## Preliminaries: data generating process

The first part of these section notes is a variation on Max Auffhammer's final exam for ARE212 in 2011. If you don't like this treatment of IV estimation, then blame him. Actually, I like this pattern: if you don't like something I do, then blame Max. He is probably at fault. I am faultless. Consider the following data generating process:

$$y_i = \beta_0 + \beta_1 x_{1i} + \beta_2 x_{2i} + \beta_3 x_{3i} + \eta_i \quad \text{with} \quad \eta_i \sim N(0, 1)$$

and  $\beta = [1 \ 2 \ -4 \ 1]'$ . Assume that the covariance matrix of the covariates, an additional instrument, and the idiosyncratic error  $(x_{1i}, x_{2i}, x_{3i}, z_i, \text{ and } \eta_i)$  is defined to be

$$\Sigma = \begin{bmatrix} 1 & 0 & \rho_{13} & 0 & 0 \\ 0 & 1 & \rho_{23} & \rho_{2z} & 0 \\ \rho_{13} & \rho_{23} & 1 & 0 & 0 \\ 0 & \rho_{2z} & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 1 \end{bmatrix}$$

where each variable is assumed to have zero mean for simplicity. Note that  $\Sigma$  is consistent with  $\eta_i$  being independently and identically distributed with constant variance. We will explore the properties of IV (e.g., weak instruments and the exclusion restriction) via Monte Carlo simulation.

The first step, then, is to figure out how to generate random data with the appropriate covariance. For this, a useful function is rmvn.chol, which returns a random  $n \times k$  multivariate normal matrix  $\mathbf{X}$ , based on the supplied covariance matrix vcov.mat. In general,  $\mathbf{X} = \mathbf{Q} + \mu'$ , where  $\mathbf{Q}'\mathbf{Q} = \Sigma$  and  $\mu$  is a vector of means for each of the k columns of  $\mathbf{X}$ . We have assumed away  $\mu = \mathbf{0}$  and so the problem becomes a simple application of matrix decomposition.

```
rmvn.chol <- function(n, vcov.mat) {
  k <- ncol(vcov.mat)
  Q <- chol(vcov.mat)
  Z <- matrix(rnorm(k*n), nrow=n, ncol=k)
  return(Z %*% Q)
}</pre>
```

It will be handy to have a simple function to generate  $\Sigma$  with three arguments representing the three non-zero correlations across the covariates and the instrument.

```
vcov.fn <- function(rho.13, rho.23, rho.2z) {
  mat <- diag(5)
  mat[3,1] <- rho.13; mat[1,3] <- rho.13
  mat[2,3] <- rho.23; mat[3,2] <- rho.23
  mat[2,4] <- rho.2z; mat[4,2] <- rho.2z
  return(mat)
}</pre>
```

The result is a way to generate the random data according to the specified data generating process. The following generates the covariance matrix and a random multivariate normal matrix with 500 observations, printing  $\Sigma$  for reference:

```
(vcov <- vcov.fn(0, 0.5, 0.5))
X <- rmvn.chol(500, vcov)
```

```
[,1] [,2] [,3] [,4] [,5]
[1,]
           0.0 0.0
[2,]
                               0
            1.0
                 0.5
[3,]
                               0
[4,]
            0.5
                      1.0
                               0
                 0.0
[5,]
                 0.0
                               1
```

A quick check to ensure that the variance of each variable is 1, as specified by  $\Sigma$ . There will be some variation around the true variance, but even with 500 observations, it's clear that we are approaching the true variance:

```
apply(X, 2, function(i){var(i)})
[1] 0.9318116 0.9494840 1.1271233 0.9727111 1.0298838
```

## Calculate bias in $\beta$ and it's standard error

We will now write a couple of functions to help examine the bias of the parameter vector and it's standard error, using an array of estimation techniques, including OLS, 2SLS, and estimation by proxy variable. First, let's put together a very simple function to calculate the parameter vector and it's standard error for both direct regression and two-stage least squares, when a first-stage covariate matrix is provided.

```
ols.results <- function(y, X, first=FALSE) {
   Xt <- t(X)
   xtxi <- solve(Xt %*% X)
   beta <- xtxi %*% Xt %*% y
   if (first == FALSE) {
      e <- y - X %*% beta
   }
   else {
      e <- y - first %*% beta
   }
   s2 <- (t(e) %*% e)/(nrow(X) - ncol(X))
   se <- sqrt(diag(xtxi)*s2)
   return(cbind(beta, se))
}</pre>
```

Now comes the serious stuff, specifically, the code that is written specifically to examine IV estimation in this example. Suppose that we do not observe  $x_{3i}$ . The composite error is then  $x_{3i} + \eta_i$ , and we estimate the parameter vector by regressing  $y_i$  on  $x_{1i}$  and  $x_{2i}$ . If  $\rho_{13} = \rho_{23} = 0$ , then there is no problem: OLS will yield consistent estimates, since the regression utilizes only exogenous variation. If, however, the covariates are correlated with the composite error, the OLS estimates will be biased.

The function est.bias returns the simulated bias in the parameter estimates and standard errors from a Monte Carlo simulation with B=10,000 repetitions. The arguments are vcov which is the variance-covariance matrix generated by vcov.fn; n which specifies the number of observations for each iteration, defaulted at 500; B is the number of iterations in the MC simulation, defaulted at 10,000; two.stage is a boolean argument indicating whether the simulation should employ two-stage least squares with  $z_i$  as the instrument for  $x_{3i}$ , defaulted to FALSE; proxy is a boolean indicating whether  $z_i$  should be treated as a proxy for  $x_{3i}$ , defaulted to FALSE. The default behavior, then, is to run a simulation where  $x_{3i}$  is left out of the OLS regression, relegated to the error term.

```
est.bias <- function(vcov, n=500, B=1000, two.stage=FALSE, proxy=FALSE) { true.beta <- c(1, 2, -4, 1)
```

```
res.beta <- mat.or.vec(3,B); res.se <- mat.or.vec(3,B)
  for (i in 1:B) {
    data <- rmvn.chol(n, vcov)</pre>
    X <- cbind(1, data[,1:3]); eta <- data[,5]</pre>
    y <- X %*% true.beta + eta
    full.ols <- ols.results(y, X)</pre>
    if (two.stage==TRUE) {
       endog <- data[,2]</pre>
       first \leftarrow cbind(1, data[,c(1,4)])
      predict <- first \%*\% solve(t(first) \%*\% first) \%*\% t(first) \%*\% endog
       exog <- cbind(1, data[,1], predict)</pre>
       limited.ols <- ols.results(y, exog, first=first)</pre>
    else if (proxy==TRUE) {
       exog <- cbind(1, data[,c(1, 4)])
       limited.ols <- ols.results(y, exog)</pre>
    }
    else {
       exog <- cbind(1, data[,1:2])
      limited.ols <- ols.results(y, exog)</pre>
    }
    res.beta[,i] <- limited.ols[,1] - true.beta[1:3]</pre>
    res.se[,i] <- limited.ols[,2] - full.ols[1:3,2]</pre>
  results <- cbind(rowMeans(res.beta), rowMeans(res.se))</pre>
  colnames(results) <- c("beta bias", "se bias")</pre>
  print(results)
We can check est.bias by first setting \rho_{13} = \rho_{23} = 0 and ensuring that the bias is very low with n = 500
and B = 10,000. The following MC simulation sets \Sigma = \mathbb{I}_5 and runs the following regression 10,000 times:
y_i = \beta_0 + \beta_1 x_{1i} + \beta_2 x_{2i} + \epsilon_i, where \epsilon_i = x_{3i} + \eta_i.
vcov <- vcov.fn(0, 0, 0)
est.bias(vcov)
         beta bias
                      se bias
[1,] 3.025788e-05 0.01850461
[2,] 7.504312e-05 0.01853065
[3,] -2.155886e-04 0.01853233
vcov < - vcov.fn(0, 0.5, 0.5)
est.bias(vcov)
         beta bias
                        se bias
[1,] -9.313589e-05 0.014414161
[2,] -3.125962e-04 0.014437845
[3,] 5.000045e-01 0.007484851
est.bias(vcov, two.stage=TRUE)
```

```
beta bias se bias
[1,] 0.0006683202 0.1239639
[2,] -0.0005169007 0.1240761
[3,] -0.0042907956 0.2877558
```

We can prove that, in general, the variance of the IV estimator is larger than the variance of an OLS estimator. Let be a matrix of instruments and  $\mathbf{P} =$ . Then the IV estimator is  $\hat{\beta}_{iv} = (\mathbf{X}'\mathbf{P}\mathbf{X})^{-1}\mathbf{X}'\mathbf{P}\mathbf{y}$ . The projection matrix  $\mathbf{P}$  is symmetric and idempotent, as is the residual maker  $(\mathbb{I}_N - \mathbf{P})$ . Note that for any  $\mathbf{X}$  and any symmetric, idempotent matrix, the following fact is true:

$$X'PX = X'PPX = X'PX = (PX)'(PX) \ge 0$$
(2)

Note also that  $\mathbb{V}(\hat{\beta}_{ols}) = (\mathbf{X}'\mathbf{X})^{-1}$  and  $\mathbb{V}(\hat{\beta}_{iv}) = (\mathbf{X}'\mathbf{P}\mathbf{X})^{-1}$ . It suffices to show that  $\mathbf{X}'\mathbf{X} \geq \mathbf{X}'\mathbf{P}\mathbf{X}$  to complete the proof. It follows from Equation (2) that:

$$\mathbf{X}'(\mathbb{I}_N - \mathbf{P})\mathbf{X} \ge 0 \Rightarrow \mathbf{X}'\mathbf{X} - \mathbf{X}'\mathbf{P}\mathbf{X} \ge 0 \Rightarrow \mathbf{X}'\mathbf{X} \ge \mathbf{X}'\mathbf{P}\mathbf{X},$$

thus proving that  $\mathbb{V}(\hat{\beta}_{iv}) \geq \mathbb{V}(\hat{\beta}_{ols})$ . The intuition is a bit more clear. The IV estimator by design restricts the use of covariation between  $\mathbf{y}$  and  $\mathbf{X}$  to *only* exogenous variation and therefore greater variance of the estimator.

```
vcov <- vcov.fn(0, 0.5, 0.5)
est.bias(vcov, proxy=TRUE)
        beta bias
                     se bias
[1,] -0.0000945208 0.09679044
[2,] 0.0002856399 0.09692606
[3,] 2.0001688753 0.08998506
vcov <- vcov.fn(0.5, 0.5, 0.5)
est.bias(vcov)
        beta bias
                        se bias
[1,] -0.0006474492 9.996362e-03
[2,] 0.5005109401 -1.006042e-04
[3,] 0.5004461721 -8.279249e-05
est.bias(vcov, two.stage=TRUE)
        beta bias
                    se bias
[1,] 0.0005228898 0.1224915
[2,] 0.4996544539 0.1125726
[3,] -0.0030798629 0.2818936
vcov <- vcov.fn(0, 0.5, 0.01)
est.bias(vcov)
        beta bias
                      se bias
[1,] -5.845659e-06 0.014421682
[2,] 9.731249e-05 0.014443940
[3,] 5.000803e-01 0.007497558
est.bias(vcov, two.stage=TRUE)
      beta bias
                   se bias
[1,] 0.18013841
                  982.6684
[2,] -0.02440537
                  745.3372
[3,] 2.06182132 13433.0801
```