Econometrics II TA Session #3

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1 Empirical Application of Binary Model: Titanic Survivors

Brief Background. "Women and children first" is a behavioral norm, which women and children are saved first in a life-threatening situation. This code was made famous by the sinking of the Titanic in 1912. An empirical application investigates characteristics of survivors of Titanic to answer whether crews obeyed the code or not.

Data. We use an open data about Titanic survivors ¹. Although this dataset contains many variables, we use only four variables: survived, age, fare, and sex. We summarize descritons of variables as follows:

- survived: a binary variable taking 1 if a passenger survived.
- age: a continuous variable representing passeger's age.
- fare: a continuous variable representing how much passeger paid.
- sex: a string variable representing passenger's sex.

Using sex, we will make a binary variable, called female, taking 1 if passeger is female. Intead of sex, we use female variable in regression.

```
dt <- read.csv(
  file = "./data/titanic.csv",
  header = TRUE, sep = ",", row.names = NULL, stringsAsFactors = FALSE)

dt$female <- ifelse(dt$sex == "female", 1, 0)
dt <- subset(dt, !is.na(survived)&!is.na(age)&!is.na(fare)&!is.na(female))

dt <- dt[,c("survived", "age", "fare", "female")]
head(dt)</pre>
```

¹data source: http://biostat.mc.vanderbilt.edu/DataSets.

```
## 5 0 25.00 151.5500 1
## 6 1 48.00 26.5500 0
```

Model. In a binary model, a dependent (outcome) variable y_i takes only two values, i.e., $y_i \in \{0,1\}$. A binary variable is sometimes called a *dummy* variable. In this application, the outcome variable is **survived**. Explanatory variables are **female**, **age**, and **fare**. The regression function is

```
\begin{split} &\mathbb{E}[survived|female, age, fare] \\ =& \mathbb{P}[survived = 1|female, age, fare] = G(\beta_0 + \beta_1 female + \beta_2 age + \beta 3 fare). \end{split}
```

The function $G(\cdot)$ is arbitrary function. In practice, we often use following three specifications:

- Linear probability model (LPM): $G(\mathbf{x}_i\beta) = \mathbf{x}_i\beta$.
- Probit model: $G(\mathbf{x}_i\beta) = \Phi(\mathbf{x}_i\beta)$ where $\Phi(\cdot)$ is the standard Gaussian cumulative function.
- Logit model: $G(\mathbf{x}_i\beta) = 1/(1 + \exp(-\mathbf{x}_i\beta))$.

1.1 Linear Probability Model

The linear probability model specifys that G(a) is linear in a, that is,

$$\mathbb{P}[survived = 1 | female, age, fare] = \beta_0 + \beta_1 female + \beta_2 age + \beta 3 fare.$$

This model can be estimated using the OLS method. In R, we can use the OLS method, running lm() function.

```
model <- survived ~ female + age + fare
LPM <- lm(model, data = dt)</pre>
```

However, lm() function does not deal with heteroskedasticity problem. To resolve it, we need to claculate heteroskedasticity-robust standard errors using the White method.

$$\hat{V}(\hat{\beta}) = \left(\frac{1}{n}\sum_i \mathbf{x}_i'\mathbf{x}_i\right)^{-1} \left(\frac{1}{n}\sum_i \hat{u}_i^2\mathbf{x}_i'\mathbf{x}_i\right) \left(\frac{1}{n}\sum_i \mathbf{x}_i'\mathbf{x}_i\right)^{-1}$$

```
# heteroskedasticity-robust standard errors
dt$"(Intercept)" <- 1
X <- as.matrix(dt[,c("(Intercept)", "female", "age", "fare")])
u <- diag(LPM$residuals^2)

XX <- t(X) %*% X
avgXX <- XX * nrow(X)^{-1}
inv_avgXX <- solve(avgXX)

uXX <- t(X) %*% u %*% X</pre>
```

```
avguXX \leftarrow uXX * nrow(X)^{-1}
vcov_b <- (inv_avgXX %*% avguXX %*% inv_avgXX) * nrow(X)^{-1}</pre>
rse b <- sqrt(diag(vcov b))
# homoskedasticity-based standard errors
se b <- sqrt(diag(vcov(LPM)))</pre>
print("The Variance of OLS"); vcov(LPM)
## [1] "The Variance of OLS"
##
                 (Intercept)
                                     female
                                                      age
## (Intercept) 9.754357e-04 -2.891381e-04 -2.333963e-05 -3.329763e-07
## female
               -2.891381e-04 7.136865e-04 2.373259e-06 -1.272800e-06
               -2.333963e-05 2.373259e-06 8.026024e-07 -4.090649e-08
## age
               -3.329763e-07 -1.272800e-06 -4.090649e-08 5.524412e-08
## fare
print("The Robust variance of OLS"); vcov_b
## [1] "The Robust variance of OLS"
##
                 (Intercept)
                                     female
                                                      age
                                                                    fare
## (Intercept) 1.133289e-03 -2.798532e-04 -2.789675e-05 2.813843e-07
## female
               -2.798532e-04 7.903766e-04 3.169092e-06 -2.401923e-06
## age
               -2.789675e-05 3.169092e-06 8.857523e-07 -3.650375e-08
                2.813843e-07 -2.401923e-06 -3.650375e-08 4.071639e-08
## fare
print("The Robust se using White method"); rse_b
## [1] "The Robust se using White method"
   (Intercept)
                      female
                                                   fare
## 0.0336643606 0.0281136372 0.0009411442 0.0002017830
print("The Robust t-value using White method"); coef(LPM)/rse_b
## [1] "The Robust t-value using White method"
## (Intercept)
                    female
                                    age
                                               fare
                 18.229508
                                           7.162302
##
      6.482874
                             -1.884168
  Using the package lmtest and sandwich is the easiest way to calculate heteroskedasticity-
robust standard errors and t-statistics.
library(lmtest) #use function `coeftest`
library(sandwich) #use function `vcovHC`
coeftest(LPM, vcov = vcovHC(LPM, type = "HCO"))[, "Std. Error"]
```

age

fare

female

(Intercept)

##

```
coeftest(LPM, vcov = vcovHC(LPM, type = "HCO"))[, "t value"]
## (Intercept) female age fare
## 6.482874 18.229508 -1.884168 7.162302
```

Finally, we summarize results of linear probability model in table 1. We will discuss interpretation of results and goodness-of-fit of LPM later.

```
# t-stats
t b <- coef(LPM)/se_b
rt b <- coef(LPM)/rse b
# p-value Pr( > |t|)
p b <- pt(abs(t b), df = nrow(X)-ncol(X), lower = FALSE)*2</pre>
rp_b <- pt(abs(rt_b), df = nrow(X)-ncol(X), lower = FALSE)*2
library(stargazer)
stargazer(
  LPM, LPM,
  se = list(se b, rse b), t = list(t b, rt b), p = list(p b, rp b),
  t.auto = FALSE, p.auto = FALSE,
  report = "vcstp", keep.stat = c("n"),
  add.lines = list(
    c("Standard errors", "Homoskedasticity-based", "Heteroskedasticity-robust")),
  title = "Results of Linear Probability Model", label = "LPM",
  type = "latex", header = FALSE, font.size = "small",
  omit.table.layout = "n", table.placement = "h"
)
```

1.2 Probit and Logit Model

Unlike LPM, the probit and logit model must be estimated using the ML method. The probability of observing y_i is

$$p_{\beta}(y_i|\mathbf{x}_i) = \mathbb{P}(y_i = 1|x_i)^{y_i}[1 - \mathbb{P}(y_i = 1|x_i)]^{1-y_i} = G(\mathbf{x}_i\beta)^{y_i}(1 - G(\mathbf{x}_i\beta))^{1-y_i}.$$

Taking logalithm yields

$$\log p_{\beta}(y_i|\mathbf{x}_i) = y_i \log(G(\mathbf{x}_i\beta)) + (1-y_i) \log(1-G(\mathbf{x}_i\beta)).$$

The log-likelihood is

$$M_n(\beta) = \sum_{i=1}^n \log p_{\beta}(y_i|\mathbf{x}_i).$$

The MLE $\hat{\beta}$ holds that the score, which is the first-order derivatives with respect to β , is equal to 0. That is $\nabla_{\beta} M_n(\hat{\beta}) = 0$. For both logit and probit model, the Hessian matrix,

Table 1: Results of Linear Probability Model

	Dependent variable: survived	
	(1)	(2)
female	0.512	0.512
	(0.027)	(0.028)
	t = 19.184	t = 18.230
	p = 0.000	p = 0.000
age	-0.002	-0.002
	(0.001)	(0.001)
	t = -1.979	t = -1.884
	p = 0.049	p = 0.060
fare	0.001	0.001
	(0.0002)	(0.0002)
	t = 6.149	t = 7.162
	p = 0.000	p = 0.000
Constant	0.218	0.218
	(0.031)	(0.034)
	t = 6.988	t = 6.483
	p = 0.000	p = 0.000
Standard errors	Homoskedasticity-based	Heteroskedasticity-robust
Observations	1,045	1,045

 $\nabla^2_{\beta\beta'}M_n(\beta)$, is always negative definite. This implies that log-likelihood function based on both models is grobally concave, and ensures that the MLE maximizes the log-likelihood function. The first-order condition of the probit model is

$$\nabla_{\beta} M_n(\hat{\beta}) = \sum_{i=1}^n \left(y_i - \Phi(\mathbf{x}_i \hat{\beta}) \right) \frac{\phi(\mathbf{x}_i \hat{\beta})}{\Phi(\mathbf{x}_i \hat{\beta}) (1 - \phi(\mathbf{x}_i \hat{\beta}))} = 0.$$

The first-order condition of the logit model is

$$\nabla_{\beta} M_n(\hat{\beta}) = \sum_{i=1}^n \left(y_i - G(\mathbf{x}_i \hat{\beta}) \right) \mathbf{x}_i' = 0.$$

Since it is hard for us to solve this condition analytically, we obtain estimators using numerical procedure.

In R, the function nlm() provides the Newton-Raphson algorithm to minimize the function

². To run this function, we need to define the log-likelihood function (LogLik) beforehand. Moreover, since we need to give initial values in augments, we use coefficients estimated by OLS. Alternatively, we often use glm() function. Using this function, we do not need to define the log-likelihood function and initial values. Since estimates of glm() are approximate to estiamtes of nlm(), we can use this function fairly. In this application, we use nlm function to minimize the log-likelihood function.

```
Y <- dt$survived
female <- dt$female; age <- dt$age; fare <- dt$fare</pre>
# log-likelihood
LnLik <- function(b, model = c("probit", "logit")) {</pre>
  xb \leftarrow b[1] + b[2] *female + b[3] *age + b[4] *fare
  if (model == "probit") {
    L <- pnorm(xb)
  } else {
    L <- 1/(1 + exp(-xb))
  }
  LL i \leftarrow Y * log(L) + (1 - Y) * log(1 - L)
  LL <- -sum(LL i)
  return(LL)
}
#Newton-Raphson
init \leftarrow c(0.218, 0.512, \rightarrow0.002, 0.001)
probit <- nlm(LnLik, init, model = "probit", hessian = TRUE)</pre>
label <- c("(Intercept)", "female", "age", "fare")</pre>
names(probit$estimate) <- label</pre>
colnames(probit$hessian) <- label; rownames(probit$hessian) <- label</pre>
b probit <- probit$estimate</pre>
vcov probit <- solve(probit$hessian); se probit <- sqrt(diag(vcov probit))</pre>
LL probit <- -probit$minimum
#glm function
model <- survived ~ female + age + fare
probit_glm <- glm(model, data = dt, family = binomial("probit"))</pre>
```

²optim() function is an another way to minimize the function. Especially, the function optim(method = "BFGS") provides the Quasi-Newton algorithm which carries on the spirit of Newton method.

```
print("The MLE of probit model using nlm"); b probit
## [1] "The MLE of probit model using nlm"
  (Intercept)
                      female
                                                    fare
                                       age
## -0.813995120 1.435384017 -0.006415761 0.005954843
print("The Variance of probit model using nlm"); vcov probit
## [1] "The Variance of probit model using nlm"
##
                 (Intercept)
                                     female
                                                       age
                                                                    fare
## (Intercept) 1.149118e-02 -3.569149e-03 -2.654781e-04 -1.375309e-05
## female
               -3.569149e-03 8.251773e-03 2.000500e-05 -5.991997e-06
## age
               -2.654781e-04 2.000500e-05 9.630856e-06 -6.874343e-07
               -1.375309e-05 -5.991997e-06 -6.874343e-07 1.103772e-06
## fare
print("The se of probit model using nlm"); se probit
## [1] "The se of probit model using nlm"
## (Intercept)
                    female
                                    age
## 0.107196925 0.090839272 0.003103362 0.001050606
print("The coefficients of probit using glm"); coef(probit glm)
## [1] "The coefficients of probit using glm"
                      female
## (Intercept)
                                                    fare
                                       age
## -0.814075240 1.435384903 -0.006413717 0.005955479
print("The se of probit using glm"); sqrt(diag(vcov(probit_glm)))
## [1] "The se of probit using glm"
## (Intercept)
                    female
                                    age
## 0.108614928 0.090860818 0.003139413 0.001056285
  Using LogLik, we can also estimate logit model by Newton-Raphson algorithm. To com-
pare result, we also use glm() function.
#Newton-Raphson
logit <- nlm(LnLik, init, model = "logit", hessian = TRUE)</pre>
names(logit$estimate) <- label</pre>
colnames(logit$hessian) <- label; rownames(logit$hessian) <- label</pre>
b_logit <- logit$estimate</pre>
vcov_logit <- solve(logit$hessian); se_logit <- sqrt(diag(vcov_logit))</pre>
LL logit <- -logit$minimum
```

```
#glm function
logit glm <- glm(model, data = dt, family = binomial("logit"))</pre>
#result
print("The MLE of logit model"); b logit
## [1] "The MLE of logit model"
## (Intercept)
                    female
                                               fare
                                   age
## -1.33719278 2.35516448 -0.01105760 0.01002878
print("The Variance of logit model"); vcov_logit
## [1] "The Variance of logit model"
##
                 (Intercept)
                                    female
                                                      age
                                                                   fare
## (Intercept)
               0.0351392692 -1.052616e-02 -8.031155e-04 -4.682750e-05
## female
               -0.0105261593 2.411636e-02 3.401375e-05 -7.818252e-06
## age
               -0.0008031155 3.401375e-05 2.939124e-05 -2.170680e-06
               -0.0000468275 -7.818252e-06 -2.170680e-06 3.448283e-06
## fare
print("The se of logit model"); se logit
## [1] "The se of logit model"
## (Intercept)
                    female
                                               fare
                                   age
## 0.187454712 0.155294438 0.005421369 0.001856955
print("The coefficients of logit using glm"); coef(logit glm)
## [1] "The coefficients of logit using glm"
## (Intercept)
                    female
                                               fare
                                   age
## -1.33727469 2.35516632 -0.01105553 0.01002942
print("The se of logit using glm"); sqrt(diag(vcov(logit_glm)))
## [1] "The se of logit using glm"
## (Intercept)
                    female
                                   age
                                               fare
## 0.187350369 0.155280058 0.005424281 0.001847912
```

As a result, table 2 summarizes results of probit model and logit model. t-statistics represents z-value which follows the standard normal distribution. Standard errors are in parentheses. We will discuss interpretation of results and goodness-of-fit later.

```
# z-value
z_probit <- b_probit/se_probit
z_logit <- b_logit/se_logit</pre>
```

```
# Pr(>|z|)
p_probit <- pnorm(abs(z_probit), lower = FALSE)*2
p_logit <- pnorm(abs(z_logit), lower = FALSE)*2

stargazer(
    probit_glm, logit_glm,
    coef = list(b_probit, b_logit), se = list(se_probit, se_logit),
    t = list(z_probit, z_logit), p = list(p_probit, p_logit),
    t.auto = FALSE, p.auto = FALSE,
    report = "vcstp", keep.stat = c("n"),
    add.lines = list(
        c("Log-Likelihood", round(LL_probit, 3), round(LL_logit, 3))),
    title = "Results of Probit and Logit model",
    label = "probit_logit",
    type = "latex", header = FALSE, font.size = "small",
    table.placement = "h", omit.table.layout = "n"
)</pre>
```

Table 2: Results of Probit and Logit model

	Dependent variable:	
	survived	
	probit	logistic
	(1)	(2)
female	1.435	2.355
	(0.091)	(0.155)
	t = 15.801	t = 15.166
	p = 0.000	p = 0.000
age	-0.006	-0.011
	(0.003)	(0.005)
	t = -2.067	t = -2.040
	p = 0.039	p = 0.042
fare	0.006	0.010
	(0.001)	(0.002)
	t = 5.668	t = 5.401
	p = 0.000	p = 0.00000
Constant	-0.814	-1.337
	(0.107)	(0.187)
	t = -7.593	t = -7.133
	p = 0.000	p = 0.000
Log-Likelihood	-530.404	-530.947
Observations Observations	1,045	1,045