Multivariate probit regression using simulated maximum likelihood

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Overview

- Introduction and motivation
- The model and the method of <u>Simulated</u>

 <u>Maximum Likelihood (SML)</u>
- The mvprobit program (ml, method lf)
- Illustrations mvprobit in action
- Further remarks about program use

Introduction and motivation

- Evaluation of probit model likelihood functions requires calculation of Normal probability distribution functions.
- Algorithms exist for accurately calculating accurate univariate and bivariate Normal pdfs, but not for trivariate or higher dimensional Normal distributions (at least not in Stata). Instead, ...
- Recent literature on calculating multivariate Normal pdfs using simulation-based methods
- Here: multivariate probit model estimated using simulated ML ('GHK' simulator): mvprobit
 - cf. triprobit at SSC-IDEAS

The model

M equation multivariate probit model:

off-diagonal elements.

$$y_{im}^* = \beta_m' X_{im} + \varepsilon_{im}$$
, $m = 1, ..., M$
 $y_{im} = 1$ if $y_{im}^* > 0$ and 0 otherwise
 ε_{im} , $m = 1, ..., M$, are error terms distributed as
multivariate normal, each with a mean of zero, and
variance-covariance matrix V , where V has values of 1
on the leading diagonal and correlations $\rho_{jk} = \rho_{kj}$ as

- Structure like a SUR model but depvars are binary (and need not have same set of *X* in every equation)
- *M* different choices at a point in time OR choices on one item at *M* points in time (panel model with free correlations)

Estimation principles (M = 3)

Log-likelihood function for a sample of *N* independent observations:

$$L = \sum_{i} w_{i} \log \Phi_{3}(\mu_{i}; \Omega)$$

 w_i is an optional weight

 $\Phi_3(\mu_i; \Omega)$ is standard trivariate normal cdf, where

$$\mu_i = (K_{i1}\beta_1'X_{i1}, K_{i2}\beta_2'X_{i2}, K_{i3}\beta_3'X_{i3})$$

with $K_{ik} = 2y_{ik} - 1$, for each $j, k = 1,...,3$.

$$\Omega$$
 has elements Ω_{jk} , where $\Omega_{jj} = 1$ for $j = 1, ..., 3$; $\Omega_{21} = \Omega_{12} = K_{i1}K_{i2}\rho_{21}$, $\Omega_{31} = \Omega_{13} = K_{i3}K_{i1}\rho_{31}$, $\Omega_{32} = \Omega_{23} = K_{i3}K_{i2}\rho_{32}$.

- Log-likelihood function depends on the trivariate standard normal distribution function $\Phi_3(.)!$
- Evaluate using Geweke-Hajivassiliou-Keane (GHK) smooth recursive conditioning simulator

The GHK simulator

- Exploits the fact that a multivariate normal distribution function can be expressed as the product of sequentially conditioned univariate normal distribution functions which can be easily and accurately evaluated.
- Trivariate case: 8 joint probabilities corresponding to the eight possible combinations of successes $(y_{im} = 1)$ and failures $(y_{im} = 0)$. Focus on Pr(every outcome is a success):

$$Pr(y_{1} = 1, y_{2} = 1, y_{3} = 1) = Pr(\varepsilon_{1} \le \beta_{1}'X_{1}, \varepsilon_{2} \le \beta_{2}'X_{2}, \varepsilon_{3} \le \beta_{3}'X_{3})$$

$$= Pr(\varepsilon_{3} \le \beta_{3}'X_{3} | \varepsilon_{2} < \beta_{2}'X_{2}, \varepsilon_{1} < \beta_{1}'X_{1})$$

$$\times Pr(\varepsilon_{2} < \beta_{2}'X_{2} | \varepsilon_{1} < \beta_{1}'X_{1}) \times Pr(\varepsilon_{1} < \beta_{1}'X_{1})$$

- Expression involves conditioning upon unobservable variables (that are correlated with each other).
- However if a good approximation for these conditional distributions can be found, then the likelihood function only requires evaluation of univariate integrals.
- How may the approximations be derived?

The GHK simulator (ctd.)

Cholesky decomposition of the covariance matrix for the errors:

$$E(\varepsilon \varepsilon') \equiv V = Cee'C$$

where C is the lower triangular Cholesky matrix corresponding to V and $e \sim \Phi_3(0, I_3)$, i.e. three uncorrelated standard normal variates.

Hence:

$$\varepsilon_{1} = C_{11}e_{1}
\varepsilon_{2} = C_{11}e_{1} + C_{22}e_{2}
\varepsilon_{3} = C_{31}e_{1} + C_{32}e_{2} + C_{33}e_{3}$$

and C_{jk} is the *jk*th element of matrix C.

 \Rightarrow rewrite Pr(three successes) as

$$\begin{aligned} &\Pr(\varepsilon_{1} \leq \beta_{1}'X_{1}, \varepsilon_{2} \leq \beta_{2}'X_{2}, \varepsilon_{3} \leq \beta_{3}'X_{3}) \\ &= \Pr[e_{3} \leq (\beta_{3}'X_{3} - C_{32}e_{2} - C_{31}e_{1})/C_{33} | \ e_{2} < (\beta_{2}'X_{2} - C_{21}e_{1})/C_{22}, \ e_{1} \leq \beta_{1}'X_{1}/C_{11}] \\ &\times \Pr[e_{2} \leq (\beta_{2}'X_{2} - C_{21}e_{1})/C_{22} | \ e_{1} \leq \beta_{1}'X_{1}/C_{11}] \times \Pr[e_{1} \leq \beta_{1}'X_{1}/C_{11}]. \end{aligned}$$

The standard normal variates, *e*, that now appear in the decomposition are uncorrelated with each other (by construction).

The first two conditional probabilities can be further rewritten as unconditional probabilities defined in terms of truncated standard normal variates:

The GHK simulator (ctd.)

Rewriting in terms of truncated standard normal variates:

$$\begin{split} \Pr(\mathbf{\epsilon}_{1} \leq \mathbf{\beta}_{1}'X_{1,} \mathbf{\epsilon}_{2} \leq \mathbf{\beta}_{2}'X_{2,} \mathbf{\epsilon}_{3} \leq \mathbf{\beta}_{3}'X_{3}) \\ &= \Pr[\mathbf{\epsilon}_{3} \leq (\mathbf{\beta}_{3}'X_{3} - C_{32}e_{2}^{*} - C_{31}e_{1}^{*})/C_{33}] \times \Pr[\mathbf{\epsilon}_{2} \leq (\mathbf{\beta}_{2}'X_{2} - C_{21}e_{1}^{*})/C_{22}] \\ &\qquad \times \Pr[\mathbf{\epsilon}_{1} \leq \mathbf{\beta}_{1}'X_{1}/C_{11}] \\ &= Q_{3} \times Q_{2} \times Q_{1}, \text{ say}, \end{split}$$

- where e_1^* and e_2^* are truncated univariate standard normal variates with upper truncation points at $\beta_1' X_1/C_{11}$ and $(\beta_2' X_2 C_{21} e_1^*)/C_{22}$ respectively.
- Computation of Q_1 is straightforward and, if one had some specific values for e_1^* and e_2^* , then one could also compute Q_2 and Q_3 , and hence the overall multivariate probability.
- Similar arguments for all the other probabilities of success/failure (and for M > 3).

The GHK simulator (ctd.)

GHK simulator:

- (i) Derive values for e_1^* , e_2^* via random draws from upper truncated standard normal distributions with truncation points as above (use random number generator, plus inversion formula);
- (ii) Recursively compute a multivariate probability value from the Qs.
- (iii) Replicate *R* times, and then
- (iv) Calculate the simulated probability as the arithmetic mean of the values of the simulated probabilities from each replication.
- *SML estimator*: ML with the multivariate normal probabilities calculated at each iteration using the GHK simulator ⇒ numerically intensive!
- SML estimator consistent as $N \to \infty$ and $R \to \infty$. (Asymptotic, like ML!)
- Simulation bias $\to 0$ when R raised with N. [$(R / \sqrt{N}) \to \infty$ sufficient.]

The mvprobit program

```
mvprobit equation1 equation2 ... equationM
   [weight] [if exp] [in range] [, draws(#) seed(#)
   beta0 atrho0 (matrix name) robust cluster(varname)
   constraints(numlist) level(#) maximize options ]
where each equation is specified as
   [eqname:] depvar [=] [varlist] [, noconstant] )
by may be used; all types of weights allowed; has standard features of estimation
   commands including access to estimated results; no limit on M (in principle)
Options
draws (#): number of random draws in simulation (R). Default = 5.
seed (#): initial value of random-number seed used in simulation process.
   (Default = 123456789).
beta0: estimates of the marginal probit regressions are reported.
atrho0 (matrix name): starting values for the off-diagonal elements of the
   correlation matrix V that differ from the default starting values (all zero).
Remaining options: same as the corresponding ones for biprobit.
```

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Prediction using mvppred

```
mvppred newvarname_prefix [if exp] [in range] [,
    statistic]
where statistic is one of
    xb the linear prediction for each equation; the default.
    stdp the standard error of the linear predictions for each equation.
    pmarg the marginal success probability for each equation.
```

pall the joint probabilities: (i) $Pr(y_{im} = 1, \text{ for all } m = 1,...,M)$, and (ii) $Pr(y_{im} = 0, \text{ for all } m = 1,...,M)$.

Only one statistic may be chosen at a time.

For statistics **xb**, **stdp**, and **pmarg**, results are stored in the variables newvarname_prefixi, for equations i = 1, ..., M.

For the **pall** statistics, results are stored in the variables newvarname_prefix1s for predicted probability (i) and newvarname_prefix0s for predicted probability (ii).

[Options for prediction restricted to the 'all successes' and 'all failures' cases = the only two cases that could be programmed without M being fixed. (Number of joint probabilities is 2^{M} .)]

Illustrations

- (1) (i) syntax, options etc., and (ii) accuracy of SML relative to ML (mvprobit vs. biprobit)
 - 'School' data (Pyndyck & Rubenfeld; Stata manuals), N = 95, M = 2
- (2) Simulated data, N = 5000, M = 4.

'School' data:

private: whether children attend a private school,

vote: whether the household head had voted for an increase in the property tax,

years: # years family has been at the present residence

logptax: log(property tax); loginc:log(income).

biprobit estimates:

- . use http://www.stata-press.com/data/r7/school.dta, clear
- . biprobit (private=years logptax loginc) (vote=years logptax loginc), nolog

Seemingly unrelated bivariate probit Number of obs = 95 Wald chi2(6) = 9.59 Log likelihood = -89.254028 Prob > chi2 = 0.1431

	Coef.	Std. Err.	z	P> z	[95% Conf.	Interval]	
private							
years	0118884	.0256778	-0.46	0.643	0622159	.0384391	
logptax	1066962	.6669782	-0.16	0.873	-1.413949	1.200557	
loginc	.3762037	.5306484	0.71	0.478	663848	1.416255	
cons	-4.184694	4.837817	-0.86	0.387	-13.66664	5.297253	
vote							
years	0168561	.0147834	-1.14	0.254	0458309	.0121188	
logptax	-1.288707	.5752266	-2.24	0.025	-2.416131	1612839	
loginc	.998286	.4403565	2.27	0.023	.1352031	1.861369	
_cons	5360573	4.068509	-0.13	0.895	-8.510188	7.438073	
							
/athrho	2764525	.2412099	-1.15	0.252	7492153	.1963102	
							
rho	2696186	.2236753			6346806	.1938267	
Likelihood ratio test of rho=0: $chi2(1) = 1.38444$ Prob > $chi2 = 0.2393$							

- . predict p11,p11
- . predict p00,p00
- . predict xbb1, xb1
- . predict xbb2, xb2
- . predict stdpb1, stdp1
- . predict stdpb2, stdp2
- . predict pmargb1, pmarg1
- . predict pmargb2, pmarg2

Predictions of joint & marginal probabilities, linear predictions, etc.

mvprobit estimates (R = 100) are close to ML estimates:

```
. mvprobit (private = years logptax loginc) (vote=years logptax loginc), nolog
> dr(100)
Multivariate probit (SML, # draws = 100)
                                         Number of obs =
                                                              95
                                         Wald chi2(6)
                                                              9.64
Log likelihood = -89.220805
                                         Prob > chi2
                                                            0.1405
               Coef. Std. Err. z P>|z| [95% Conf. Interval]
private
           -.0118233 .0256205 -0.46 0.644 -.0620386
     years
                                                          .0383921
    logptax | -.1033056 .6672673 -0.15 0.877 -1.411126 1.204514
    loginc
           .3695001 .5303571 0.70 0.486 -.6699806 1.408981
     cons
             -4.140149 4.837923 -0.86
                                        0.392 -13.6223 5.342006
vote
           -.0172153 .0148029 -1.16 0.245 -.0462285
     years
                                                          .011798
    logptax | -1.280732 .5725493 -2.24 0.025 -2.402908 -.1585563
    loginc | .9956743 .437904 2.27 0.023 .1373982 1.85395
                       4.055359 -0.14 0.890 -8.511157
             -.5627991
                                                          7.385558
     _cons
                                -1.17 0.241
     rho21 | -.2739382 .2216667 -1.24 0.217 -.6356397
                                                         .1863855
Likelihood ratio test of rho21 = 0:
           chi2(1) = 1.45088 Prob > chi2 = 0.2284
 . mvppred pall, pall
 (Pr(all zeros), Pr(all ones) will be stored in variables pall0s, pall1s)
 . mvppred xbm, xb
 (xb will be stored in variables xbmi, i = 1, ..., \#eqs)
                                                                Predictions
 . mvppred stdpm, stdp
 (stdp will be stored in variables stdpmi, i = 1,..., #eqs)
 . mvppred pmarqm, pmarq
                                                                             14
 (pmarg will be stored in variables pmargmi, i = 1,..., #eqs)
```

SML predictions very similar to ML counterparts

Variable	Obs	Mean	Std. Dev.	Min	Max
pall1s	95	.0513848	.0293697	.0006823	.1675037
p11	95	.0514965	.0295284	.0006783	.1691212
pall0s	95	.3252403	.1496049	.0397381	.8772917
p00	95	.3241522	.1485598	.040815	.882799
xbm1	95	-1.273431	.2017621	-1.930628	8744448
xbb1	95	-1.275218	.2041617	-1.937996	8695227
xbm2	95	.3308476	.4381363	-1.365069	1.519954
xbb2	95	.3313479	.4383708	-1.37215	1.52224
stdpm1	95	.3404043	.1805338	.185824	1.046698
stdpb1	95	.3405953	.1807651	.1859334	1.049172
stdpm2	95	.2541217	.1181758	.141525	.7976922
stdpb2	95	.2546185	.1186652	.1415696	.8023056
pmargm1	95	.1057509	.0322633	.0267645	.190938
pmargb1	95	.1055308	.032576	.0263119	.1922807
pmargm2	95	.6216642	.1502225	.0861157	.9357387
pmargb2	95	.6218135	.1500823	.0850083	.9360256

Less similarity between SML and ML with smaller *R*, we found. More on choice of *R* below.

N = 95 is 'small', so potential finite sample biases anyway. Raise N and M in simulated data example ...

Simulated data: M = 4, N = 5000

. ge y4 = y4s>0

```
. set seed 12309
. set obs 5000
obs was 0, now 5000
. matrix R = (1, .25, .5, .75 \setminus .25, 1, .75, .5 \setminus .5, .75, 1, .75 \setminus .75, .5, .7
> 5, 1)
. drawnorm u1 u2 u3 u4, corr(R)
. corr u*
                                                              Correlation structure (V)
(obs=5000)
                      u1
                              u2
                                         u3
                                                   u4
                  1.0000
          u1
          u2
                  0.2587
                           1.0000
          u3
                  0.5077
                           0.7483
                                     1.0000
          u4
                  0.7523
                           0.5093
                                     0.7589
                                              1.0000
. \text{ qe } x1 = uniform() - .5
. qe x2 = uniform() + 1/3
. qe x3 = 2*uniform() + .5
. ge x4 = .5*uniform() - 1/3
. * Equations
                                                            Equations
. qe y1s = .5 + 4*x1 + u1
. qe y2s = 3 + .5*x1 - 3*x2 + u2
. qe y3s = 1 - 2*x1 + .4*x2 - .75*x3 + u3
. qe y4s = -6 + 1*x1 - .3*x2 + 3*x3 - .4*x4 + u4
. qe y1 = y1s>0
. ge y2 = y2s>0
. qe y3 = y3s>0
```

mvprobit estimates (R = 75)

Estimates quite close to those in the 'true' model

NB warning message at iteration 1 not a problem as model later converged

LR test of MVP against *M* independent univariate probits (*V* identity matrix)

. mvprobit (y1=x1) (y2=x1 x2) (y3 = x1 x2 x3) (y4=x1 x2 x3 x4), dr(75)Iteration 0: log likelihood = -8681.8526 Warning: cannot do Cholesky factorization of rho matrix Iteration 1: log likelihood = -7922.4199 Iteration 2: log likelihood = -7749.5212 Iteration 3: log likelihood = -7746.5769 Iteration 4: log likelihood = -7746.5734 Iteration 5: log likelihood = -7746.5734 Multivariate probit (SML, # draws = 75) Number of obs = Wald chi2(10) = 5561.10 Log likelihood = -7746.5734 Prob > chi2 Coef. Std. Err. z P>|z| [95% Conf. Interval] 3.991634 .0962586 41.47 0.000 3.80297 cons .5078066 .0233257 21.77 .5413448 .0704179 7 69 0 000 -2.848419 .0781794 -36.43 0.000 -3.001648 -2.695191 cons | 2.867846 .0734467 39.05 0.000 -2.011164 .0708565 -28.38 0.000 -2.15004 -1.872288 8.32 0.000 .5341271 .0642228 .4082527 .0311735 0.000 -.7438451 -23.86 .9133876 .0725944 12.58 0.000 .7711051 1.05567 1.125271 .0891551 12.62 0.000 .9505303 -.3030878 .0826195 -3.67 0.000 x3 2.898115 .0779738 37.17 0.000 2.745289 -.7486084 -.4352364 .1598866 -2.72 0.006 cons | -5.751499 .1685475 -34.12 0.000 -6.081846 -5.421152 /atrho21 | .2621885 .0317031 8.27 0.000 .2000516 .3243253 /atrho31 | .5646645 .0345949 16.32 0.000 .4968597 /atrho41 | .9396437 .0571792 16.43 0.000 .8275746 1.051713 /atrho32 | .9737444 .0406156 23.97 0.000 .8941392 /atrho42 | .5670195 .0424459 13.36 0.000 /atrho43 | 1.007126 .0537097 18.75 0.000 .9018571 1.112395 rho21 | .2563413 .0296198 8.65 0.000 .1974249 .3134127 rho31 .5114301 .0255462 20.02 0.000 rho41 | .7350585 .0262846 27.97 0.000 .6791715 rho32 | .7503451 .0177483 42.28 0.000 .7134322 .7831052 rho42 | .513167 .0312682 16.41 0.000 .4493034 .5718126 rho43 | .7645708 .0223127 34.27 0.000 .7172009 .8049075 Likelihood ratio test of rho21 = rho31 = rho41 = rho32 = rho42 = rho43 = 0:

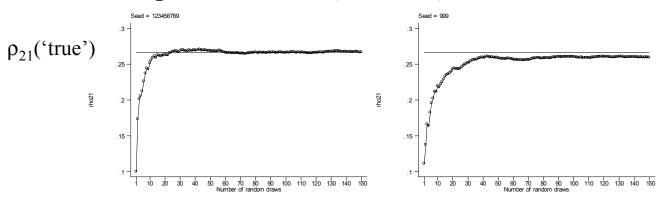
Using mvprobit: further remarks

- Choosing number of random draws: higher *R* increases accuracy but also run-time;
- Choosing seed (different random numbers give different simulated probabilities)

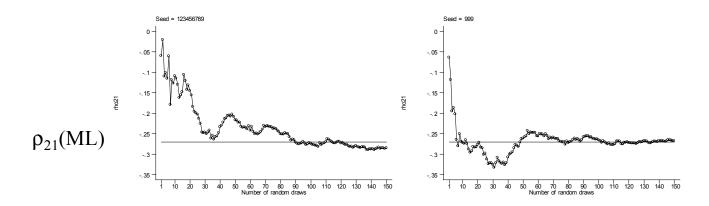
Re-estimated the earlier models for each value of R = 1, ..., 150, and several alternative seed values

SML estimate of ρ_{21} : R = 1, ..., 150; seed = 123456789, 999

(a) two-equation model, generated data (N = 5000)



(a) two-equation model, 'School' data (N = 95)



With 'large' N, SML estimator OK regardless of seed when $R > \sqrt{N}$

Using mvprobit: further remarks

Run-time:

- Four-equation model took c. 2.25 hours using Stata 7/SE on Pentium P4/1.4Ghz; c. 5.3 hours using Stata 7/IC on Sun Solaris)
- increases linearly in R (for given N, M)
- increasing number of explanatory variables, *ceteris paribus*, has no big effect
- atrho option use reduced run-time relatively little (since got good estimates of *V* within a few iterations anyway)
- Runtime increases substantially as M increases (several days/weeks if N = 1000s and M > 6 or 7!)

Capacity:

- R*M temporary variables created: set memory
- For 'large' models (many covariates, many M): set matsize