

Modeling unobserved heterogeneity in Stata

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Plan of the talk

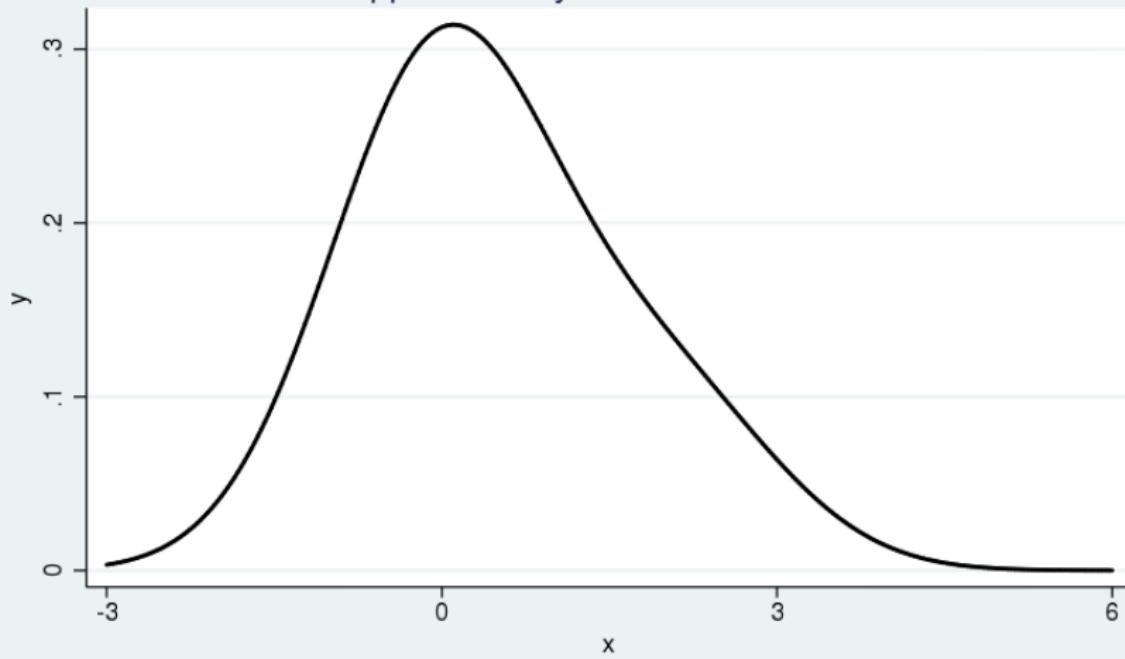
Concepts and terminology

Finite mixture models with **fmm**

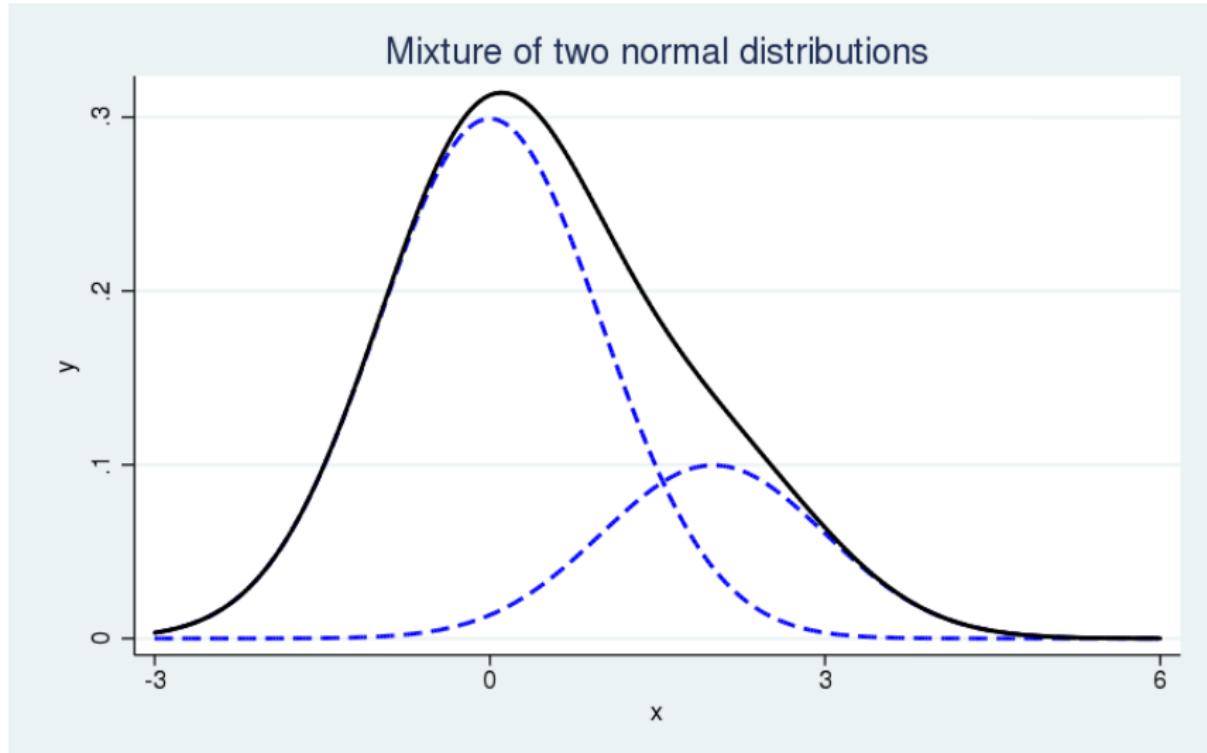
Latent class models with **gsem** ... **lclass()**

Observed distribution for a whole population:

Approximately normal distribution



Unobserved distributions of the two underlying subpopulations:



Unobserved heterogeneity refers to differences among individuals or observations that cannot be measured by regressors.

Latent class models

Latent – unobserved, hidden

Class – subpopulation, group, type, component, density, distribution

Finite mixture models

Finite – number of classes determined a priori

Mixture – of distributions, densities, regression models

Mixture of distributions:

The observed \mathbf{y} are assumed to come from g distinct distributions f_1, f_2, \dots, f_g in proportions or with probabilities $\pi_1, \pi_2, \dots, \pi_g$.

We can write a simple mixture model as

$$f(\mathbf{y}) = \sum_{i=1}^g \pi_i f_i(\mathbf{y} | \mathbf{x}' \boldsymbol{\beta}_i)$$

where π_i is the probability for the i th class, and

$f_i(\cdot)$ is the conditional probability density function (pdf) for the observed response in the i th class model.

(continued)

$$f(\mathbf{y}) = \sum_{i=1}^g \pi_i f_i(\mathbf{y} | \mathbf{x}' \boldsymbol{\beta}_i)$$

We use the multinomial logistic distribution to model the probabilities for the latent classes.

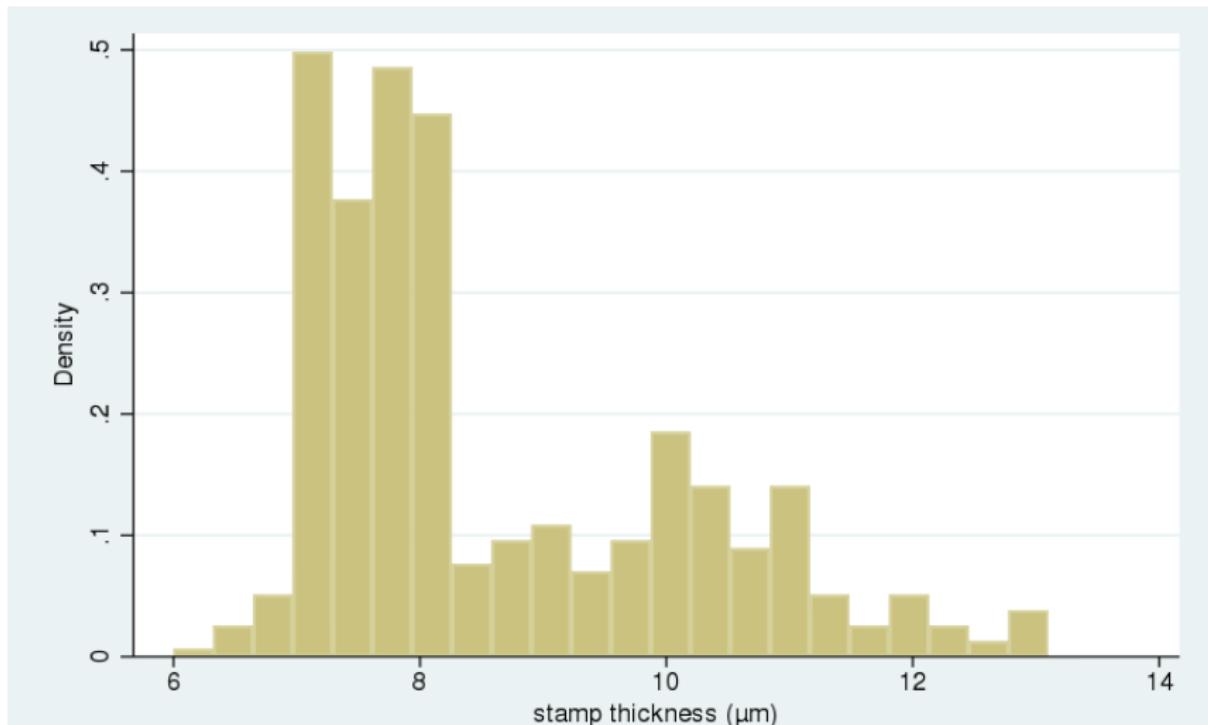
$$\pi_i = \frac{\exp(\gamma_i)}{\sum_{j=1}^g \exp(\gamma_j)}$$

where γ_i is the linear prediction for the i th latent class.

By convention, the first latent class is the base category, $\gamma_1 = 0$.

Example: Postal stamp thickness

```
. webuse stamp  
. gen thick = thickness*100  
. label var thick "stamp thickness ( $\mu\text{m}$ )"  
. histogram thick
```



We want to model the empirical distribution as a mixture of two normal distributions:

$$f(\mathbf{y}) = \pi_1 \times N(\mu_1, \sigma_1^2) + \pi_2 \times N(\mu_2, \sigma_2^2)$$

This is as simple as typing:

```
. fmm 2 : regress thick
```

where

fmm 2 means we have two components

and **regress** is a keyword for “normal distribution”

Finite mixture model		Number of obs	=	485				
Log likelihood = -748.75749								
<hr/>								
		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]		
<hr/>								
1.Class		(base outcome)						
<hr/>								
2.Class								
	_cons		-.4498027	.124093	-3.62	0.000	-.6930205	-.2065848
<hr/>								

Class : 1
Response : thick
Model : regress

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
thick						
_cons		7.609076	.0297275	255.96	0.000	7.550811 7.667341
var(e.thick)		.206297	.022201			.1670665 .2547395

Class : 2
Response : thick
Model : regress

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
thick						
_cons		10.16013	.1427942	71.15	0.000	9.880254 10.44
var(e.thick)		1.441319	.2583438			1.014354 2.048003

Recall we use the multinomial logistic distribution to model the probabilities for the latent classes:

$$\pi_i = \frac{\exp(\gamma_i)}{\sum_{j=1}^g \exp(\gamma_j)}$$

In simple cases, we can calculate latent class probabilities by hand:

```
. di      1 / ( 1 + exp(-.4498027) )
. di exp(-.4498027) / ( 1 + exp(-.4498027) )

.61059232
.38940768
```

This is a little bit easier:

```
. di      1 / ( 1 + exp(_b[2.Class:_cons]) )
. di exp(_b[2.Class:_cons]) / ( 1 + exp(_b[2.Class:_cons]) )

.61059232
.38940768
```

You can also use **predict** and **summarize**:

```
. predict pr*, classposteriorpr
```

```
. des pr1 pr2
```

variable name	storage type	display format	value label	variable label
pr1	float	%9.0g		Predicted posterior probability (1.Class)
pr2	float	%9.0g		Predicted posterior probability (2.Class)


```
. su pr1 pr2
```

Variable	Obs	Mean	Std. Dev.	Min	Max
pr1	485	.6105923	.4519458	1.53e-30	.9829751
pr2	485	.3894077	.4519458	.0170249	1

estat lcprob is your friend:

```
. estat lcprob
```

```
Latent class marginal probabilities           Number of obs      =      485
```

Delta-method			
	Margin	Std. Err.	[95% Conf. Interval]
<hr/>			
Class			
1 .6105923	.0295055	.5514633	.6666385
2 .3894077	.0295055	.3333615	.4485367
<hr/>			

Note that when you have a mixture of distributions, the posterior probability of being in a given class is the same for all observations with the same value.

```
. su pr1 pr2 if thick==8
```

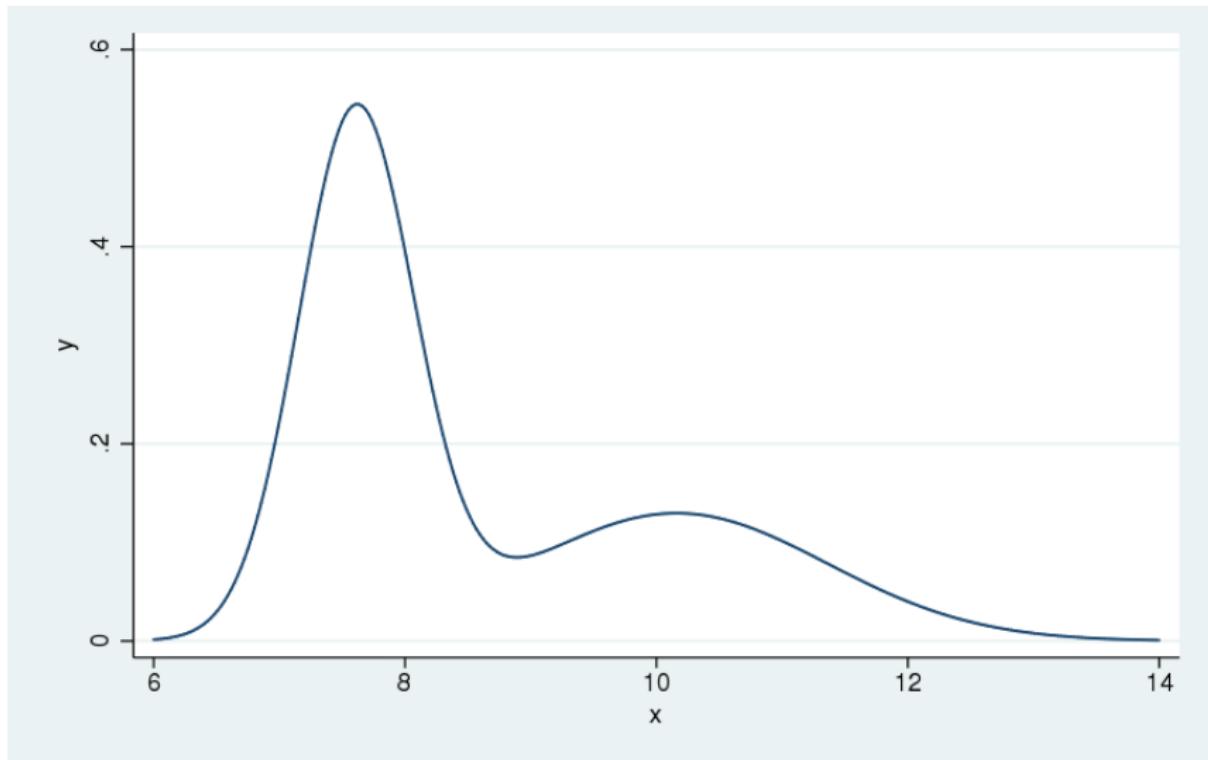
Variable	Obs	Mean	Std. Dev.	Min	Max
-----+-----					
pr1	37	.93524	0	.93524	.93524
pr2	37	.06476	0	.06476	.06476

This makes it easy to plot the estimated mixture density.

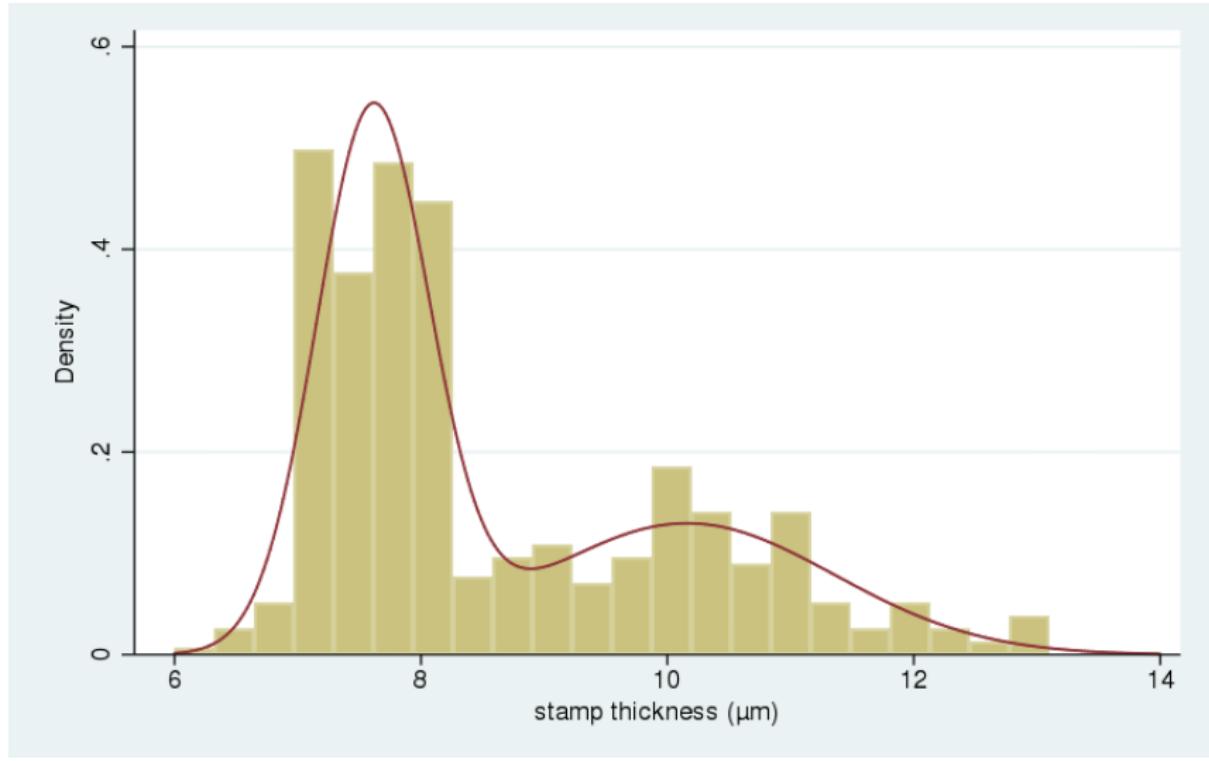
This is our estimated mixture density:

$$\hat{f}(\mathbf{y}) = .61 \times N(7.61, .21) + .39 \times N(10.16, 1.44)$$

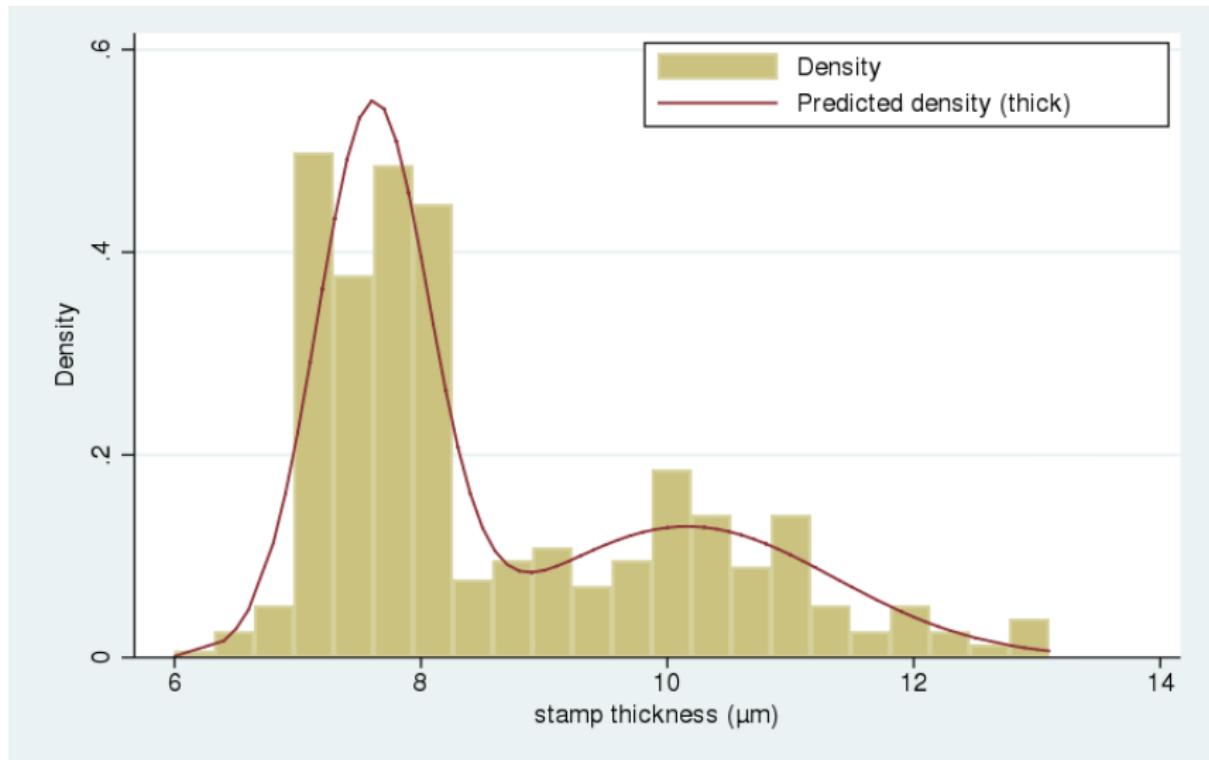
```
. twoway ///
    function .61*normalden(x,7.61,sqrt(.21)) + .39*normalden(x,10.16,sqrt(1.44)), range(6 14)
```



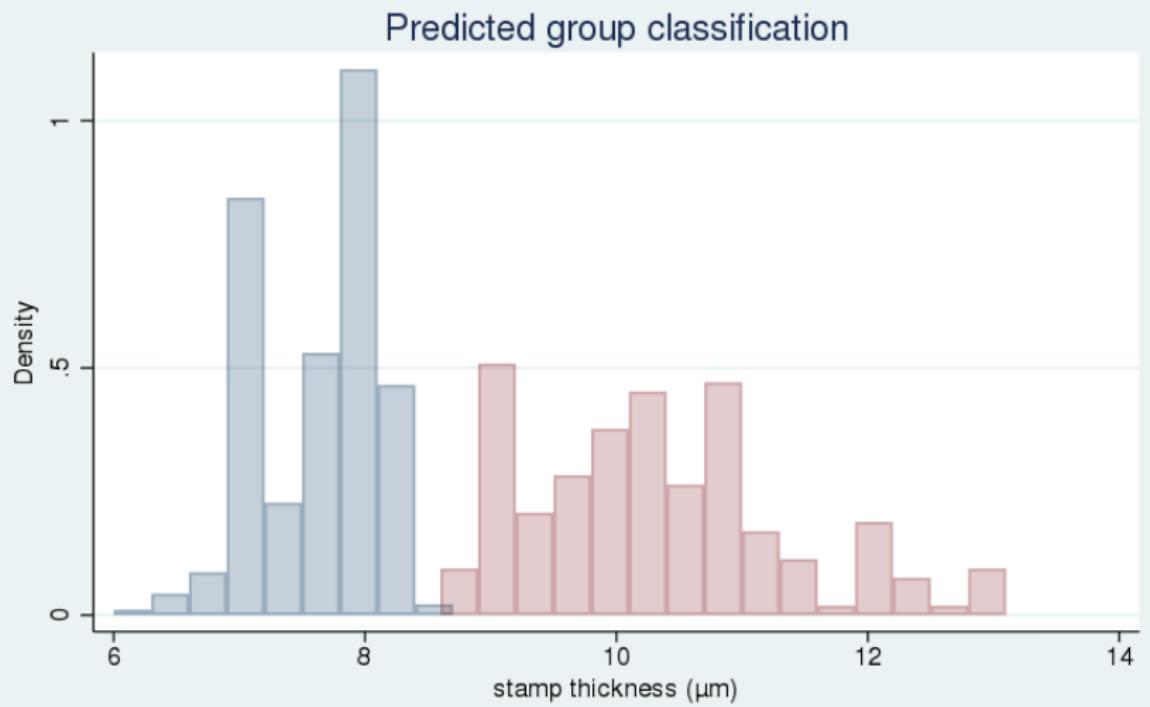
```
. histogram thick, addplot( ///
    function .61*normalden(x,7.61,sqrt(.21)) + .39*normalden(x,10.16,sqrt(1.44)) range(6 14) ///
) legend(off)
```



```
. predict den, density marginal  
. histogram thick, addplot(line den thick) legend(ring(0) pos(2))
```



```
. gen group = pr1 > .5  
. twoway histogram thick if group ... ///  
    histogram thick if !group ...
```



When we add covariates, we fit a mixture of “models”.

Here, we fit a mixture of two linear regression models.

```
. use chol  
(Fictional cholesterol data)
```



```
. describe
```

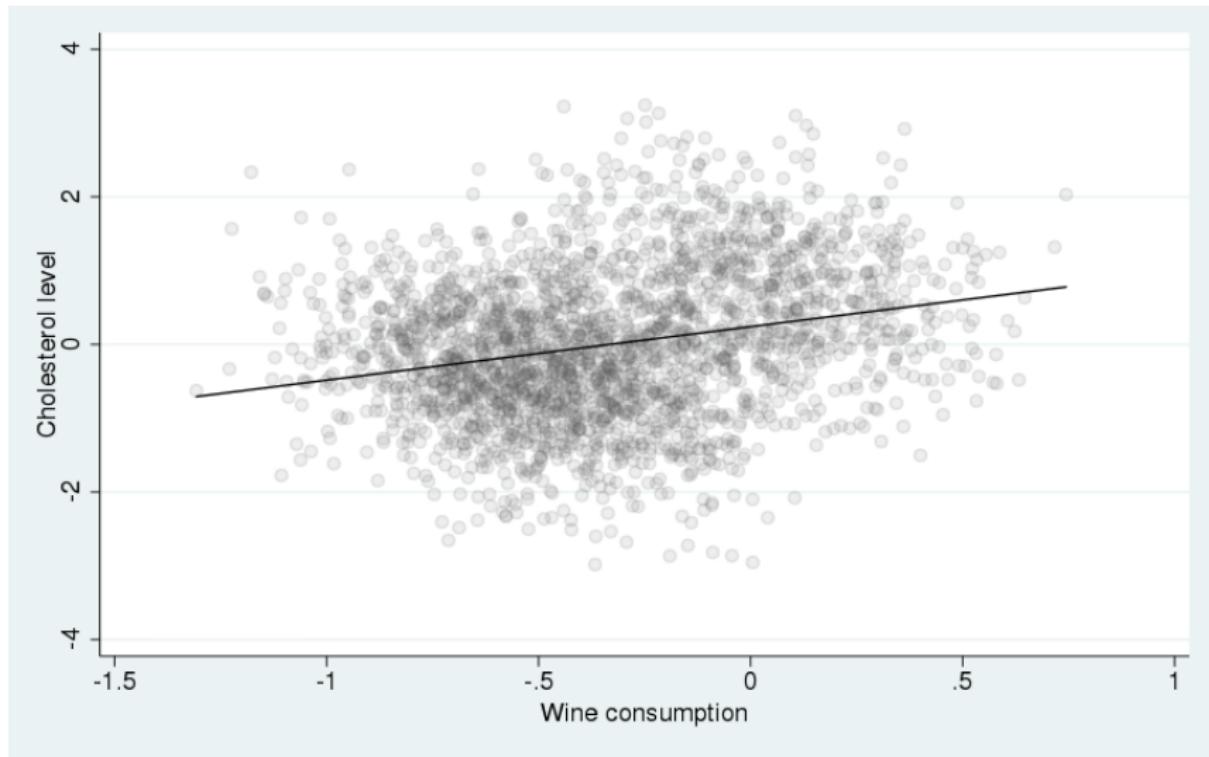
variable	name	storage	display	value	variable label
		type	format	label	
chol		float	%9.0g		Standardized cholesterol level
wine		float	%9.0g		Mean-centered monthly wine consumption
pchol		float	%9.0g		=1 if either parent has high cholesterol level

```
. regress chol wine
```

Source	SS	df	MS	Number of obs	=	2,500
				F(1, 2498)	=	171.27
Model	160.343489	1	160.343489	Prob > F	=	0.0000
Residual	2338.65652	2,498	.936211577	R-squared	=	0.0642
				Adj R-squared	=	0.0638
Total	2499.00001	2,499		Root MSE	=	.96758

chol	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]
wine	.7243775	.0553511	13.09	0.000	.6158387 .8329162
_cons	.2408989	.0267081	9.02	0.000	.1885266 .2932712

```
. twoway (scatter chol wine ...) (lfit chol wine ...) ...
```



```
. fmm 2, lcprob(pchol): regress chol wine
```

Finite mixture model Number of obs = 2,500
Log likelihood = -3062.7143

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
1.Class		(base outcome)				
2.Class						
	pchol	7.473592	.8977705	8.32	0.000	5.713994 9.23319
	_cons	-3.228661	.3939579	-8.20	0.000	-4.000804 -2.456518

Class : 1
Response : chol
Model : regress

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
<hr/>						
chol						
	wine	-.6850974	.0783981	-8.74	0.000	-.8387549 -.5314399
	_cons	-.7401758	.0443478	-16.69	0.000	-.8270959 -.6532557
<hr/>						
	var(e.chol)	.6152073	.0219867		.5735887	.6598457
<hr/>						

Class : 2
Response : chol
Model : regress

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
<hr/>						
chol						
	wine	-.4798618	.1319125	-3.64	0.000	-.7384056 -.221318
	_cons	.8343004	.0323813	25.76	0.000	.7708342 .8977667
<hr/>						
	var(e.chol)	.6720669	.0383181		.601009	.7515261
<hr/>						

```
. predict c*, classposteriorpr
```

```
. su c?
```

Variable	Obs	Mean	Std. Dev.	Min	Max
c1	2,500	.6743291	.4538173	6.22e-09	1
c2	2,500	.3256709	.4538173	4.24e-11	1

```
. estat lcprob
```

Latent class marginal probabilities Number of obs = 2,500

Delta-method				
	Margin	Std. Err.	[95% Conf. Interval]	
Class				
1	.6743291	.0055936	.6632719	.6851956
2	.3256709	.0055936	.3148044	.3367281

```
. predict xb*
```

```
. su xb?
```

Variable	Obs	Mean	Std. Dev.	Min	Max
xb1	2,500	-.5123399	.2395686	-1.249959	.1553361
xb2	2,500	.9938833	.1678007	.477234	1.461543

```
. estat lcmean
```

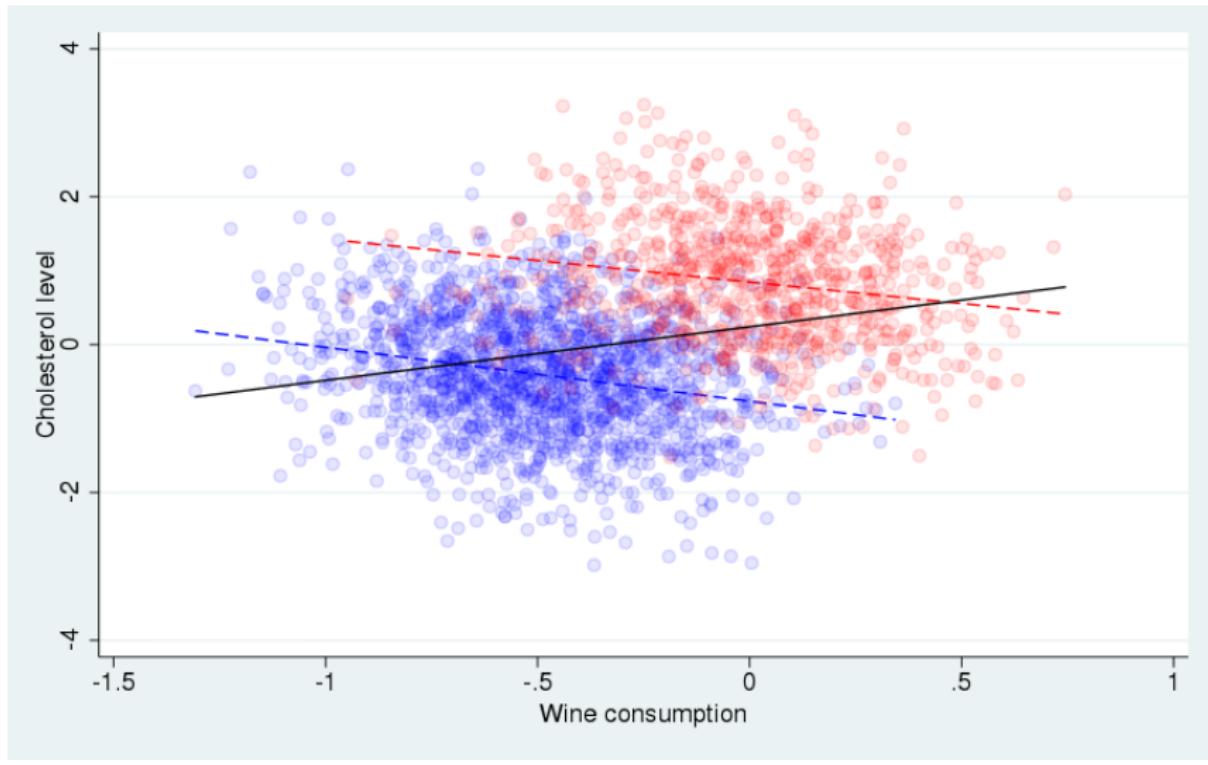
Latent class marginal means Number of obs = 2,500

Delta-method						
	Margin	Std. Err.	z	P> z	[95% Conf. Interval]	
1	chol	-.5123399	.024033	-21.32	0.000	-.5594438 -.465236
2	chol	.9938833	.0601744	16.52	0.000	.8759435 1.111823

```

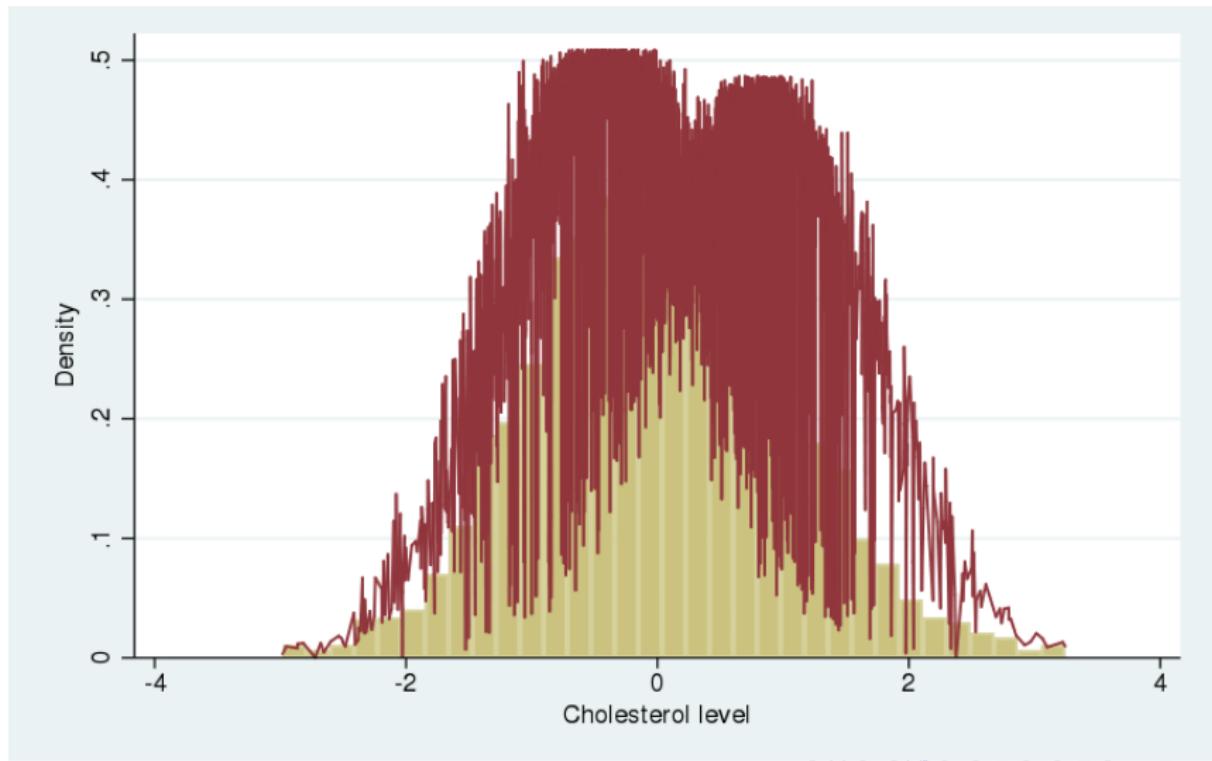
. gen grp = c1 > .5
. twoway (scatter chol wine if grp ...) (scatter chol wine if !grp ...) ///
    ( lfit chol wine if grp ...) ( lfit chol wine if !grp ...) ///
    ( lfit chol wine ... ) ...

```

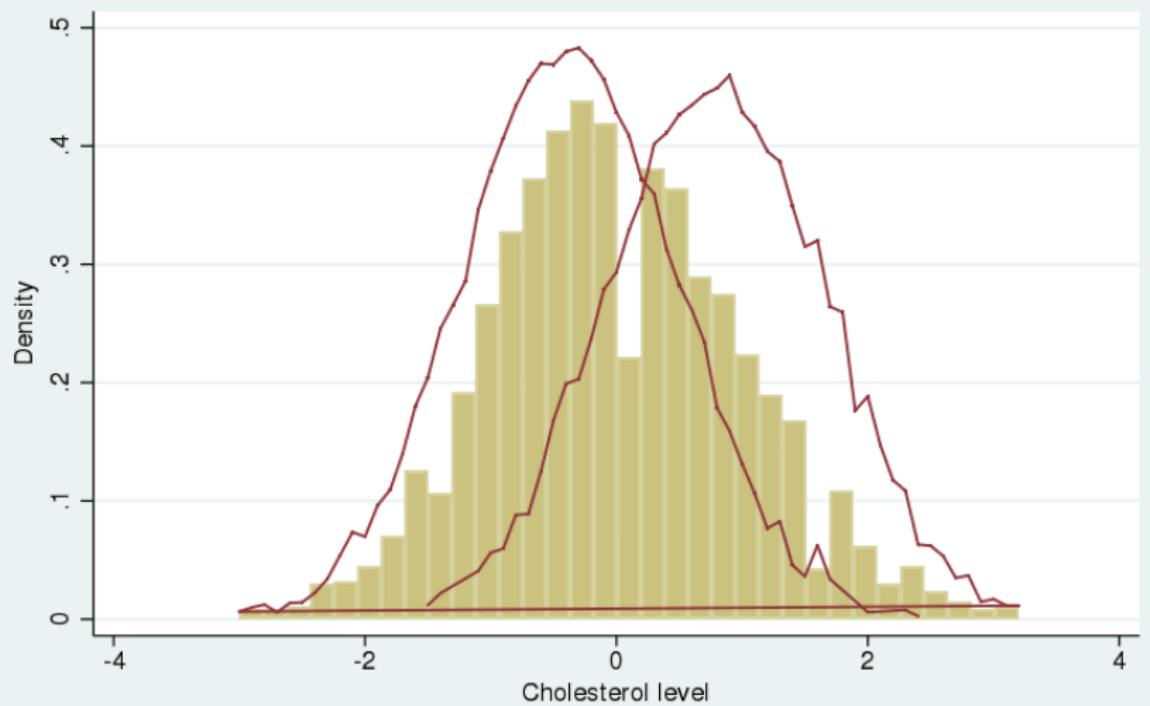


Some tips ...

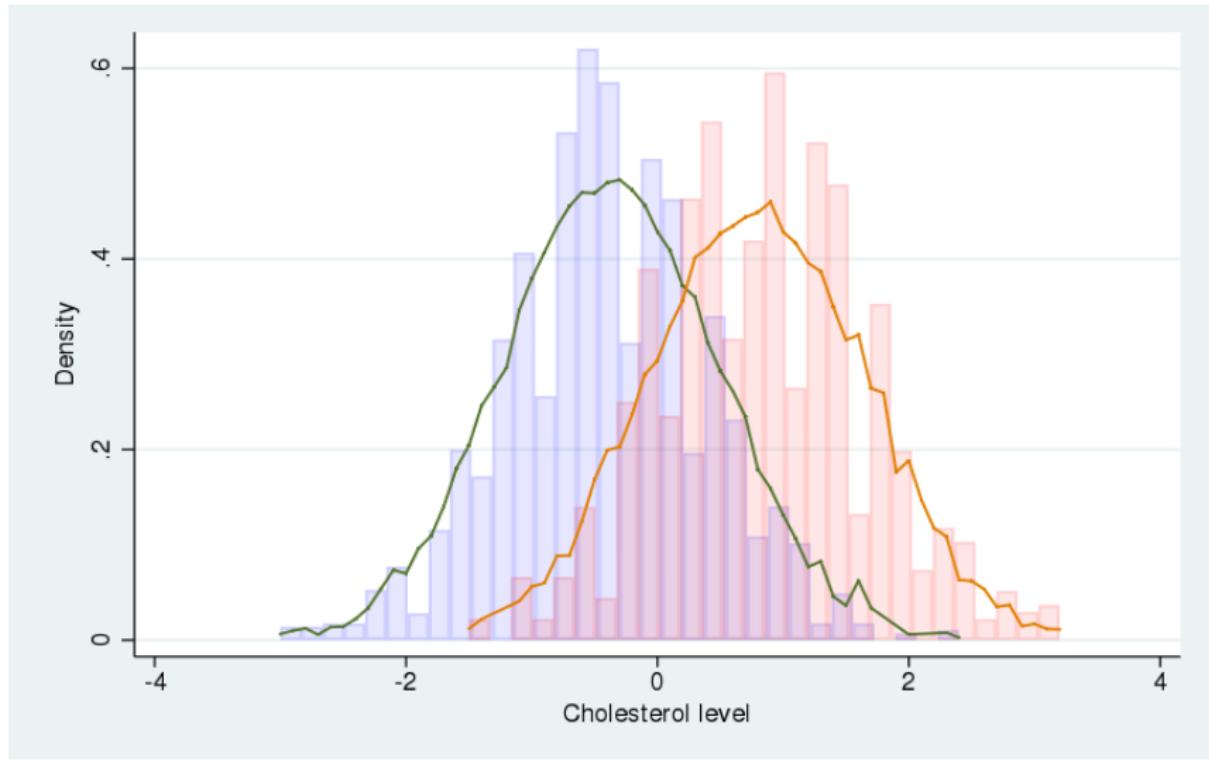
```
. predict den, density marginal  
. sort chol  
. histogram chol, addplot(line den chol) legend(off)
```



```
. gen chol12 = round(chol,.1)
. sort grp chol12
. by grp chol12 : egen den2 = mean(den)
. twoway (histogram chol12) (line den2 chol12) ...
```



```
. twoway ///
>     (histogram chol2 if grp, color(blue%10)) (line den2 chol2 if grp) ///
>     (histogram chol2 if !grp, color(red%10)) (line den2 chol2 if !grp) ...
```



regress is one of the many **fmm** keywords

Linear outcomes: **regress**, **truncreg**, **intreg**, **tobit**, and **ivregress**

Binary outcomes: **logit**, **probit**, and **cloglog**

Ordinal outcomes: **ologit** and **oprobit**

Nominal outcomes: **mlogit**

Count outcomes: **poisson**, **nbreg**, and **tpoisson**

Generalized linear models: **glm** with family **beta**, **exponential**,
gamma, **lognormal**, and more

Fractional outcomes: **betareg**

Survival outcomes: **streg**

fmm has two syntaxes.

You have seen the simple syntax

```
. fmm 3 : regress y x1 x2 x3
```

You can also use the hybrid syntax

```
. fmm : (regress y x1 x2 x3) (regress y x1 x2 x3) (regress y x1 x2 x3)
```

In the simple syntax, each component gets the same regressors:

```
. fmm 3 : regress y x1 x2 x3  
. fmm   : (regress y x1 x2 x3) (regress y x1 x2 x3) (regress y x1 x2 x3)
```

In the hybrid syntax, each component can have different regressors

```
. fmm : (regress y x1 x2) (regress y x2 x3) (regress y x3, noconstant)
```

In the hybrid syntax, you can also fit mixtures of different models or distributions:

```
. fmm : (regress y) (glm y, family(lognormal)) (tobit y, ll(0))
```

```
. fmm : (regress y x1) (glm y, family(lognormal)) (tobit y x1 x2, ll(0))
```

You can also model the latent class probabilities using the **lcprob()** option.

Simple syntax

```
. fmm 3, lcprob(w1 w2) : regress y x1 x2 x3
```

Hybrid syntax

```
. fmm : (regress y x1) (regress y x2 x3, lcprob(w1 w2)) (regress y x1 x3, lcprob(w2))
```

There is one special **fmm** keyword **pointmass** that allows one or more components to be a degenerate distribution taking on a single integer value with probability one.

This distribution cannot be used by itself and is always combined with other **fmm** keywords, most often to model zero-inflated outcomes.

This means you can use **fmm** in place of **zip** and **zinb**, and as an alternative to **zioprobit**.

Example: zero-inflated Poisson model

The zero-inflated Poisson model is a model in which the distribution of the outcome is a two-component mixture. One component is a distribution that is all zero. The other component is a Poisson distribution.

```
. webuse fish, clear
```

```
. describe
```

variable name	storage type	display format	value label	variable label
livebait	byte	%9.0g		1 if visitor uses live bait
camper	byte	%9.0g		1 if visitor is camping
persons	byte	%9.0g		number of persons accompanying the visitor
child	byte	%9.0g		number of children accompanying the visitor
count	int	%9.0g		number of fish caught

```
. su count
```

Variable	Obs	Mean	Std. Dev.	Min	Max
count	250	3.296	11.63503	0	149



```
. zip count persons livebait, inflate(child camper)  
  
. fmm : (poisson count persons livebait) (pointmass count, lcprob(child camper))
```

Finite mixture model
Number of obs = 250
Log likelihood = -850.70142

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]

1.Class						
	child	1.602571	.2797719	5.73	0.000	1.054228 2.150913
	camper	-1.015698	.365259	-2.78	0.005	-1.731593 -.2998039
	_cons	-.4922872	.3114562	-1.58	0.114	-1.10273 .1181558

2.Class		(base outcome)				

Class : 2
Response : count
Model : poisson

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]

count						
	persons	.8068853	.0453288	17.80	0.000	.7180424 .8957281
	livebait	1.757289	.2446082	7.18	0.000	1.277866 2.236713
	_cons	-2.178472	.2860289	-7.62	0.000	-2.739078 -1.617865

```
. estat lcprob
```

Latent class marginal probabilities Number of obs = 250

		Delta-method		
		Margin	Std. Err.	[95% Conf. Interval]
Class				
1		.4786335	.0341083	.4125554 .5454678
2		.5213665	.0341083	.4545322 .5874446

```
. estat lcmean
```

Latent class marginal means Number of obs = 250

Expression : Predicted mean (number of fish caught in class 2.Class),
predict(outcome(count) class(2))

		Delta-method				
		Margin	Std. Err.	z	P> z	[95% Conf. Interval]
2						
	count	6.490014	.2361623	27.48	0.000	6.027144 6.952884

With **pointmass()** you can run any imaginable inflated model (not that they all make sense though)

```
. fmm : (poisson y x1) (poisson y x2) (pointmass y)

. fmm : (poisson y x1) (poisson y x2) (pointmass y) (pointmass y, value(5))

. fmm : (ologit y) (pointmass y, value(1))

. fmm : (ologit y) (pointmass y, value(2)) (pointmass y, value(4))

. fmm : (mlogit y x1 x2 x3) (pointmass y, value(3))
```

With **fmm** you cannot fit mixture models for multiple responses. In other words, each class can have only one dependent variable.

. use chol, clear							
. fmm : (regress chol) (regress wine)							
Class	:	1					
Response	:	chol					
Model	:	regress					
		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
chol							
_cons		-.0450922	848.3992	-0.00	1.000	-1662.877	1662.787
var(e.chol)		.0584758	206.5966			.	.
Class	:	2					
Response	:	wine					
Model	:	regress					
		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
wine							
_cons		-.3325602	.0069923	-47.56	0.000	-.3462649	-.3188556
var(e.wine)		.1222298	.0034571			.1156383	.129197

If you want each class to have two outcomes, **chol** and **wine**, you need to go through **gsem**.

FMM in gsem:

A model with categorical latent variables and categorical observed variables is called a latent class model.

A model with categorical latent variables and continuous observed variables is called a latent profile model.

A finite mixture model can be either.

Start with a mixture of two normal distributions.

```
fmm 2: regress chol
```

Different ways of doing the same in **gsem**:

```
gsem (chol <-) , lclass(Class 2) lcinvariant(none)
```

```
gsem (chol <-) (chol <-) , lclass(Class 2) lcinvariant(none)
```

```
gsem (1: chol <-) (2: chol <-) , lclass(Class 2) lcinvariant(none)
```

The default model is linear regression so these are equivalent:

```
fmm 2: regress chol  
  
gsem (chol <-) , lclass(Class 2) lcinviant(none)  
  
gsem (chol <-, regress) , lclass(Class 2) lcinviant(none)  
  
gsem (chol <-, family(gaussian)) , lclass(Class 2) lcinviant(none)  
  
gsem (chol <-, family(gaussian) link(identity)) , lclass(Class 2) lcinviant(none)
```

Adding covariates is easy.

Below, both class models receive **the same** covariates:

```
fmm 2: regress chol wine
```

```
gsem (chol <- wine) , lclass(Class 2) lcinvariant(none)
```

```
gsem (chol <- wine) (chol <- ) , lclass(Class 2) lcinvariant(none)
```

You have to be more explicit if you want the model for class 2 to have only the constant term:

```
fmm : (regress chol wine) (regress chol)
```

```
gsem (1: chol <- wine) (2: chol <- ) , lclass(Class 2) lcinvariant(none)
```

Now you should be able to figure out how to fit a mixture model for multiple responses.

Pseudo-**fmm** syntax:

```
fmm 2: regress chol wine =
```

gsem syntax (real):

```
gsem (chol wine <-), lclass(Class 2) lcinvariant(none)
```

```
gsem (chol <-) (wine <-), lclass(Class 2) lcinvariant(none)
```

If you want class 1 to have two dependent variables and class 2 to have one dependent variable, you have to be explicit about it:

```
gsem (1: chol wine <-) (2: chol <- pchol), lcclass(Class 2) lcinvariant(none)
```

Generalized structural equation model Number of obs = 2,500
Log likelihood = -3217.3559

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
1.Class	(base outcome)					
2.Class						
	_cons	2.817545	.2573915	10.95	0.000	2.313067 3.322023

Class : 1

Response : chol

Family : Gaussian

Link : identity

Response : wine

Family : Gaussian

Link : identity

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
chol	_cons	-.7664867	.131116	-5.85	0.000	-1.023469 -.5095042
wine	_cons	-.4772524	.0243019	-19.64	0.000	-.5248833 -.4296215
var(e.chol)		.2887896	.1144431		.1328198	.6279142
var(e.wine)		.0100204	.0032932		.0052619	.0190822

Class : 2

Response : chol

Family : Gaussian

Link : identity

		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
chol						
pchol		.4683264	.0184701	25.36	0.000	.4321258 .5045271
_cons		.0322942	.0206101	1.57	0.117	-.0081007 .0726892
var(e.chol)		.7804484	.0254232		.7321771	.8319021

For zero-inflated models in **gsem**, use **family(pointmass)**:

```
webuse fish, clear

fmm : (pointmass count) (poisson count persons livebait)

gsem ///

(1: count <- , family(pointmass)) ///
(2: count <- persons livebait, family(poisson)) ///
, ///
lclass(Class 2) lcinvariant(none)
```

To model class probabilities, add a class equation:

```
fmm : (pointmass count, lcprob(child camper)) (poisson count persons livebait)
gsem ///
(1: count <-, family(pointmass)) ///
(2: count <- persons livebait, family(poisson)) ///
(Class <- child camper) ///
, ///
lclass(Class 2) lcinvariant(none)
```

With two classes, you can specify which class receives the predictors:

```
(1.Class <- child camper)
```

With more than two classes, you can specify different predictors for different classes:

```
(2.Class <- x1 x2) (3.Class <- x2 x3) ...
```

Last but not least, **gsem** allows you to specify more than one categorical latent variable:

```
gsem (x1 x2 x3 <- _cons), logit lclass(C 2) lclass(D 3)
```

In expanded notation, you get terms such as

```
(2.C#3.D: x1 <- _cons)
```

Questions?