Fundamentals of Statistical Modeling (VT21)

Andrea Discacciati Karolinska Institutet Stockholm, Sweden

Lab 2

Load the dataset and the mlci command

```
. version 14
. use https://raw.githubusercontent.com/anddis/fsm/master/data/lab2.dta, clear
. run https://raw.githubusercontent.com/anddis/fsm/master/do/mlci.do
```

Install the qplot command (you need to be connected to the Internet)

. net sj 16-3 gr42_7

```
package gr42_7 from http://www.stata-journal.com/software/sj16-3
```

```
TITLE

SJ16-3 gr42_7. Update: Quantile plots

DESCRIPTION/AUTHOR(S)

Update: Quantile plots
by Nicholas J. Cox, Durham University,
Department of Geography, Durham, UK

Support: n.j.cox@durham.ac.uk
After installation, type help qplot

INSTALLATION FILES

gr42_7/qplot.ado
gr42_7/qplot.sthlp

(type net install gr42_7)
```

```
. net install gr42_7 checking gr42_7 consistency and verifying not already installed... all files already exist and are up to date.
```

Install the rcsgen command (you need to be connected to the Internet)

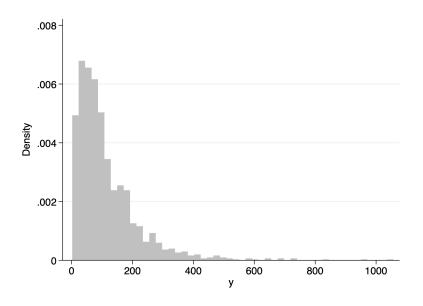
```
. cap net install http://fmwww.bc.edu/RePEc/bocode/r/rcsgen.pkg
```

Exercise 1

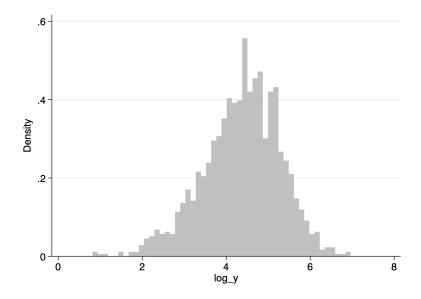
This dataset contains information on the blood concentration of a biomarker (y) in a random sample of 1432 subjects. Take a look at the histogram. What can we say about the distribution of this biomarker?

Plot also the histogram of log(y). How does the distribution of the biomarker after logarithmic transform look like?

```
. hist y, bin(50) name(p1, replace)
(bin=50, start=2.2592716, width=21.079531)
. graph export p1.png, replace
(file p1.png written in PNG format)
```



```
. gen log_y = log(y)
. hist log_y, bin(50) name(p2, replace)
(bin=50, start=.8150425, width=.12294848)
. graph export p2.png, replace
(file p2.png written in PNG format)
```



Exercise 2

We assume that f(y) is gamma (see Lab 1). Estimate the parameters α and β using the gammaden() function. Fix the location parameter g (the third argument of the gammaden() function) to be equal to 0. Constrain α and β to be positive.

Note: the parameters α and β are not interpretable. We can reparametrise the gamma distribution so that one parameter is equal to its mean. This is described in the Extra material for Lab 2.

```
Iteration 0:     log likelihood = -13891.173
Iteration 1:     log likelihood = -8165.6417
Iteration 2:     log likelihood = -8160.8897
Iteration 3:     log likelihood = -8160.8781
Iteration 4:     log likelihood = -8160.8781
```

Maximum likelihood estimation

Log likelihood = -8160.8781 Number of obs = 1,432

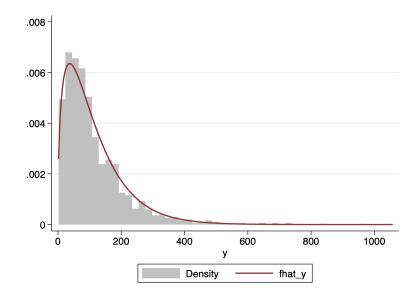
	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]
/theta1 /theta2	.3906123 4.349872	.033984	11.49 107.83	0.000	.3240049 4.270804	.4572196 4.42894

```
. mlci exp /theta1
```

Plot the estimated density $\hat{f}(y)$ over the sample histogram

```
. gen fhat_y = gammaden(exp(_b[/theta1]), exp(_b[/theta2]), 0, y)
```

- . tw (hist y, bin(50)) (line fhat_y y, sort), name(p3, replace) legend(rows(1))
- . graph export p3.png, replace
 (file p3.png written in PNG format)



Exercise 3

We assume that f(y) is log-normal distributed. That is, we assume that the biomarker is standard normal distributed after we apply the transform

$$G(y) = (\log(y) - \mu)/\sigma$$

The derivative of G(y) with respect to y is

$$G'(y) = g(y) = 1/(y\sigma).$$

Estimate the parameters μ and σ . Constrain σ to be positive.

^{1.477885 95%} CI: 1.382654, 1.579676

[.] mlci exp /theta2

^{77.46854 95%} CI: 71.57918, 83.84246

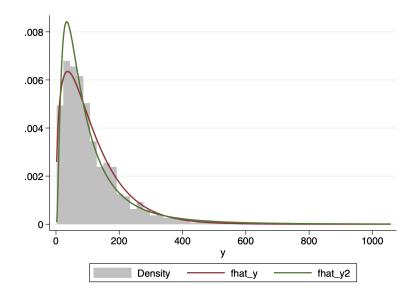
```
. local f = "normalden(`G´)*`g´"
. mlexp (log(`f'))
               log likelihood = -21814.225
initial:
alternative:
               log\ likelihood = -12440.421
               log likelihood = -10178.274
rescale:
rescale eq:
               log likelihood = -8264.4966
               log likelihood = -8264.4966
Iteration 0:
               log likelihood = -8198.7786
Iteration 1:
Iteration 2:
               log likelihood = -8159.1222
               log likelihood = -8158.8813
Iteration 3:
Iteration 4:
               log\ likelihood = -8158.8812
Maximum likelihood estimation
Log likelihood = -8158.8812
                                                  Number of obs
                                                                            1,432
                             Std. Err.
                    Coef.
                                             z
                                                  P>|z|
                                                            [95% Conf. Interval]
                 4.365484
                             .0242269
                                                  0.000
                                                               4.318
                                                                         4.412968
         /mu
                                         180.19
      /theta
                 -.0868798
                             .0186859
                                          -4.65
                                                  0.000
                                                           -.1235034
                                                                        -.0502561
```

. mlci exp /theta .9167873 95% CI: .8838186, .9509858

Compare the likelihood with that from the gamma model

Plot the estimated density $\hat{f}(y)$ over the sample histogram

```
. gen fhat_y2 = normalden((log(y) - _b[/mu]) / exp(_b[/theta]))*(1 / y / exp(_b[/theta])) . tw (hist y, bin(50)) (line fhat_y fhat_y2 y, sort), name(p4, replace) legend(rows(1)) . graph export p4.png, replace (file p4.png written in PNG format)
```



Exercise 4

We make the transform G(y) more flexible using polynomials. Consider the transform

$$G(y) = (\log(y) + \eta \log(y)^2 - \mu)/\sigma$$

The derivative of G(y) with respect to y is

$$G'(y) = g(y) = (1 + 2\eta \log(y)) / (\sigma y)$$

Estimate the parameters μ, σ, η . Constrain σ to be positive.

```
. local sigma = "exp({theta})"
. local G = "(\log(y) + \{eta\} * \log(y)^2 - \{mu\}) / `sigma`"
. local g = "(1 + \{eta\}*2*log(y)) / ('sigma'*y)"
. local f = "normalden(`G´)*`g´"
. mlexp (log(`f´))
initial:
               log likelihood = -21814.225
               log likelihood = -62186.361
alternative:
rescale:
               log likelihood = -21814.225
               log likelihood = -21814.225
rescale eq:
Iteration 0:
               log\ likelihood = -21814.225
                                             (not concave)
Iteration 1:
               log likelihood = -10891.752
                                             (not concave)
               log likelihood = -8663.3365
Iteration 2:
Iteration 3:
               log\ likelihood = -8355.3282
               \log likelihood = -8210.2164
Iteration 4:
               \log likelihood = -8167.6715
Iteration 5:
Iteration 6:
               \log likelihood = -8151.9226
               \log = -8142.8803
Iteration 7:
Iteration 8:
               log likelihood = -8139.5952
               \log = -8138.8787
Iteration 9:
Iteration 10: log likelihood = -8138.5163
Iteration 11:
               log\ likelihood = -8138.5033
              log likelihood = -8138.5022
Iteration 12:
Iteration 13: log likelihood = -8138.5022
Maximum likelihood estimation
Log likelihood = -8138.5022
                                                 Number of obs
                                                                          1,432
                    Coef.
                            Std. Err.
                                            z
                                                 P>|z|
                                                           [95% Conf. Interval]
                 .1804595
                             .0689773
                                                 0.009
                                                           .0452665
                                                                        .3156524
        /eta
                                          2.62
                 7.956257
                             1.37386
                                          5.79
                                                 0.000
                                                           5.263542
                                                                        10.64897
         /mu
```

. mlci exp /theta 2.307437 95% CI: 1.464886, 3.634595

/theta

Plot the estimated density $\hat{f}(y)$ over the sample histogram

.2318207

.8361374

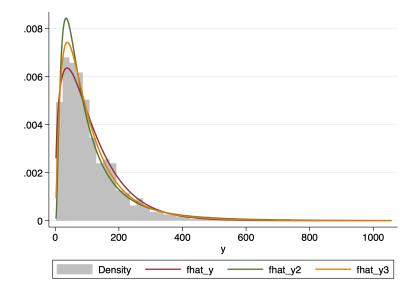
```
. gen fhat_y3 = normalden((log(y)+_b[/eta]*log(y)^2 - _b[/mu])/exp(_b[/theta])) * ///
> (1+_b[/eta]*2*log(y)) / (exp(_b[/theta]) * y)
. tw (hist y, bin(50)) (line fhat_y fhat_y2 fhat_y3 y, sort), name(p5, replace) legend(rows(1))
. graph export p5.png, replace
(file p5.png written in PNG format)
```

0.000

.3817772

3.61

1.290498



Exercise 5

Instead of a quadratic term, we add two restricted cubic splines transforms of $\log(y)$: $V_2(\log(y))$ and $V_3(\log(y))$. We consider the transform

$$G(y) = (\log(y) + \eta_1 V_2(\log(y)) + \eta_2 V_3(\log(y)) - \mu) / \sigma$$

The derivative of G(y) with respect to y is

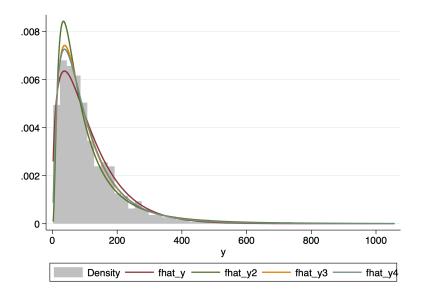
$$G'(y) = g(y) = (1 + \eta_1 v_2(\log(y)) + \eta_2 v_3(\log(y))) / (\sigma y)$$

Estimate the parameters μ , σ , η_1 , η_2 . Constrain σ to be positive. Jointly test the 2 parameters η_1 , η_2 to assess whether adding the 2 RCS transforms improves the fit of this model with respect to the "basic" log-normal model (see Exercise 3).

```
. rcsgen log_y, gen(V) dgen(v) df(3)
Variables V1 to V3 and v1 to v3 were created
. local sigma = "exp({theta})"
. local G = (\log(y) + \{eta1\} * V2 + \{eta2\} * V3 - \{mu\}) / sigma''
. local g = "(1+\{eta1\}*v2+\{eta2\}*v3)/(`sigma`*y)"
. local f = "normalden(`G')*`g'"
. mlexp (log(`f´))
initial:
               log likelihood = -21814.225
               log likelihood = -21814.225
final:
               log likelihood = -21814.225
rescale:
               \log likelihood = -21814.225
Iteration 0:
                                             (not concave)
Iteration 1:
               log likelihood = -15749.959
                                             (not concave)
               log likelihood = -11575.393
Iteration 2:
                                             (not concave)
               log likelihood = -8930.8791
Iteration 3:
                                             (not concave)
               \log = -8353.802
Iteration 4:
               log likelihood = -8226.7656
Iteration 5:
               log likelihood = -8161.3477
Iteration 6:
               log\ likelihood = -8141.3375
Iteration 7:
               log likelihood = -8137.886
Iteration 8:
               log likelihood = -8137.333
Iteration 9:
Iteration 10: log likelihood = -8137.3305
Iteration 11: log likelihood = -8137.3305
Maximum likelihood estimation
Log likelihood = -8137.3305
                                                  Number of obs
                                                                           1,432
                                                            [95% Conf. Interval]
                    Coef.
                            Std. Err.
                                            z
                                                 P>|z|
       /eta1
                -.0080713
                             .0421601
                                         -0.19
                                                 0.848
                                                           -.0907036
                                                                          .074561
       /eta2
                -.0259272
                             .0451895
                                         -0.57
                                                 0.566
                                                            -.114497
                                                                         .0626426
         /mu
                 5.045942
                             .2441727
                                         20.67
                                                 0.000
                                                            4.567372
                                                                        5.524512
                  .2911407
                             .1009571
                                          2.88
                                                 0.004
                                                            .0932684
                                                                          .489013
      /theta
```

Plot the estimated density $\hat{f}(y)$ over the sample histogram

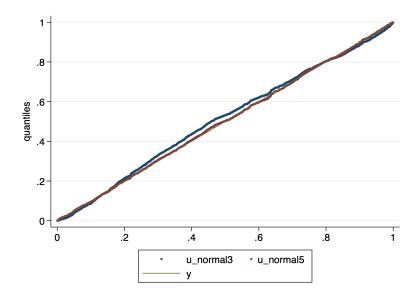
```
. gen fhat_y4 = normalden((log(y)+_b[/eta1]*V2+_b[/eta2]*V3 - _b[/mu])/exp(_b[/theta])) * ///
> (1+_b[/eta1]*v2+_b[/eta2]*v3) / (exp(_b[/theta]) * y)
. tw (hist y, bin(50)) (line fhat_y fhat_y2 fhat_y3 fhat_y4 y, sort), name(p6, replace) legend(rows(1))
. graph export p6.png, replace
(file p6.png written in PNG format)
```



Exercise 6

Let's assess the goodness-of-fit of the log-normal model with RCS transforms (see Exercise 5) and of the log-normal model (see Exercise 3) using a quantile plot.

```
. gen u_normal5 = normal((log(y)+_b[/eta1]*V2+_b[/eta2]*V3 - _b[/mu])/exp(_b[/theta]))
. // Re-fit log-normal model (Exercise 3)
. local sigma = "exp({theta})"
. local G = "(log(y) - {mu}) / `sigma`"
. local g = "(1 / y / sigma')"
. local f = "normalden(`G')*`g'"
. mlexp (log(`f´))
initial:
               log likelihood = -21814.225
               log likelihood = -12440.421
alternative:
rescale:
               log likelihood = -10178.274
               log likelihood = -8264.4966
rescale eq:
Iteration 0:
               log likelihood = -8264.4966
               log likelihood = -8198.7786
Iteration 1:
Iteration 2:
               log\ likelihood = -8159.1222
Iteration 3:
               log likelihood = -8158.8813
               log likelihood = -8158.8812
Iteration 4:
Maximum likelihood estimation
                                                 Number of obs
Log likelihood = -8158.8812
                                                                           1,432
                    Coef.
                            Std. Err.
                                                 P>|z|
                                                            [95% Conf. Interval]
                                            z
         /mu
                 4.365484
                             .0242269
                                        180.19
                                                 0.000
                                                               4.318
                                                                        4.412968
                -.0868798
                             .0186859
                                                 0.000
                                                          -.1235034
                                                                       -.0502561
      /theta
                                         -4.65
. gen u_normal3 = normal((log(y) - _b[/mu])/exp(_b[/theta]))
. qplot u_normal3 u_normal5, addplot(function y = x, lw(medthin)) name(p7, replace) ///
  msym(Oh Oh) msize(tiny tiny)
. graph export p7.png, replace
(file p7.png written in PNG format)
```



Extra: Exercise 7 (more on transforms of random variables)

We now assume that f(y) is gamma-distributed after square root transform.

$$G(y) = \sqrt{y}$$

The derivative is

$$G'(y) = g(y) = 0.5/\sqrt{y}$$

Estimate the parameters α and β using the gammaden() function. Fix the location parameter g to be equal to 0. Constrain α and β to be positive. Compare the likelihood with that form the log-normal and gamma models

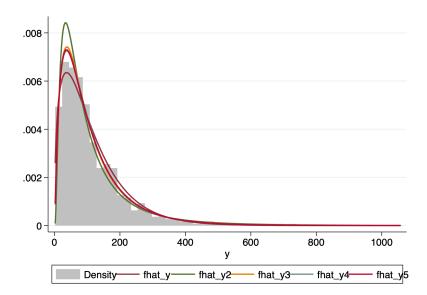
```
. local G = "sqrt(y)"
. local g = "(0.5 / sqrt(y))"
. local f = \text{"gammaden}(\exp(\{\text{theta1}\}), \exp(\{\text{theta2}\}), 0, G') * G') * G')
. mlexp (log(`f´))
initial:
                log likelihood = -18140.526
                \log = -11624.987
alternative:
                \log likelihood = -8442.1703
rescale:
                \log = -8442.1703
rescale eq:
Iteration 0:
                log\ likelihood = -8442.1703
Iteration 1:
                \log likelihood = -8185.3624
                \log \text{ likelihood} = -8138.5993
Iteration 2:
                log likelihood = -8138.2943
Iteration 3:
                \log = -8138.2942
Iteration 4:
Maximum likelihood estimation
Log likelihood = -8138.2942
                                                   Number of obs
                                                                              1,432
                                                   P>|z|
                                                              [95% Conf. Interval]
                     Coef.
                              Std. Err.
                                              z
     /theta1
                  1.652508
                              .0362406
                                           45.60
                                                   0.000
                                                              1.581478
                                                                           1.723538
     /theta2
                  .6290655
                              .0380415
                                           16.54
                                                   0.000
                                                              .5545055
                                                                           .7036256
```

```
. mlci exp /theta1
5.220056 95% CI: 4.862136, 5.604323
. mlci exp /theta2
1.875857 95% CI: 1.74108, 2.021067
```

Plot the estimated density $\hat{f}(y)$ over the sample histogram. Visually compare the estimated density from the lognormal + splines model with the density from the gamma model after square root transform. What do you

conclude?

```
. gen fhat_y5 = gammaden(exp(_b[/theta1]), exp(_b[/theta2]), 0, sqrt(y))*(.5 / sqrt(y))
. tw (hist y, bin(50)) (line fhat_y fhat_y2 fhat_y3 fhat_y4 fhat_y5 y, sort), name(p8, replace) legend(rows(1))
. graph export p8.png, replace
(file p8.png written in PNG format)
```



Extra: Exercise 8 (more on goodness of fit)

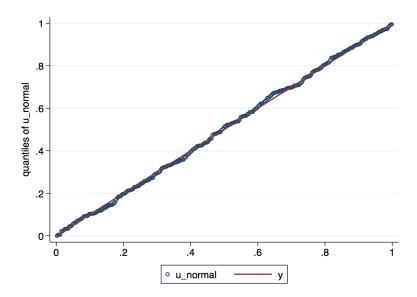
Let's go back to the normal distributed variable (Exercise 1, Lab 1).

. use https://raw.githubusercontent.com/anddis/fsm/master/data/lab1.dta, clear

Assume that $f(y_n)$ is normal and estimate the parameters μ and σ . Generate the transform $u = \hat{F}(y_n)$. Draw the estimated quantile plot using the qplot command.

```
. local f = "normalden(y_n, {mu}, exp({theta}))"
. mlexp(ln(`f'))
               log likelihood =
initial:
                                     -<inf>
                                             (could not be evaluated)
feasible:
               log likelihood = -32398.765
rescale:
               log likelihood = -1981.1218
               \log \text{ likelihood} = -1440.3171
rescale eq:
               log likelihood = -1440.3171
Iteration 0:
                                             (not concave)
Iteration 1:
               \log likelihood = -1112.6119
               log likelihood = -1085.7986
Iteration 2:
Iteration 3:
               log\ likelihood = -1059.4172
               \log = -1059.332
Iteration 4:
               log likelihood = -1059.3319
Iteration 5:
Maximum likelihood estimation
Log likelihood = -1059.3319
                                                                             300
                                                 Number of obs
                             Std. Err.
                                                            [95% Conf. Interval]
                    Coef.
                                            z
                                                 P>|z|
                                                 0.000
                                                            177.5557
                                                                        179.4265
         /mu
                 178,4911
                             .4772459
                                        374.00
      /theta
                 2.112168
                             .0408248
                                         51.74
                                                 0.000
                                                            2.032153
                                                                        2.192183
```

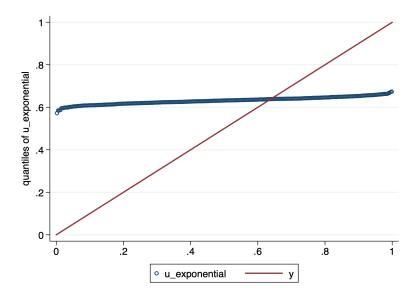
```
. mlci exp /theta
8.266141 95% CI: 7.630494, 8.954739
.
. gen u_normal = normal((y_n-_b[/mu])/exp(_b[/theta]))
. qplot u_normal, addplot(function y = x) name(p1, replace)
. graph export p0.png, replace
(file p0.png written in PNG format)
```



Assume now that $f(y_n)$ is exponential and estimate the parameter λ . Generate the transform $u = \hat{F}(y_n)$. Draw the estimated quantile plot using the qplot command.

```
. local f = "exp({theta})*exp(-y_n * exp({theta}))"
. mlexp(ln(`f'))
initial:
                log likelihood = -53547.324
log likelihood = -32628.094
alternative:
                \log likelihood = -2180.7534
rescale:
                \log likelihood = -2180.7534
Iteration 0:
                log likelihood = -1857.2088
log likelihood = -1855.3682
Iteration 1:
Iteration 2:
                \log likelihood = -1855.3616
Iteration 3:
                log likelihood = -1855.3616
Iteration 4:
Maximum likelihood estimation
Log likelihood = -1855.3616
                                                     Number of obs
                                                                                   300
                                                     P>|z|
                                                                 [95% Conf. Interval]
                      Coef.
                               Std. Err.
                                                z
      /theta
                  -5.184539
                                .057735
                                           -89.80
                                                     0.000
                                                               -5.297697
                                                                              -5.07138
. mlci exp /theta
            95% CI: .0050031, .0062738
. gen u_exponential = 1-exp(-y_n * exp(_b[/theta]))
. qplot u_exponential, addplot(function y = x) name(p2, replace)
```

. graph export p00.png, replace (file p00.png written in PNG format)



What can you conclude about the goodness of fit of the normal and exponential model?

Extra: Exercise 9 (binary variables)

Assume that y_{ber} follows a Bernoulli distribution. We want to estimate the probability of "success" ($y_{ber} = 1$). Estimate the probability η while constraining it to be bounded between 0 and 1. First, write down the likelihood by hand. Then, use the binomialp() function.

Are the results you obtain identical to those obtained from logistic regression?

```
. local eta = "invlogit({theta})"
. local f = "`eta´^y_ber * (1-\text{`eta'})^(1-y_ber)"
. mlexp (ln(`f'))
initial:
               log\ likelihood = -207.94415
               log likelihood = -199.7231
alternative:
rescale:
               log\ likelihood = -199.7231
               log likelihood = -199.7231
Iteration 0:
               log likelihood = -199.70172
Iteration 1:
Iteration 2:
               log likelihood = -199.70172
Maximum likelihood estimation
Log likelihood = -199.70172
                                                  Number of obs
                                                                              300
                    Coef.
                             Std. Err.
                                            z
                                                 P>|z|
                                                            [95% Conf. Interval]
      /theta
                  .4754237
                             .1187479
                                          4.00
                                                 0.000
                                                            .2426821
                                                                         .7081653
 mlci invlogit /theta
 .6166667
            95% CI: .5603745, .6699956
. local eta = "invlogit({theta})"
. local f = "binomialp(1, y_ber, `eta´)"
. mlexp (ln(`f'))
               log likelihood = -207.94415
initial:
alternative:
               log likelihood = -199.7231
               log likelihood = -199.7231
rescale:
               log likelihood = -199.7231
Iteration 0:
               log likelihood = -199.70172
Iteration 1:
               log likelihood = -199.70172
Iteration 2:
Maximum likelihood estimation
Log likelihood = -199.70172
                                                  Number of obs
                                                                              300
                             Std. Err.
                                                            [95% Conf. Interval]
                    Coef.
                                                 P>|z|
                                            z
```

/theta	.4754237	.1187479	4.00	0.000	. 2426821		.7081653						
. mlci invlogit /theta .6166667 95% CI: .5603745, .6699956													
logit y_ber													
<pre>Iteration 0: log likelihood = -199.70172 Iteration 1: log likelihood = -199.70172</pre>													
Logistic regre		Number o	f obs	=	300								
		LR chi2(0)	=	0.00								
				Prob > c	hi2	=							
Log likelihood	1 = -199.7017	2		Pseudo R	2	=	0.0000						
y_ber	Coef.	Std. Err.	z	P> z	[95%	Conf.	Interval]						
_cons	.4754237	.1187479	4.00	0.000	.2426	5821	.7081653						