 3 and 4.

Task 1

Objective

Build a logistic model to classify the images into malignant/benign, and write down your likelihood function, its gradient and Hessian matrix.

Task 1 - Build a logistic model

Define the “Diagnosis” variable will be coded as 1 for malignant cases and 0 for benign cases.

Given n i.i.d. observations with p predictors, we consider a logistic regression model

$$P(Y_i = 1 \mid \mathbf{x}_i) = \frac{e^{\mathbf{x}_i^\top \boldsymbol{\beta}}}{1 + e^{\mathbf{x}_i^\top \boldsymbol{\beta}}}, \quad i = 1, \dots, n \quad (1)$$

where $\boldsymbol{\beta} = (\beta_0, \beta_1, \dots, \beta_p)^\top \in \mathbb{R}^{p+1}$ is the parameter vector, $\mathbf{x}_i = (1, X_{i1}, \dots, X_{ip})^\top$ is the vector of predictors in the i -th observation, and $Y_i \in \{0, 1\}$ is the binary response in the i -th observation.

Task 1 - Build a logistic model

Let $\mathbf{y} = (Y_1, Y_2, \dots, Y_n)^\top$ denote the response vector,
 $\mathbf{X} = (\mathbf{x}_1, \mathbf{x}_2, \dots, \mathbf{x}_n)^\top \in \mathbb{R}^{n \times (p+1)}$ denote the design matrix. The
observed likelihood of $\{(Y_1, \mathbf{x}_1), (Y_2, \mathbf{x}_2), \dots, (Y_n, \mathbf{x}_n)\}$ is

$$L(\beta; \mathbf{y}, \mathbf{X}) = \prod_{i=1}^n \left[\left(\frac{e^{\mathbf{x}_i^\top \beta}}{1 + e^{\mathbf{x}_i^\top \beta}} \right)^{Y_i} \left(\frac{1}{1 + e^{\mathbf{x}_i^\top \beta}} \right)^{1-Y_i} \right].$$

Task 1 - Build a logistic model

Maximizing the likelihood is equivalent to maximizing the log-likelihood function:

$$f(\beta; \mathbf{y}, \mathbf{X}) = \sum_{i=1}^n \left[Y_i \mathbf{x}_i^\top \beta - \log \left(1 + e^{\mathbf{x}_i^\top \beta} \right) \right]. \quad (2)$$

The estimates of model parameters are

$$\hat{\beta} = \arg \max_{\beta} f(\beta; \mathbf{y}, \mathbf{X}),$$

and the optimization problem is

$$\max_{\beta} f(\beta; \mathbf{y}, \mathbf{X}). \quad (3)$$

Task 1 - Build a logistic model

Denote $p_i = P(Y_i = 1 \mid \mathbf{x}_i)$ as given in (1) and $\mathbf{p} = (p_1, p_2, \dots, p_n)^\top$. The gradient of f is

$$\begin{aligned}\nabla f(\beta; \mathbf{y}, \mathbf{X}) &= \mathbf{X}^\top (\mathbf{y} - \mathbf{p}) \\ &= \sum_{i=1}^n (Y_i - p_i) \mathbf{x}_i \\ &= \begin{pmatrix} \sum_{i=1}^n (Y_i - p_i) \\ \sum_{i=1}^n (Y_i - p_i) X_{i1} \\ \vdots \\ \sum_{i=1}^n (Y_i - p_i) X_{ip} \end{pmatrix}.\end{aligned}$$

Task 1 - Build a logistic model

Denote $w_i = p_i(1 - p_i) \in (0, 1)$ and $\mathbf{W} = \text{diag}(w_1, \dots, w_n)$. The Hessian matrix of f is given by

$$\begin{aligned}\nabla^2 f(\beta; \mathbf{y}, \mathbf{X}) &= -\mathbf{X}^\top \mathbf{W} \mathbf{X} \\ &= -\sum_{i=1}^n w_i \mathbf{x}_i \mathbf{x}_i^\top \\ &= -\begin{pmatrix} \sum_{i=1}^n w_i & \sum_{i=1}^n w_i X_{i1} & \cdots & \sum_{i=1}^n w_i X_{i1} \\ \sum_{i=1}^n w_i X_{i1} & \sum_{i=1}^n w_i X_{i1}^2 & \cdots & \sum_{i=1}^n w_i X_{i1} X_{ip} \\ \vdots & \vdots & \ddots & \vdots \\ \sum_{i=1}^n w_i X_{ip} & \sum_{i=1}^n w_i X_{ip} X_{i1} & \cdots & \sum_{i=1}^n w_i X_{ip}^2 \end{pmatrix}.\end{aligned}$$

Task 1 - Build a logistic model

Next, we show that the Hessian matrix $\nabla^2 f(\beta; \mathbf{y}, \mathbf{X})$ is a negative-definite matrix if \mathbf{X} has full rank.

Proof. For any $(p + 1)$ -dimensional nonzero vector α , given that \mathbf{X} has full rank, $\mathbf{X}\alpha$ is also a nonzero vector. Since \mathbf{W} is positive-definite, we have

$$\begin{aligned}\alpha^\top \nabla^2 f(\beta; \mathbf{y}, \mathbf{X}) \alpha &= \alpha^\top (-\mathbf{X}^\top \mathbf{W} \mathbf{X}) \alpha \\ &= -(\mathbf{X}\alpha)^\top \mathbf{W} (\mathbf{X}\alpha) \\ &< 0.\end{aligned}$$

Thus, $\nabla^2 f(\beta; \mathbf{y}, \mathbf{X})$ is negative-definite. □

Hence, the optimization problem (3) is a well-defined problem.

Task 2

Objective

Develop a Newton-Raphson algorithm to estimate your model

Task 2 - Newton-Raphson algorithm

The target function f given in task 1:

$$f(\beta; \mathbf{y}, \mathbf{X}) = \sum_{i=1}^n \left[Y_i \mathbf{x}_i^\top \beta - \log \left(1 + e^{\mathbf{x}_i^\top \beta} \right) \right]. \quad (4)$$

Task 2 - Newton-Raphson algorithm

Task 2 - Newton-Raphson algorithm

```
NewtonRaphson <- function(dat, func, start, tol = 1e-8, maxiter) {  
  i <- 0  
  cur <- start  
  stuff <- func(dat, cur)  
  res <- c(0, stuff$f, cur)  
  prevf <- -Inf  
  X <- cbind(rep(1, nrow(dat)), as.matrix(dat[, -1]))  
  y <- dat[, 1]  
  warned <- 0  
  while (abs(stuff$f - prevf) > tol && i < maxiter) {  
    i <- i + 1  
    prevf <- stuff$f  
    prev <- cur  
    d <- -solve(stuff$Hess) %*% stuff$grad  
    cur <- prev + d  
    lambda <- 1  
    maxhalv <- 0  
    while (func(dat, cur)$f < prevf && maxhalv < 50) {
```

Task 2 - Newton-Raphson algorithm

Data preprocessing and data partition.

```
bc_df <- read.csv("breast-cancer.csv")[-c(1, 33)] %>% # remove  
  mutate(diagnosis = ifelse(diagnosis == "M", 1, 0)) # code  
bc_df[, -1] <- scale(bc_df[, -1]) # predictors are standardized  
  
set.seed(1)  
indexTrain <- createDataPartition(y = bc_df$diagnosis, p = 0.8,  
  Training <- bc_df[indexTrain, ]  
Test <- bc_df[-indexTrain, ]  
  
glm(diagnosis ~ ., family = binomial(link = "logit"), data = bc_df)  
  
logisticstuff <- function(dat, betavec) {  
  dat <- as.matrix(dat)  
  n <- nrow(dat)  
  p <- ncol(dat) - 1  
  X <- cbind(rep(1, n), dat[, -1]) # design matrix  
  y <- dat[, 1] # response vector
```


Task 2 - Newton-Raphson algorithm

Task 3

Task 4

Discussions

- ▶ There is much freedom when designing the simulations.
- ▶ In our algorithm, we have 5 parameters. n , p , ratio, c , corr
- ▶ More parameters can be adjusted.

Limitations and Future Work

- ▶ Limitation: We reproduced high-dimensional scenarios, but we still don't know the solution.
- ▶ Future Work: We may adjust other parameters to investigate further.

Reference

1. Li Y, Hong HG, Ahmed SE, Li Y. Weak signals in high-dimensional regression: Detection, estimation and prediction. Appl Stochastic Models Bus Ind. 2018;1–16. <https://doi.org/10.1002/asmb.2340>

Q&A

- ▶ Thanks for listening!
- ▶ Any questions?