## Econometrics-HW5

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library(tidyverse)

```
## -- Attaching packages ------ tidyverse 1.3.1 --
## v ggplot2 3.3.5
                     v purrr
                               0.3.4
## v tibble 3.1.6
                     v dplyr
                               1.0.8
## v tidyr
           1.2.0
                     v stringr 1.4.0
## v readr
            2.1.2
                     v forcats 0.5.1
## Warning: package 'tidyr' was built under R version 4.0.5
## Warning: package 'readr' was built under R version 4.0.5
## Warning: package 'dplyr' was built under R version 4.0.5
## -- Conflicts ----- tidyverse_conflicts() --
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                   masks stats::lag()
apple <- read_csv("apple.csv")</pre>
## Rows: 660 Columns: 17
## -- Column specification -------
## Delimiter: ","
## chr (1): state
## dbl (16): id, educ, date, regprc, ecoprc, inseason, hhsize, male, faminc, ag...
## i Use `spec()` to retrieve the full column specification for this data.
## i Specify the column types or set `show_col_types = FALSE` to quiet this message.
head(apple)
## # A tibble: 6 x 17
##
       id educ
                date state regprc ecoprc inseason hhsize male faminc
    <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
                                            <dbl> <dbl> <dbl> <dbl> <dbl> <
## 1 10002
                              1.19
             16 111597 SD
                                    1.19
                                                1
                                                      4
                                                            0
                                                                  45
                                                                        43
## 2 10004
                              0.59
                                                            0
                                                                  65
                                                                        37
             16 121897 KS
                                    0.79
                                                0
                                                      1
## 3 10034
             18 111097 MI
                              0.59
                                                            0
                                                                  65
                                                                        44
                                    0.99
                                                1
                                                      3
## 4 10035
            12 111597 TN
                              0.89
                                    1.09
                                                                  55
                                                                        55
## 5 10039
          15 122997 NY
                              0.89
                                    1.09
                                                0
                                                      1
                                                                  25
                                                                        22
## 6 10041
                              0.59
                                    0.79
            12 112297 WV
                                                1
                                                                        34
## # ... with 6 more variables: reglbs <dbl>, ecolbs <dbl>, numlt5 <dbl>,
## # num5_17 <dbl>, num18_64 <dbl>, numgt64 <dbl>
#create dummy variables
apple$ecobuy <- ifelse(apple$ecolbs>0,1,0)
```

```
attach(apple)
```

#### a. Derive the log-likelihood function.

 $\max l_i = p_1^{regprc_i} p_2^{ecoprc_i} (1 - p_1 - p_2)^{age_i}$  Take log on the above eqn:  $\max \sum_i^N \log l_i = \sum_i^N regprc_i \log p_i + \sum_i^N regprc_i \log p_i$  $ecoprc_i \log p_2 + age_i \log(1 - p_1 - p_2)$ 

#### b. Estimate the parameters of interest via M-estimation

```
source('~/Dropbox/My Mac (Wei's MacBook Air)/Downloads/PhD-Coursework/22Spring/Econometrics/My_solution
source('~/Dropbox/My Mac (Wei's MacBook Air)/Downloads/PhD-Coursework/22Spring/Econometrics/My_solution
 source('~/Dropbox/My Mac (Wei's MacBook Air)/Downloads/PhD-Coursework/22Spring/Econometrics/My_solution
X=cbind(regprc,ecoprc,age)
thetahat=cbind(0.2,0.6,0.2)
out <- qderivfun(ecobuy, X, t(thetahat))</pre>
out1 <- qderivfun2(ecobuy,X,t(thetahat))</pre>
out1
## $H
##
                          [,2]
                                        [,3]
               [,1]
## [1,] -2207533.70 -11158.73
                                  -1304947.8
        -11158.73 -301384.07
## [2,]
                                  -533228.8
## [3,] -1304947.80 -533228.77 -194413608.1
##
## $AOhat
##
               [,1]
                          [,2]
                                        [,3]
## [1,] -3344.74802 -16.90717
                                  -1977.1936
        -16.90717 -456.64253
## [2,]
                                  -807.9224
## [3,] -1977.19364 -807.92238 -294566.0729
c. Make t-test:
ava_theta <- solve(out1$A0hat)%*% out$B0hat %*% solve(out1$A0hat)/length(ecobuy)
ava_theta
##
                [,1]
                              [,2]
## [1,] 2.860107e-05 8.278539e-05 2.523338e-05
## [2,] 8.278539e-05 2.547686e-04 7.605291e-05
## [3,] 2.523338e-05 7.605291e-05 3.348403e-05
se_theta <- diag(ava_theta)^(1/2)
```

```
## [1] 0.005347996 0.015961471 0.005786539
```

Now, make some t-tests on theta:

se\_theta

```
t_1 <- log(thetahat)/se_theta
t_2 <- thetahat/se_theta
t_1
```

```
[,1]
                        [,2]
                                   [,3]
## [1,] -300.9422 -32.00367 -278.1348
```

```
t_2
## [,1] [,2] [,3]
```

No matther what, abs of t test values are way larger than 1.96. So we can react the zero null for all variables.

#### d. Use probit command in R:

## [1,] 37.39718 37.59052 34.56298

```
probitout <- glm(ecobuy~regprc+ecoprc+age,family=binomial(link="probit"))</pre>
probitout
##
## Call: glm(formula = ecobuy ~ regprc + ecoprc + age, family = binomial(link = "probit"))
## Coefficients:
## (Intercept)
                     regprc
                                   ecoprc
                                                    age
       1.26121
                    2.03914
                                 -2.33504
##
                                              -0.00432
##
## Degrees of Freedom: 659 Total (i.e. Null); 656 Residual
## Null Deviance:
                         873.8
## Residual Deviance: 813.5
                                 AIC: 821.5
```

The conclusions will not be changed.

# e. Compute the likelihood ratio test and justify whether reject the null hypothesis.

```
probitout_red <- glm(ecobuy~ecoprc+age,family=binomial(link="probit"))</pre>
probitout_red
## Call: glm(formula = ecobuy ~ ecoprc + age, family = binomial(link = "probit"))
##
## Coefficients:
## (Intercept)
                     ecoprc
                                      age
##
      1.491366
                  -0.911912
                               -0.003958
##
## Degrees of Freedom: 659 Total (i.e. Null); 657 Residual
## Null Deviance:
                        873.8
## Residual Deviance: 843.3
                                 AIC: 849.3
#compute likelihood ratio
library(epiDisplay)
## Loading required package: foreign
## Warning: package 'foreign' was built under R version 4.0.5
## Loading required package: survival
## Loading required package: MASS
## Warning: package 'MASS' was built under R version 4.0.5
## Attaching package: 'MASS'
```

```
## The following object is masked from 'package:dplyr':
##
## select

## Loading required package: nnet

## Warning: package 'nnet' was built under R version 4.0.5

##
## Attaching package: 'epiDisplay'

## The following object is masked from 'package:ggplot2':
##
## alpha

lrtest(probitout,probitout_red)

## Likelihood ratio test for MLE method
## Chi-squared 1 d.f. = 29.71658 , P value = 5.000524e-08
```

Actually from the command we can derive Chi-squared number from lrtest command without replying on dchisq. It's chi-squared value is high, and P-value is way lower than 0.05, therefore, we reject the null hypothesis.