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 ~ factor(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))
label <- c("(Intercept)", "factor(female)1", "age", "fare")</pre>
names(rse b) <- label</pre>
# homoskedasticity-based standard errors
se b <- sqrt(diag(vcov(LPM)))</pre>
print("The Variance of OLS"); vcov(LPM)
## [1] "The Variance of OLS"
##
                     (Intercept) factor(female)1
                                                                          fare
                                                            age
## (Intercept)
                    9.754357e-04 -2.891381e-04 -2.333963e-05 -3.329763e-07
## factor(female)1 -2.891381e-04
                                    7.136865e-04 2.373259e-06 -1.272800e-06
## age
                   -2.333963e-05
                                    2.373259e-06 8.026024e-07 -4.090649e-08
## fare
                   -3.329763e-07 -1.272800e-06 -4.090649e-08 5.524412e-08
print("The Robust variance of OLS"); vcov b
## [1] "The Robust variance of OLS"
##
                 (Intercept)
                                     female
                                                      age
## (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
               -2.789675e-05 3.169092e-06 8.857523e-07 -3.650375e-08
## age
## fare
                2.813843e-07 -2.401923e-06 -3.650375e-08 4.071639e-08
print("The Robust se using White method"); rse b
## [1] "The Robust se using White method"
       (Intercept) factor(female)1
##
                                                               fare
                                                age
                      0.0281136372
      0.0336643606
                                       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) factor(female)1
                                                               fare
                                                age
##
                         18.229508
                                          -1.884168
                                                           7.162302
```

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"]
##
       (Intercept) factor(female)1
                                                age
                                                                fare
##
      0.0336643606
                      0.0281136372
                                       0.0009411442
                                                        0.0002017830
coeftest(LPM, vcov = vcovHC(LPM, type = "HCO"))[, "t value"]
##
       (Intercept) factor(female)1
                                                                fare
                                                age
##
          6.482874
                          18.229508
                                                            7.162302
                                          -1.884168
```

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"),
  covariate.labels = c("Female = 1"),
  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

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

Table 1: Results of Linear Probability Model

	Dependent variable: survived		
	(1)	(2)	
Female = 1	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	

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, $\nabla^2_{\beta\beta'} M_n(\beta)$, 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 (LnLik) 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
# 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, -0.002, 0.001)
probit <- nlm(LnLik, init, model = "probit", hessian = TRUE)</pre>
label <- c("(Intercept)", "factor(female)1", "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
```

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.

```
#glm function
model <- survived ~ factor(female) + age + fare</pre>
probit_glm <- glm(model, data = dt, family = binomial("probit"))</pre>
#result
print("The MLE of probit model using nlm"); b probit
## [1] "The MLE of probit model using nlm"
##
       (Intercept) factor(female)1
                                                                fare
                                                 age
                                                         0.005954843
##
      -0.813995120
                       1.435384017
                                       -0.006415761
print("The Variance of probit model using nlm"); vcov probit
## [1] "The Variance of probit model using nlm"
##
                      (Intercept) factor(female)1
                                                                           fare
                                                             age
## (Intercept)
                    1.149118e-02 -3.569149e-03 -2.654781e-04 -1.375309e-05
## factor(female)1 -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) factor(female)1
                                                                fare
                                                 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"
       (Intercept) factor(female)1
                                                                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) factor(female)1
##
                                                                fare
                                                 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>
label <- c("(Intercept)", "factor(female)1", "age", "fare")</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
#qlm 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) factor(female)1
                                                age
                                                                fare
##
       -1.33719278
                        2.35516448
                                                          0.01002878
                                        -0.01105760
print("The Variance of logit model"); vcov_logit
## [1] "The Variance of logit model"
##
                      (Intercept) factor(female)1
                                                                           fare
                                                             age
## (Intercept)
                    0.0351392692 -1.052616e-02 -8.031155e-04 -4.682750e-05
## factor(female)1 -0.0105261593
                                     2.411636e-02 3.401375e-05 -7.818252e-06
## age
                   -0.0008031155
                                     3.401375e-05 2.939124e-05 -2.170680e-06
## fare
                   -0.0000468275
                                    -7.818252e-06 -2.170680e-06 3.448283e-06
print("The se of logit model"); se_logit
## [1] "The se of logit model"
       (Intercept) factor(female)1
##
                                                                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) factor(female)1
                                                age
                                                                fare
##
       -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) factor(female)1
                                                                fare
                                                age
##
       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</pre>
z logit <- b logit/se logit</pre>
\# Pr(>|z|)
p_probit <- pnorm(abs(z_probit), lower = FALSE)*2</pre>
p_logit <- pnorm(abs(z_logit), lower = FALSE)*2</pre>
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"),
  covariate.labels = c("Female = 1"),
  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"
)
```

1.3 Interpretaions

In the linear probability model, interepretations of coefficients are straight-forward. The coefficient β_1 is the change in survival probability given a one-unit increase in continuous variable x. In the case of discrete variable, the coefficient β_1 is the difference in survival probability between two groups. However, when we use the probit or logit model, it is hard for us to interepret results because the partial effect is not constant across other covariates. As an illustration, the partial effect of continuous variable age is

$$\partial_{age} \mathbb{P}[survived = 1 | female, age, fare] = \begin{cases} \beta_2 & \text{if LPM} \\ \phi(\mathbf{x}_i \beta) \beta_2 & \text{if Probit} \ . \\ \frac{\exp(-\mathbf{x}_i \beta)}{(1 + \exp(-\mathbf{x}_i \beta))^2} \beta_2 & \text{if Logit} \end{cases}$$

The partial effect of dummy variable female is

$$\begin{split} \mathbb{P}[survived = 1 | female = 1, age, fare] - \mathbb{P}[survived = 1 | female = 0, age, fare] \\ = \begin{cases} \beta_1 & \text{if LPM} \\ \Phi(\beta_0 + \beta_1 + \beta_2 age + \beta_3 fare) - \Phi(\beta_0 + \beta_1 + \beta_2 age + \beta_3 fare) & \text{if Probit} \\ \Lambda(\beta_0 + \beta_1 + \beta_2 age + \beta_3 fare) - \Lambda(\beta_0 + \beta_1 + \beta_2 age + \beta_3 fare) & \text{if Logit} \end{cases} \end{split}$$
 where $\Lambda(a) = 1/(1 + \exp(-a)).$

Table 2: Results of Probit and Logit model

	Dependent variable: survived		
	probit	logistic	
	(1)	(2)	
Female = 1	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	1,045	1,045	

The first solution is to compute the partial effect at interesting values of \mathbf{x}_i . We often use the sample average of covariates ("average" person) to plugin in the partial effect formula. This is sometimes called *marginal effect at means*. However, since it is unclear what the sample average of dummy variable represents, the marginal effect at means may be hard to explain.

The second solution is to compute the average value of partial effect across the population, that is,

$$\partial_{x_{ij}}\mathbb{P}[y_i=1|\mathbf{x}_i]=\beta_j\mathbb{E}[g(\mathbf{x}_i\beta)],$$

or, in the case of discrete variable,

$$\mathbb{E}[\mathbb{P}[y_i=1|x_{ij}=1,\mathbf{x}_{i,-k}]-\mathbb{P}[y_i=1|x_{ij}=0,\mathbf{x}_{i,-k}]].$$

This is called *average marginal effect* (AME). When we use dummy variables as explanatory variables, we should use this solution.

Standard errors of average marginal effect can be obtained by the Delta method. Let $h_{ij}(\hat{\beta})$ be marginal (partial) effect of the variable x_j for unit i. Then, AME is $h_j(\hat{\beta}) = \mathbb{E}[h_{ij}(\hat{\beta})]$. The Delta method implies that $h_j(\hat{\beta}) \stackrel{d}{\to} N(h_j(\beta), \nabla_{\beta} h_j(\hat{\beta}) V(\beta) (\nabla_{\beta} h_j(\hat{\beta}))')$, where V is variance of $\hat{\beta}$, and

$$\nabla_{\beta}h_{j}(\hat{\beta}) = \begin{pmatrix} \frac{\partial h_{j}(\hat{\beta})}{\partial \beta_{1}} & \dots & \frac{\partial h_{j}(\hat{\beta})}{\partial \beta_{k}} \end{pmatrix}$$

When you use the nlm function to obtain MLE, we need to calculate standard errors manually. The DeltaAME function is a function returing average marginal effect and its standard errors.

```
DeltaAME <- function(b, X, vcov, jbin = NULL, model = c("probit", "logit")) {
  Xb <- numeric(nrow(X))</pre>
  for (i in 1:length(b)) {
     Xb \leftarrow Xb + b[i] * X[,i]
  }
  if (model == "probit") {
    dens <- dnorm(Xb)</pre>
    grad <- -Xb * dens
  } else {
    dens \leftarrow \exp(-Xb)/(1 + \exp(-Xb))^2
    grad <- dens * (-1+2*exp(-Xb)/(1+exp(-Xb)))
  }
  ame <- mean(dens) * b
  if (!is.null(jbin)) {
    for (i in jbin) {
      val1 <- X[,-i] %*% matrix(b[-i], ncol = 1) + b[i]</pre>
      val0 <- X[,-i] %*% matrix(b[-i], ncol = 1)</pre>
      if (model == "probit") {
        amed <- mean(pnorm(val1) - pnorm(val0))</pre>
      } else {
        amed <- mean((1/(1 + \exp(-val1))) - (1/(1 + \exp(-val0))))
      ame[i] <- amed
    }
  }
  e <- NULL
  for (i in 1:length(b)) {
    e <- c(e, rep(mean(X[,i] * grad), length(b)))
  }
  Jacob <- matrix(e, nrow = length(b), ncol = length(b))</pre>
```

```
for (i in 1:nrow(Jacob)) {
  Jacob[i,] <- b[i] * Jacob[i,]</pre>
}
diag(Jacob) <- diag(Jacob) + rep(mean(dens), length(b))</pre>
if (!is.null(jbin)) {
  for (i in jbin) {
    val1 \leftarrow X[,-i] \% matrix(b[-i], ncol = 1) + b[i]
    val0 <- X[,-i] %*% matrix(b[-i], ncol = 1)</pre>
    de <- NULL
    if (model == "probit") {
      for (j in 1:length(b)) {
          if (j != i) {
           dep <- X[,j] * (dnorm(val1) - dnorm(val0))</pre>
           de <- c(de, mean(dep))</pre>
          } else {
           dep <- dnorm(val1)</pre>
           de <- c(de, mean(dep))</pre>
      }
    } else {
      for (j in 1:length(b)) {
          if (j != i) {
           dep <- X[,j] *
             ((\exp(-val1)/(1 + \exp(-val1))^2) - (\exp(-val0)/(1 + \exp(-val0))^2))
           de <- c(de, mean(dep))</pre>
          } else {
           dep \leftarrow exp(-val1)/(1 + exp(-val1))^2
           de <- c(de, mean(dep))</pre>
      }
    }
    Jacob[i,] <- de</pre>
  }
}
label <- names(b)</pre>
colnames(Jacob) <- label; rownames(Jacob) <- label</pre>
vcov_ame <- Jacob %*% vcov %*% t(Jacob)</pre>
se_ame <- sqrt(diag(vcov_ame))</pre>
z ame <- ame/se ame
p ame <- pnorm(abs(z ame), lower = FALSE)*2
```

```
return(list(AME = ame[-1], SE = se ame[-1], zval = z ame[-1], pval = p ame[-1]))
}
X <- as.matrix(dt[,c("(Intercept)", "female", "age", "fare")])</pre>
ame probit <- DeltaAME(b probit, X, vcov probit, jbin = 2, model = "probit")</pre>
ame_logit <- DeltaAME(b_logit, X, vcov_logit, jbin = 2, model = "logit")</pre>
print("AME of probit estimates"); ame probit$AME
## [1] "AME of probit estimates"
## factor(female)1
                                                fare
                                age
       0.508541457
                      -0.001824620
                                        0.001693537
print("AME of logit estimates"); ame logit$AME
## [1] "AME of logit estimates"
## factor(female)1
                                                fare
                                age
                                        0.001653639
##
       0.507384641
                      -0.001823282
print("SE of AME of probit estimates"); ame probit$SE
## [1] "SE of AME of probit estimates"
## factor(female)1
                                                fare
                                age
                                       0.0002874017
      0.0285474135
                      0.0008786651
print("SE of AME of logit estimates"); ame_logit$SE
## [1] "SE of AME of logit estimates"
## factor(female)1
                                                fare
                                age
                                       0.0002948759
##
      0.0287277842
                      0.0008897546
  When we use the glm function, we can use the function margins in the library margins
to obtain the average marginal effect.
library(margins)
summary(margins(probit glm))
     factor
##
                AME
                         SE
                                  Z
                                         р
                                              lower
                                                      upper
##
        age -0.0018 0.0009 -2.0520 0.0402 -0.0036 -0.0001
##
       fare 0.0017 0.0003 5.8398 0.0000 0.0011
                                                     0.0023
    female1 0.5085 0.0286 17.7611 0.0000 0.4524 0.5647
summary(margins(logit glm))
##
     factor
                AME
                         SE
                                  Z
                                         р
                                              lower
                                                      upper
##
        age -0.0018 0.0009 -2.0480 0.0406 -0.0036 -0.0001
```

```
## fare 0.0017 0.0003 5.6338 0.0000 0.0011 0.0022 ## female1 0.5074 0.0287 17.6652 0.0000 0.4511 0.5637
```

Table 3 shows results of linear probability model, probit model, and logit model. In the probit and logit model, coefficients report average marginal effects, t-statistics report z-statistics which follows the standard normal distribution.

All specifications shows that the survival probability of female is about 50% point higher than of male, which is statistically significant. Moreover, the survival probability is decreasing in age, which implies children are more likely to survive. However, the size of coefficient is small. Overall, crews obeyed the code of "women and children first", but the survival probability of children is not largely different from of adult.

```
stargazer(
  LPM, probit_glm, logit_glm,
  coef = list(coef(LPM), ame_probit$AME, ame_logit$AME),
  se = list(rse_b, ame_probit$SE, ame_logit$SE),
  t = list(rt_b, ame_probit$zval, ame_logit$zval),
  p = list(rp_b, ame_probit$pval, ame_logit$pval),
  t.auto = FALSE, p.auto = FALSE,
  omit = c("Constant"), covariate.labels = c("Female = 1"),
  report = "vcstp", keep.stat = c("n"),
  omit.table.layout = "n", table.placement = "t",
  title = "Titanic Survivors: LPM, Probit (AME), and Logit (AME)",
  label = "titanic",
  type = "latex", header = FALSE
)
```

Table 3: Titanic Survivors: LPM, Probit (AME), and Logit (AME)

	Dependent variable: survived			
	OLS	probit	logistic	
	(1)	(2)	(3)	
$\overline{\text{Female}} = 1$	0.512	0.509	0.507	
	(0.028)	(0.029)	(0.029)	
	t = 18.230	t = 17.814	t = 17.662	
	p = 0.000	p = 0.000	p = 0.000	
age	-0.002	-0.002	-0.002	
	(0.001)	(0.001)	(0.001)	
	t = -1.884	t = -2.077	t = -2.049	
	p = 0.060	p = 0.038	p = 0.041	
fare	0.001	0.002	0.002	
	(0.0002)	(0.0003)	(0.0003)	
	t = 7.162	` /	t = 5.608	
	p = 0.000	p = 0.000	p = 0.00000	
Observations	1,045	1,045	1,045	