

```

* Propensity Score Matching in Stata
* Copyright 2013 by Ani Katchova

clear all
set more off

* Download and install Stata ado files for pscore
* net install st0026_2

use C:\Econometrics\Data\matching_earnings

* Define treatment, outcome, and independent variables
global treatment TREAT
global ylist RE78
global xlist AGE EDUC MARR
global breps 5

* For difference-in-differences, outcome is the differences in outcomes after and
before
* global ylist REDIFF

describe $treatment $ylist $xlist
summarize $treatment $ylist $xlist

bysort $treatment: summarize $ylist $xlist

* Regression with a dummy variable for treatment (t-test)
reg $ylist $treatment

* Regression with a dummy variable for treatment controlling for x
reg $ylist $treatment $xlist

* Propensity score matching with common support
pscore $treatment $xlist, pscore(myscore) blockid(myblock) comsup

* Matching methods

* Nearest neighbor matching
attnd $ylist $treatment $xlist, pscore(myscore) comsup boot reps($breps) dots

* Radius matching
attr $ylist $treatment $xlist, pscore(myscore) comsup boot reps($breps) dots
radius(0.1)

* Kernel Matching
atrk $ylist $treatment $xlist, pscore(myscore) comsup boot reps($breps) dots

* Stratification Matching
atts $ylist $treatment $xlist, pscore(myscore) blockid(myblock) comsup boot
reps($breps) dots

```

```

. * Propensity Score Matching in Stata
. * Copyright 2013 by Ani Katchova
.
. clear all

. set more off

.
. * Download and install Stata ado files for pscore
. * net install st0026_2
.
. use C:\Econometrics\Data\matching_earnings

.
. * Define treatment, outcome, and independent variables
. global treatment TREAT

. global ylist RE78

. global xlist AGE EDUC MARR

. global breps 5

.
. * For difference-in-differences, outcome is the differences in outcomes after and
before
. * global ylist REDIFF
.
. describe $treatment $ylist $xlist

```

variable name	storage type	display format	value label	variable label
TREAT	float	%9.0g		
RE78	float	%9.0g		
AGE	float	%9.0g		
EDUC	float	%9.0g		
MARR	float	%9.0g		

```

. summarize $treatment $ylist $xlist

```

Variable	Obs	Mean	Std. Dev.	Min	Max
TREAT	2675	.0691589	.2537716	0	1
RE78	2675	20502.38	15632.52	0	121174
AGE	2675	34.22579	10.49984	17	55
EDUC	2675	11.99439	3.053556	0	17
MARR	2675	.8194393	.3847257	0	1

```

.
. bysort $treatment: summarize $ylist $xlist

```

-> TREAT = 0

Variable	Obs	Mean	Std. Dev.	Min	Max
RE78	2490	21553.92	15555.35	0	121174
AGE	2490	34.8506	10.44076	18	55
EDUC	2490	12.11687	3.082435	0	17
MARR	2490	.8662651	.3404357	0	1

-> TREAT = 1

Variable	Obs	Mean	Std. Dev.	Min	Max
RE78	185	6349.145	7867.405	0	60307.9
AGE	185	25.81622	7.155019	17	48
EDUC	185	10.34595	2.01065	4	16
MARR	185	.1891892	.3927217	0	1

.
 . * Regression with a dummy variable for treatment (t-test)
 . reg \$ylist \$treatment

Source	SS	df	MS	Number of obs =	2675
Model	3.9811e+10	1	3.9811e+10	F(1, 2673) =	173.41
Residual	6.1365e+11	2673	229573201	Prob > F =	0.0000
				R-squared =	0.0609
				Adj R-squared =	0.0606
Total	6.5346e+11	2674	244375675	Root MSE =	15152

RE78	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]
TREAT	-15204.78	1154.614	-13.17	0.000	-17468.8 -12940.75
_cons	21553.92	303.6414	70.98	0.000	20958.53 22149.32

.
 . * Regression with a dummy variable for treatment controlling for x
 . reg \$ylist \$treatment \$xlist

Source	SS	df	MS	Number of obs =	2675
Model	1.3388e+11	4	3.3470e+10	F(4, 2670) =	171.99
Residual	5.1958e+11	2670	194600040	Prob > F =	0.0000
				R-squared =	0.2049
				Adj R-squared =	0.2037
Total	6.5346e+11	2674	244375675	Root MSE =	13950

RE78	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]
TREAT	-6901.379	1213.245	-5.69	0.000	-9280.375 -4522.384
AGE	176.6396	27.60029	6.40	0.000	122.5195 230.7597

EDUC		1900.046	91.77385	20.70	0.000	1720.091	2080.001
MARR		4937.016	800.0326	6.17	0.000	3368.269	6505.762
_cons		-11901.44	1703.792	-6.99	0.000	-15242.33	-8560.557

```

.
. * Propensity score matching with common support
. pscore $treatment $xlist, pscore(myscore) blockid(myblock) comsup

```

```

*****
Algorithm to estimate the propensity score
*****

```

The treatment is TREAT

TREAT		Freq.	Percent	Cum.
0		2,490	93.08	93.08
1		185	6.92	100.00
Total		2,675	100.00	

Estimation of the propensity score

```

Iteration 0:  log likelihood = -672.64954
Iteration 1:  log likelihood = -445.24322
Iteration 2:  log likelihood = -417.65611
Iteration 3:  log likelihood = -414.43392
Iteration 4:  log likelihood = -414.37065
Iteration 5:  log likelihood = -414.37062

```

Probit regression	Number of obs	=	2675
	LR chi2(3)	=	516.56
	Prob > chi2	=	0.0000
Log likelihood = -414.37062	Pseudo R2	=	0.3840

TREAT		Coef.	Std. Err.	z	P> z	[95% Conf. Interval]
AGE		-.0479858	.0060942	-7.87	0.000	-.0599302 - .0360415
EDUC		-.1686459	.0192597	-8.76	0.000	-.2063942 - .1308977
MARR		-1.469566	.0995124	-14.77	0.000	-1.664607 -1.274526
_cons		2.729172	.3230556	8.45	0.000	2.095994 3.362349

Note: the common support option has been selected
The region of common support is [.000914, .80332316]

Description of the estimated propensity score
in region of common support

Estimated propensity score				

	Percentiles	Smallest		
1%	.0010694	.000914		
5%	.0014751	.000914		
10%	.0020168	.000914	Obs	2332
25%	.0048323	.0009183	Sum of Wgt.	2332
50%	.0177077		Mean	.0796709
		Largest	Std. Dev.	.144896
75%	.0534804	.7537369		
90%	.3188462	.7681876	Variance	.0209948
95%	.4279398	.7897533	Skewness	2.41066
99%	.645166	.8033232	Kurtosis	8.272405

Step 1: Identification of the optimal number of blocks
Use option detail if you want more detailed output

The final number of blocks is 10

This number of blocks ensures that the mean propensity score
is not different for treated and controls in each blocks

Step 2: Test of balancing property of the propensity score
Use option detail if you want more detailed output

The balancing property is satisfied

This table shows the inferior bound, the number of treated
and the number of controls for each block

Inferior	TREAT		
of block	0	1	Total
of pscore			
-----+-----	-----	-----	-----
.000914	671	1	672
.00625	316	6	322
.0125	379	8	387
.025	341	14	355

.05		135	8		143
.1		110	12		122
.2		137	61		198
.4		51	51		102
.6		7	23		30
.8		0	1		1
-----+-----+-----					
Total		2,147	185		2,332

Note: the common support option has been selected

End of the algorithm to estimate the pscore

```
.
. * Matching methods
.
. * Nearest neighbor matching
. attnd $ylist $treatment $xlist, pscore(myscore) comsup boot reps($breps) dots
```

The program is searching the nearest neighbor of each treated unit.
This operation may take a while.

ATT estimation with Nearest Neighbor Matching method
(random draw version)
Analytical standard errors

n. treat.	n. contr.	ATT	Std. Err.	t
185	431	-6715.251	1200.953	-5.592

Note: the numbers of treated and controls refer to actual
nearest neighbour matches

Bootstrapping of standard errors

```
command:      attnd RE78 TREAT AGE EDUC MARR , pscore(myscore) comsup
statistic:    attnd      = r(attnd)
.....
```

Bootstrap statistics	Number of obs	=	2675
	Replications	=	5

Variable	Reps	Observed	Bias	Std. Err.	[95% Conf. Interval]		
attnd	5	-6715.251	329.3118	917.7546	-9263.347	-4167.156	(N)
					-7521.834	-5499.719	(P)
					-7521.834	-5499.719	(BC)

Note: N = normal
P = percentile
BC = bias-corrected

ATT estimation with Nearest Neighbor Matching method
(random draw version)
Bootstrapped standard errors

n. treat.	n. contr.	ATT	Std. Err.	t
185	431	-6715.251	917.755	-7.317

Note: the numbers of treated and controls refer to actual nearest neighbour matches

```
.
. * Radius matching
. attr $ylist $treatment $xlist, pscore(myscore) comsup boot reps($breps) dots
radius(0.1)
```

The program is searching for matches of treated units within radius.
This operation may take a while.

ATT estimation with the Radius Matching method
Analytical standard errors

n. treat.	n. contr.	ATT	Std. Err.	t
185	2147	-1.33e+04	652.019	-20.324

Note: the numbers of treated and controls refer to actual matches within radius

Bootstrapping of standard errors

```
command:      attr RE78 TREAT AGE EDUC MARR , pscore(myscore) comsup radius(.1)
statistic:    attr          = r(attr)
.....
```

```
Bootstrap statistics                                Number of obs    =      2675
                                                    Replications      =          5
```

Variable	Reps	Observed	Bias	Std. Err.	[95% Conf. Interval]		
attr	5	-13251.41	612.6666	1070.465	-16223.5	-10279.32	(N)
					-14296.53	-11596.59	(P)
					-14296.53	-11994	(BC)

```
Note:  N    = normal
       P    = percentile
       BC   = bias-corrected
```

ATT estimation with the Radius Matching method Bootstrapped standard errors

n. treat.	n. contr.	ATT	Std. Err.	t
185	2147	-1.33e+04	1070.465	-12.379

Note: the numbers of treated and controls refer to actual matches within radius

```
.
. * Kernel Matching
. attk $ylist $treatment $xlist, pscore(myscore) comsup boot reps($breps) dots
```

The program is searching for matches of each treated unit.
 This operation may take a while.

ATT estimation with the Kernel Matching method

n. treat.	n. contr.	ATT	Std. Err.	t
185	2147	-7009.315	.	.


```

184      2148    -6497.606      .      .

```

```

-----

```

Bootstrapping of standard errors

```

command:      atts RE78 TREAT AGE EDUC MARR , pscore(myscore) blockid(myblock) comsup
statistic:    atts      = r(atts)
.....

```

note: label truncated to 80 characters

```

Bootstrap statistics                                Number of obs    =      2675
                                                    Replications      =         5

```

Variable		Reps	Observed	Bias	Std. Err.	[95% Conf. Interval]	
-----+-----							
atts		5	-6497.605	235.7813	484.0005	-7841.406	-5153.805 (N)
						-7072.771	-5824.891 (P)
						-7072.771	-5988.378 (BC)

```

Note:  N   = normal
       P   = percentile
       BC  = bias-corrected

```

ATT estimation with the Stratification method
 Bootstrapped standard errors

n. treat.	n. contr.	ATT	Std. Err.	t

184	2148	-6497.605	484.000	-13.425
