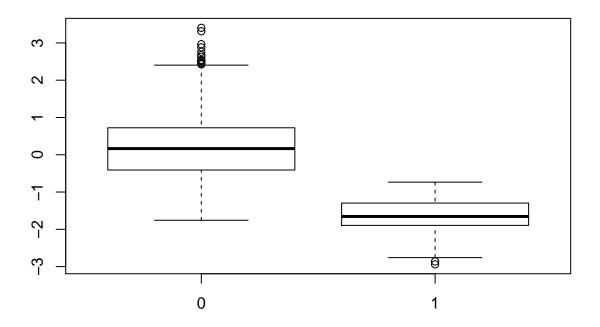
## R Notebook

```
options(scipen=999)
set.seed(1234)
LR2 <- read.table(file="./LR2.csv", header = TRUE, sep = ",")
names(LR2)
## [1] "y" "x"
attach(LR2)
plot(x,y)
          0.8
    9.0
    0.4
    0.2
    0.0
                                                 \infty
                  -2
                                     0
         -3
                           -1
                                              1
                                                       2
                                                                3
                                       Χ
boxplot(x~y)
```



## Assignment 2

## Exercise 1

$$\Pr(Y = 1|X = x) = \Phi(\beta_0 + \beta_1 x)$$
$$\Phi(x) = \int_{-\infty}^{x} \frac{1}{\sqrt{2\pi}} \exp^{-\frac{1}{2}t^2} dt$$

```
phi_pdf <- function(x) {
  cons <- sqrt(2*pi)^-1

# ab <- seq(-10,x,by = 0.001)
  cons*exp(-(x^2/2))
}
#PDF
phi_pdf(0) #0.3989423</pre>
```

## [1] 0.3989423

dnorm(0,0,1)#0.3989423

## [1] 0.3989423

$$\Phi(z) = P(Z \le z), Z \sim \mathcal{N}(0, 1)$$

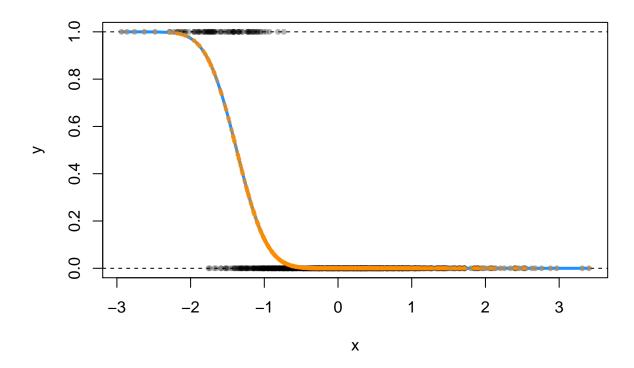
Thus  $\Phi(\beta_0 + \beta_1 x) = P(Z \le z)$ 

Write an R function that computes the maximum likelihood estimate, \mathscr{L} \left( \beta\_0, \beta\_1 \right), along with bootstrapped errors.

```
# objective function
probit_mle_b <- function(x,y,...) {
    opts <- list(...)</pre>
```

```
# probit link
probit <- function(b,x,y) {</pre>
 n <- length(y)
 11 <- 0
  for(i in 1:n) {
    z \leftarrow b[1]+b[2]*x[i]
    z <- pnorm(z, mean=0, sd=1, log.p = FALSE)
    11 \leftarrow 11 + \log(z)*(y[i]==1) + \log(1-z)*(y[i]==0)
 return(-11)
# mle
obj = optim(c(0,0), probit, x=x, y=y)
coef1 <- obj$par[1]</pre>
coef2 <- obj$par[2]</pre>
return_list <- list(</pre>
  model = obj,
  fitted = pnorm(coef1+coef2*x,0,1),
  coefficients = c(coef1,coef2)
)
## Bootstrap
B <- 100
b_boot = matrix(rep(0,2*B),B,2)
n \leftarrow length(y) #n=1000
for (i in 1:B) {
  # indices for the i-th bootstrap subsample
  ind_ = sample(n,n,replace=TRUE)
  # input vector in the subsample
  xb = x[ind_]
  # output vector in the subsample
  yb = y[ind_]
  # compute the maximum likelihood estimates
  obj = optim(c(0,0), probit, x=xb, y=yb)
  b_boot[i,1] = obj$par[1]
  b_{boot[i,2]} = obj par[2]
\texttt{return\_list\$standard\_errors} \leftarrow \texttt{c(sd(b\_boot[,1]),sd(b\_boot[,2]))}
return_list$boots <- b_boot
if(!is.null(opts)) {
```

```
opts <- unlist(opts)</pre>
    return_list$response = ifelse(pnorm(coef1+coef2*opts,0,1)>1/2,1,0)
  return(
    return_list
}
# Apply the probit estimator to LR2
est <- probit_mle_b(LR2$x,LR2$y,x)</pre>
est$coefficients
## [1] -4.242606 -3.086030
#-4.242606 -3.086030
sum(diag(prop.table(table(est$response,y))))
## [1] 0.957
#0.957
train <- LR2[1:800,]
test <- LR2[801:1000,]
est.2 <- probit_mle_b(train$x,train$y,test$x)</pre>
est.2$coefficients
## [1] -4.557228 -3.213275
#-4.557228 -3.213275
sum(diag(prop.table(table(est.2$response,test$y))))
## [1] 0.94
#0.94
glm.est <- glm(y~x,family=binomial(link = "probit"))</pre>
## Warning: glm.fit: fitted probabilities numerically 0 or 1 occurred
plot(x,y, pch=20, col=scales::alpha("black",alpha = 0.3))
abline(h=1, lty=2)
abline(h=0, lty=2)
y0 <- sort(predict(glm.est,list(x),type="response"), decreasing = TRUE)
y1 <- sort(est$fitted,TRUE)
lines(sort(x),y0,lwd=3,col="dodgerblue")
points(sort(x),y1,pch=20,cex=0.7,col=scales::alpha("darkorange",0.5))
```



## Exercise 2

Consider the model. Let X and U be two independent uniformly distributed random variables and let Y be given by the equation

$$Y = I\left(U \le \frac{1}{1 + \exp(-\beta_0 - \beta_1 X - \beta_2 X^3 - \beta_3 \log(X))}\right)$$