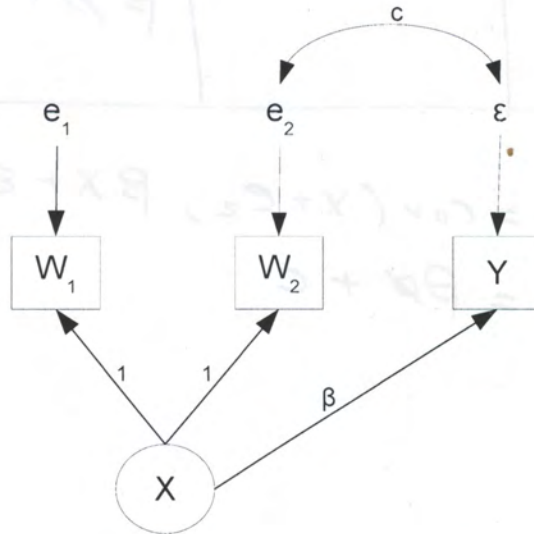


STA 431 Quiz 7

1. In this double measurement path diagram, data for W_2 and Y were collected in the same setting and at the same time. They were influenced by common omitted variables, and so their error terms are correlated.



- (a) (1 point) Write the model equations. Just write three equations and nothing else.

$$W_1 = X + e_1$$

$$W_2 = X + e_2$$

$$Y = \beta X + \epsilon$$
 (Subscripts i are optional)

- (b) (1 point) Using the notation $Var(X) = \phi$, $Var(e_1) = \omega_1$, $Var(e_2) = \omega_2$ and $Var(\epsilon) = \psi$, what is the parameter vector θ for this model?

$$\theta = (\phi, \omega_1, \omega_2, \beta, c, \psi)$$
 (Of course order does not matter)

- (c) (1 point) Does this model pass the test of the Parameter Count Rule? Answer Yes or No and give the numbers.

Yes: 6 equations in 6 unknowns

$$w_1 = X + e_1, \quad Y = \beta X + \epsilon$$

$$w_2 = X + e_2$$

(d) (3 points) Give the covariance matrix of the observable data as a function of the model parameters. Show any necessary calculations.

$\Sigma =$

	w_1	w_2	Y
w_1	$\sigma + w_1$	\emptyset	$\beta \emptyset$
w_2		$\emptyset + w_2$	$\beta \emptyset + c$
Y			$\beta^2 \emptyset + \psi$

$$\text{cov}(w_2, Y) = \text{cov}(X + e_2, \beta X + \epsilon)$$

$$= \beta \emptyset + c$$

(Not even this work is actually required.)

(e) (2 points) Is the parameter β identifiable? If Yes, show it. If No, give two parameter vectors that have *different* values of β , but produce the same covariance matrix. A simple numerical example would be best.

Yes, identifiable: $\beta = \sigma_{13} / \sigma_{12}$

2. (2 points) For the R part of the assignment, you analyzed the Pig Birth data. In a path diagram of the model (which you do *not* have to give here), there would be a curved double-headed arrow connecting the measurement error terms for Questionnaire One.

What is the value of the test statistic for testing whether the arrow is present? The answer is a number on your printout. Write the number in the space below. On your printout, circle the number and write "Question 2" beside it. *Do not answer this question if you do not have a printout for Question 3 of this week's assignment.*

$$z = 5.389$$

Please attach your printout to the quiz paper. The printout should show your *complete R input and output*. Make sure your name and student number appear on the printout.

Assignment 7, Question 3

```

> # a.
> pigs = read.table("http://www.utstat.toronto.edu/brunner/openSEM/data/openpigs2.data.txt")
> head(pigs); dim(pigs)
  nbreedersQ1 ngivebirthQ1 nbreedersQ2 ngivebirthQ2
1           69           45           89           67
2           52           24           85           41
3           48           33           68           34
4            7            4           30           30
5           41           37           54           35
6           35           23           57           48
[1] 114  4
>
> # b.
> S = var(pigs); S
      nbreedersQ1 ngivebirthQ1 nbreedersQ2 ngivebirthQ2
nbreedersQ1  691.3607   581.6943   348.5299   272.6710
ngivebirthQ1  581.6943   562.6853   260.0286   239.2279
nbreedersQ2  348.5299   260.0286   770.9685   370.8930
ngivebirthQ2  272.6710   239.2279   370.8930   334.1683
> # c. and d. are handwritten.
>
> # e.
>
> # install.packages("lavaan", dependencies = TRUE) # Only need to do this once
> library(lavaan)
This is lavaan 0.6-11
lavaan is FREE software! Please report any bugs.
>
> pigmod =
+ '#####
+ # Latent variable model
+ # -----
+ Lbirth ~ beta*Lbreeders
+ #
+ # Measurement model
+ # -----
+ Lbreeders =~ 1*nbreedersQ1 + 1*nbreedersQ2
+ Lbirth    =~ 1*ngivebirthQ1 + 1*ngivebirthQ2
+ #
+ # Variances
+ # -----
+ Lbreeders =~ phi*Lbreeders          # Var(Lbreeders) = phi
+ Lbirth    =~ psi*Lbirth             # Var(epsilon) = psi
+ nbreedersQ1 =~ omega1*nbreedersQ1 # Var(e1) = omega1
+ ngivebirthQ1 =~ omega2*ngivebirthQ1 # Var(e2) = omega2
+ nbreedersQ2 =~ omega3*nbreedersQ2 # Var(e3) = omega3
+ ngivebirthQ2 =~ omega4*ngivebirthQ2 # Var(e4) = omega4
+ #
+ # Covariances (between error terms)
+ # -----
+ nbreedersQ1 =~ omega12*ngivebirthQ1 # Cov(e1,e2) = omega12
+ nbreedersQ2 =~ omega34*ngivebirthQ2 # Cov(e3,e4) = omega34
+ # Reliabilities of number of breeding sows for the two questionnaires
+ # -----
+ rel1 := omega1/(phi+omega1)
+ rel2 := omega3/(phi+omega3)
+ '##### End of pigmod #####
>
> fit1 = lavaan(pigmod, data=pigs)
> summary(fit1)

```

lavaan 0.6-11 ended normally after 120 iterations

Estimator ML
Optimization method NLMINB
Number of model parameters 9

Number of observations 114

Model Test User Model:

Test statistic 0.087
Degrees of freedom 1
P-value (Chi-square) 0.768

Parameter Estimates:

Standard errors Standard
Information Expected
Information saturated (h1) model Structured

Latent Variables:

	Estimate	Std.Err	z-value	P(> z)
Lbreeders =~				
nbreedersQ1	1.000			
nbreedersQ2	1.000			
Lbirth =~				
ngivebirthQ1	1.000			
ngivebirthQ2	1.000			

Regressions:

	Estimate	Std.Err	z-value	P(> z)
Lbirth ~				
Lbredrs (beta)	0.757	0.054	14.047	0.000

Covariances:

	Estimate	Std.Err	z-value	P(> z)
.nbreedersQ1 ~~				
.ngvbrQ1 (om12)	308.539	57.249	5.389	0.000
.nbreedersQ2 ~~				
.ngvbrQ2 (om34)	101.406	45.513	2.228	0.026

Question 2

Variances:

	Estimate	Std.Err	z-value	P(> z)
Lbredrs (phi)	357.145	64.936	5.500	0.000
.Lbirth (psi)	33.634	10.861	3.097	0.002
.nbrdrQ1 (omg1)	330.683	67.114	4.927	0.000
.ngvbrQ1 (omg2)	321.255	54.160	5.932	0.000
.nbrdrQ2 (omg3)	416.335	80.763	5.155	0.000
.ngvbrQ2 (omg4)	93.000	35.566	2.615	0.009

Defined Parameters:

	Estimate	Std.Err	z-value	P(> z)
rel1	0.481	0.070	6.877	0.000
rel2	0.538	0.070	7.706	0.000

```
>
> # h.
>
> # MOM of beta: Really need to calculate Sigma to see this.
> 0.5*(S[1,4]+S[2,3])/S[1,3]
[1] 0.7642093
```

```

>
> # i.
> parameterEstimates(fit1)
      lhs op          rhs  label  est   se    z  pvalue  ci.lower  ci.upper
1     Lbirth ~      Lbreeders  beta  0.757 0.054 14.047 0.000    0.651    0.862
2     Lbreeders =~  nbreedersQ1  1.000 0.000    NA    NA    1.000    1.000
3     Lbreeders =~  nbreedersQ2  1.000 0.000    NA    NA    1.000    1.000
4     Lbirth =~    ngivebirthQ1  1.000 0.000    NA    NA    1.000    1.000
5     Lbirth =~    ngivebirthQ2  1.000 0.000    NA    NA    1.000    1.000
6     Lbreeders ~~      Lbreeders  phi 357.145 64.936  5.500 0.000 229.873 484.417
7     Lbirth  ~~      Lbirth    psi  33.634 10.861  3.097 0.002  12.348  54.920
8     nbreedersQ1 ~~  nbreedersQ1  omega1 330.683 67.114  4.927 0.000 199.143 462.224
9     ngivebirthQ1 ~~  ngivebirthQ1  omega2 321.255 54.160  5.932 0.000 215.103 427.407
10    nbreedersQ2 ~~  nbreedersQ2  omega3 416.335 80.763  5.155 0.000 258.042 574.628
11    ngivebirthQ2 ~~  ngivebirthQ2  omega4  93.000 35.566  2.615 0.009  23.291 162.709
12    nbreedersQ1 ~~  ngivebirthQ1  omega12 308.539 57.249  5.389 0.000 196.334 420.744
13    nbreedersQ2 ~~  ngivebirthQ2  omega34 101.406 45.513  2.228 0.026  12.201 190.610
14      rel1 := omega1/(phi+omega1)  rel1  0.481 0.070  6.877 0.000    0.344    0.618
15      rel2 := omega3/(phi+omega3)  rel2  0.538 0.070  7.706 0.000    0.401    0.675

```

```

> # j. k. and L. See above.
>
> # m. Compare precision of measurement.
> # H0: omega1=omega3 and omega2=omega4
> # For Wald tests: Wtest = function(L,Tn,Vn,h=0) # H0: L theta = h
> source("http://www.utstat.utoronto.ca/~brunner/Rfunctions/Wtest.txt")
>
> # Fit model with bootstrap SEs
> fit2 = lavaan(pigmod, data=pigs, se = "bootstrap")
> summary(fit2) # I didn't ask for this explicitly, so no questions on it.
lavaan 0.6-11 ended normally after 120 iterations

```

```

Estimator          ML
Optimization method NLMINB
Number of model parameters 9

Number of observations 114

```

Model Test User Model:

```

Test statistic          0.087
Degrees of freedom      1
P-value (Chi-square)    0.768

```

Parameter Estimates:

```

Standard errors          Bootstrap
Number of requested bootstrap draws 1000
Number of successful bootstrap draws 1000

```

Latent Variables:

```

      Estimate Std.Err z-value P(>|z|)
Lbreeders =~
  nbreedersQ1  1.000
  nbreedersQ2  1.000
Lbirth =~
  ngivebirthQ1 1.000
  ngivebirthQ2 1.000

```

Regressions:

	Estimate	Std.Err	z-value	P(> z)
Lbirth ~ Lbredrs (beta)	0.757	0.071	10.721	0.000

Covariances:

	Estimate	Std.Err	z-value	P(> z)
.nbreedersQ1 ~ .ngvbrQ1 (om12)	308.539	54.937	5.616	0.000
.nbreedersQ2 ~ .ngvbrQ2 (om34)	101.406	37.255	2.722	0.006

Variances:

	Estimate	Std.Err	z-value	P(> z)
Lbredrs (phi)	357.145	81.988	4.356	0.000
.Lbirth (psi)	33.634	11.319	2.972	0.003
.nbrdrQ1 (omg1)	330.683	60.901	5.430	0.000
.ngvbrQ1 (omg2)	321.255	52.548	6.114	0.000
.nbrdrQ2 (omg3)	416.335	86.653	4.805	0.000
.ngvbrQ2 (omg4)	93.000	30.053	3.095	0.002

Defined Parameters:

	Estimate	Std.Err	z-value	P(> z)
rel1	0.481	0.085	5.686	0.000
rel2	0.538	0.082	6.537	0.000

```

>
> # Wald test of H0: omegal=omega3 and omega2=omega4
> That = coef(fit2); That
  beta    phi    psi  omegal  omega2  omega3  omega4  omegal2  omega34
0.757 357.145 33.634 330.683 321.255 416.335 93.000 308.539 101.406
> VV = vcov(fit2)
> LL = rbind(c(0, 0, 0, 1, 0,-1, 0, 0, 0),
+           c(0, 0, 0, 0, 1, 0,-1, 0, 0))
> Wtest(LL,That,VV)
      W      df      p-value
2.393441e+01 2.000000e+00 6.349046e-06
>
> # Follow-up
> # omegal vs. omega3
> L13 = rbind(c(0, 0, 0, 1, 0,-1, 0, 0, 0))
> Wtest(L13,That,VV)
      W      df      p-value
0.7262899 1.0000000 0.3940884
> # omega2 vs. omega4
> L24 = rbind(c(0, 0, 0, 0, 1, 0,-1, 0, 0))
> Wtest(L24,That,VV)
      W      df      p-value
1.338262e+01 1.000000e+00 2.539661e-04
>

```