

Homework #3

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Reweighting

Matching

- (a) To run OLS on $Y_{i2} - Y_{i1} = \gamma_2 - \gamma_1 + \beta D_{i2} + \epsilon_{i2} - \epsilon_{i1}$, the identifying restriction is that $E[D_{i2}(\epsilon_{i2} - \epsilon_{i1})] = 0$; that is, program participation is uncorrelated with the change in error terms $\epsilon_{i2} - \epsilon_{i1}$.
- (b) $Y_{i2} = Y_{i1} + (\gamma_2 - \gamma_1) + \beta D_{i2} + \epsilon_{i2} - \epsilon_{i1}$
- (c) Taking expectations,

$$E[Y_{i2}|Y_{i1}, D_{i2} = 1] = Y_{i1} + (\gamma_2 - \gamma_1) + \beta + E[\epsilon_{i2} - \epsilon_{i1}|Y_{i1}, D_{i1} = 1]$$

$$E[Y_{i2}|Y_{i1}, D_{i2} = 0] = Y_{i1} + (\gamma_2 - \gamma_1) + E[\epsilon_{i2} - \epsilon_{i1}|Y_{i1}, D_{i1} = 0]$$

the difference of these two identifies β if

$$E[\epsilon_{i2} - \epsilon_{i1}|Y_{i1}, D_{i2} = 1] - E[\epsilon_{i2} - \epsilon_{i1}|Y_{i1}, D_{i2} = 0] = 0$$

$$E\left[E[\epsilon_{i2} - \epsilon_{i1}|D_{i2} = 1] - E[\epsilon_{i2} - \epsilon_{i1}|D_{i2} = 0] \middle| Y_{i1}\right] = 0$$

$$E\left[D_{i2}(\epsilon_{i2} - \epsilon_{i1}) \middle| Y_{i1}\right] = 0$$

- (d) This assumption is stricter than that of the assumption in part (a); it is the same restriction in part (a), but conditional on Y_{i1} .
- (e) The Smith and Todd (2005) matched differences-in-differences estimator requires that selection is independent on the differences in untreated outcomes between time periods, conditional on the propensity. Formally,

$$E\left[D_{i2}(\gamma_2 + \epsilon_{i2} - \gamma_1 - \epsilon_{i1}) \middle| P_i\right] = E[D_{i2}]E\left[(\gamma_2 + \epsilon_{i2} - \gamma_1 - \epsilon_{i1}) \middle| P_i\right]$$

$$\implies E\left[D_{i2}(\epsilon_{i2} - \epsilon_{i1}) \middle| P_i\right] = 0$$

which is the same condition from above, except instead of conditioning on Y_{i1} we are conditioning on P_i , the conditions are not really “weaker” than the ones above.

2SLS

- (a) The moment conditions defining the 2SLS estimator are

$$\begin{aligned} E[Z_i \epsilon_i] &= 0 \\ E[Z_i^2 \epsilon_i] &= 0 \end{aligned}$$

they imply each other, since functions of uncorrelated variables are uncorrelated with each other.

- (b) The moment conditions defining the control function estimator are

$$\begin{aligned} E[Z_i \epsilon_i] &= 0 \\ E[Z_i v_i] &= 0 \\ E[\epsilon_i | v_i] &= \rho \frac{\sigma_\epsilon}{\sigma_v} v_i \\ E\left[(Y_i - \beta_0 - \beta_1 X_i - \beta_2 X_i^2 - \rho \frac{\sigma_\epsilon}{\sigma_v} v_i) X_i\right] &= 0 \\ E\left[(Y_i - \beta_0 - \beta_1 X_i - \beta_2 X_i^2 - \rho \frac{\sigma_\epsilon}{\sigma_v} v_i) X_i^2\right] &= 0 \\ E\left[(Y_i - \beta_0 - \beta_1 X_i - \beta_2 X_i^2 - \rho \frac{\sigma_\epsilon}{\sigma_v} v_i) v_i\right] &= 0 \end{aligned}$$

- (c) The control function estimator relies on stronger conditions than the 2SLS estimator. The only condition in the 2SLS estimator is that the instrument, Z , is uncorrelated with the error term ϵ . However, the control function estimator relies on specifying the exogenous relationship between X and Z , in particular that the conditional expectation of X given Z is linear.
- (d) The control function would work better when Z is weak, because it brings to bear more moment restrictions than the 2SLS estimator, whereas the 2SLS estimator only relies on the exogeneity of Z .
- (e) The table below contains the estimates for β_1 and β_2 under the 2SLS and control function methods.
- (i) Overall, for all values of γ , the control function estimate appears to be more efficient (the spread of the estimates is much lower), because the control function's "first-stage" is correctly specified, and the estimation involves more moment restrictions.
 - (ii) As γ_1 shrinks, the mean 2sls estimate falls (away from the true value of 1.0), although the median appears to remain unbiased. The interquartile range of both estimators increases, although the IQR for the 2sls estimate increases much faster than that of the control function. The control function estimates appear to remain both median and mean-unbiased.
 - (iii) If one wanted to test the hypothesis that $\rho \frac{\sigma_\epsilon}{\sigma_v} = 0$, one would need an additional term in the variance of the coefficient on v_i . In particular, by using the two-stage estimate formulas for the asymptotic variance of the estimate of $\rho \frac{\sigma_\epsilon}{\sigma_v}$, one would

see that in the “sandwich” estimator, one would replace the term

$$E[s_i(\theta_0, \gamma^*)s_i(\theta_0, \gamma^*)']$$

where $\theta_0 = (\beta_0, \beta_1, \beta_2, \rho_{\sigma_v^2}^{\sigma_\epsilon})$ and $\gamma^* = (\gamma_0, \gamma_1)$, and s_i being the score function (gradient of the squared residual with respect to θ_0), with this term

$$E[g_i(\theta_0, \gamma^*)g_i(\theta_0, \gamma^*)']$$

s. t. $g_i(\theta_0, \gamma^*) \equiv s_i(\theta_0, \gamma^*) + E[\nabla_\gamma s_i(\theta_0, \gamma^*)] r_i(\theta_0, \gamma^*)$

where r_i is the asymptotic variance of $\sqrt{N}(\hat{\gamma} - \gamma^*)$ and can be obtained from the “first-stage” regression. To construct an estimate of the standard error, one can simply plug in the population analogues into this sandwich estimator and run a t-test with the modified standard error.

Statistic	β_1^{2sls}	β_2^{2sls}	β_1^{cf}	β_2^{cf}
$\gamma = 0.3$				
mean	1.003234	1.076318	.9938653	1.000297
sd	.3412886	3.060352	.1081107	.0204986
p50	.9992083	.9904965	.9961268	1.000156
iqr	.1585197	.3704991	.1449331	.0277689
$\gamma = 0.2$				
mean	.7418318	2.253272	.9936412	.999546
sd	8.550667	47.84852	.1688314	.0207697
p50	.9829823	1.005342	.9991214	.9994298
iqr	.315193	.7726767	.2297587	.0277564
$\gamma = 0.1$				
mean	.8100506	.839673	.9693889	.9988507
sd	7.576605	17.05286	.5007757	.0224328
p50	1.029302	1.044092	1.005126	.9992241
iqr	.7045021	1.448808	.4600121	.0299107

Bootstrap OLS

- See the regression results in table below. The clustered standard error on D is 0.0925.
- The p value for the null hypothesis that the coefficient on D equals zero is 0.008.
- See column 2 of the table below. Using a block bootstrap our new standard error on D is 0.0965 and the p value is 0.003.
- The p value of the symmetric percentile t-test is 0.005.
- Using the Wild bootstrap with Rademacher weights, the p value of the null hypothesis is 0.009.
- Since we have only have 20 clusters, we need to use a bootstrap to correct the standard errors. When the limiting distribution is normal a regular bootstrap will give us correct standard errors (our answer in b). The refinement we did in part c, will give us a slightly better estimate. However, the most correct standard errors when we do not

Table 1: Bootstrap

	(1) Standard Reg	(2) Bootstrap
D	0.276*** (0.09248)	0.276*** (0.09651)
X	0.147 (0.11505)	0.147 (0.12219)
X2	0.216*** (0.02678)	0.216*** (0.02602)
Constant	0.392*** (0.09661)	0.392*** (0.10193)
R^2	0.381	0.381
Observations	200	200

Standard errors in parentheses

* p<0.10, ** p<0.05, *** p<0.01

have a normal limiting distribution or are concerned about heteroskedasticity is the wild bootstrap imposing the null used in CGM.

Bootstrap Probit See attached Stata code.

Table 2: Comparison of p-Values on Null Hypothesis of $D = 0$ from Various Tests

	Cluster Robust Wald Test	Symmetric Bootstrap Percentile t-test	Cluster Robust Score Test	Clustered Score Bootstrap
p-value:	0.005	0.003	0.005	0.018