

Treatment Effects Using Stata

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StataCorp LP

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México, D.F.

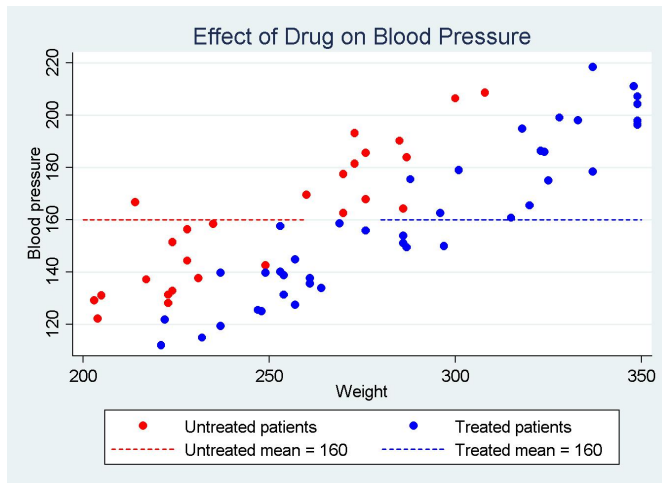
Motivation

We are interested in the outcomes of receiving a treatment in scenarios where researchers have observational data.

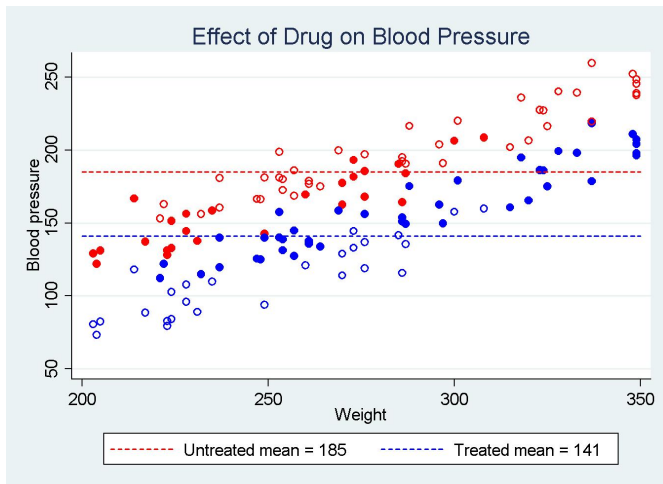
For instance:

- The impact on public education outcomes for schools that received a transfer and those that did not.
- Employment outcomes for individuals that participated in a job training program and those that did not.
- The effect on birth weight for babies of mothers that smoked relative to those of mothers that did not.

Observed Effect of Statin on Blood Pressure



Potential Outcomes of Statin on Blood Pressure



How We Approach Treatment Effects

- We cannot observe individuals in both states simultaneously
 - ▶ Design a random experiment
 - ▶ We cannot do this because of technical or ethical concerns
- We need to account for covariates that are correlated with the treatment
- We will think of the problem in terms of models that govern the treatment result and the outcome

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Notation and Definitions

- The potential outcome is denoted by the random variable y_τ with $\tau \in \{0, 1, \dots, K\}$. The potential realizations will be denoted by:
 - ▶ y_{0i} is the outcome individual i if they do not receive the treatment, where $i = 1 \dots n$
 - ▶ y_{ki} is the potential outcome for individual i if they receive different discrete levels of the treatment, where $k = 1 \dots K$
 - ▶ Usually people think about the binary case where there are only two levels y_{0i} and y_{1i}

- Potential outcome mean

$$POM = E(y_\tau)$$

- Average treatment effect

$$ATE = E(y_{ki} - y_{0i})$$

- Average treatment effect on the treated

$$ATET = E(y_{ki} - y_{0i} | \tau = k)$$

- From now on we will focus on binary treatments. All results are valid for multivariate treatments unless explicitly noted.

Assumptions

- We will be dealing with a cross-sectional random sample of n individuals

- **Overlap:**

$$0 < P(\tau_i = 1 | X_i = x) < 1$$

- **Conditional Independence:** Conditional on the covariates, X , the potential outcomes, y_0 , y_1 , and the treatment, τ , are independent

General Framework Illustrated with a Linear Example

OUTCOME MODEL:

$$y_0 = x\beta_0 + \varepsilon_0$$

$$y_1 = x\beta_1 + \varepsilon_1$$

$$y = \tau y_1 + (1 - \tau) y_0$$

TREATMENT MODEL:

$$\tau = \begin{cases} 1 & \text{if } w\gamma + \eta > 0 \\ 0 & \text{otherwise} \end{cases}$$

- w refers to the covariates that determine the treatment
- y_0 and y_1 are not observed. Only y , x , w , and τ are observed
- The random disturbances η , ε_0 , and ε_1 are independent
- The functional forms for the outcome model do not need to be linear
- All the estimators we will see arise from combinations of the outcome model and the treatment model

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$$y_1 = x\beta_1 + \varepsilon_1$$

$$y = \tau y_1 + (1 - \tau) y_0$$

TREATMENT MODEL:

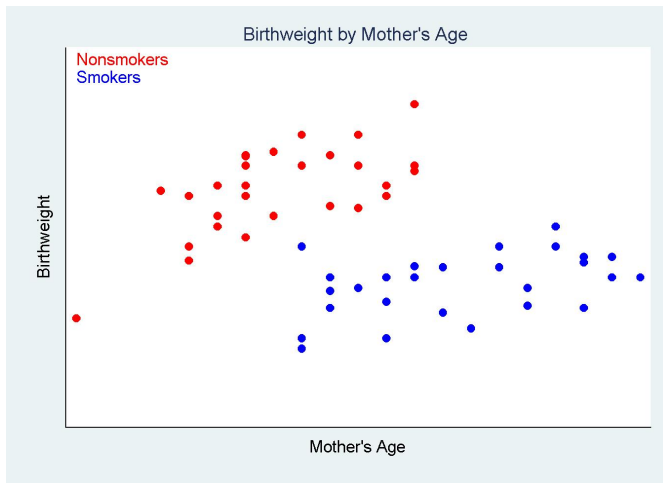
$$\tau = \begin{cases} 1 & \text{if } w\gamma + \eta > 0 \\ 0 & \text{otherwise} \end{cases}$$

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Estimators Discussed Today

- Regression Adjustment (RA)
- Inverse Probability Weighting (IPW)
- Augmented Inverse Probability Weighting (AIPW)
- Inverse Probability Weighted Regression Adjustment (IPWRA)
- Nearest Neighbor Matching
- Propensity Score Matching

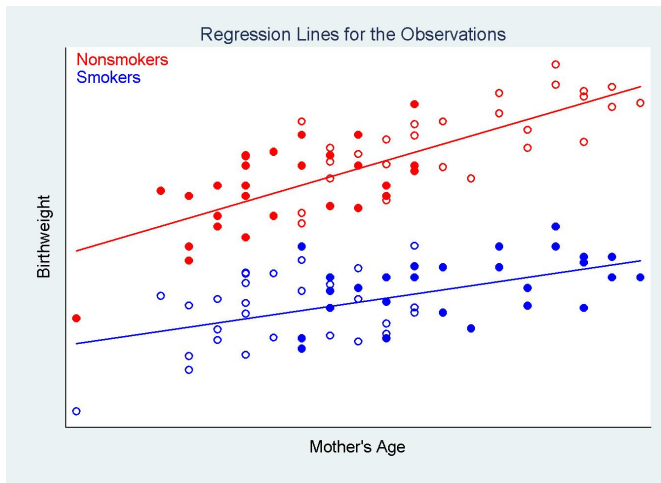
Effect of Smoking Mothers on Birthweight



Regression Adjustment (RA)

- We model the potential outcome and do not say anything about the treatment mechanism
- A conditional expectation is estimated for the treatment and control groups.
- The results from the estimations are used to compute POMs and thereafter ATEs, and ATETs.

Graphical Representation of RA Estimation



Models for the Potential Outcome

Outcome Model	$E(y x, z, \tau)$
linear	$x\beta_\tau$
logit	$\exp(x\beta_\tau) / \{1 - \exp(x\beta_\tau)\}$
probit	$\Phi(x\beta_\tau)$
poisson	$\exp(x\beta_\tau)$
hetprobit	$\Phi(x\beta_\tau/z\alpha_\tau)$

Data from Cattaneo (2010) Journal of Econometrics

```
. describe bweight lbweight mbsmoke prenatal fbaby mmarried mage fage alcohol
```

variable name	storage type	display format	value label	variable label
bweight	int	%9.0g		infant birth weight (grams)
lbweight	byte	%9.0g		1 if low birthweight baby
mbsmoke	byte	%9.0g	mbsmoke	1 if mother smoked
prenatal	byte	%9.0g		trimester of first prenatal care visit
fbaby	float	%9.0g	YesNo	1 if first baby
mmarried	byte	%10.0g	mmarried	1 if mother married
mage	byte	%9.0g		mother's age
fage	byte	%9.0g		father's age
alcohol	byte	%9.0g		1 if alcohol consumed during pregnancy

RA Linear Outcome Average Treatment Effect (ATE)

```
. teffects ra (bweight prenatal1 mmarried mage fbaby) (mbsmoke)
Iteration 0: EE criterion = 7.734e-24
Iteration 1: EE criterion = 1.196e-25
Treatment-effects estimation      Number of obs      =      4642
Estimator      : regression adjustment
Outcome model  : linear
Treatment model: none
```

	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE						
mbsmoke (smoker vs nonsmoker)	-239.6392	23.82402	-10.06	0.000	-286.3334	-192.945
POmean						
mbsmoke nonsmoker	3403.242	9.525207	357.29	0.000	3384.573	3421.911

RA Average Treatment Effect on the Treated (ATET)

```
. teffects ra (bweight prenatal1 mmarried mage fbaby) (mbSmoke), atet
Iteration 0: EE criterion = 7.629e-24
Iteration 1: EE criterion = 2.697e-26
Treatment-effects estimation      Number of obs      =      4642
Estimator      : regression adjustment
Outcome model  : linear
Treatment model: none
```

	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATET						
mbSmoke (smoker vs nonsmoker)	-223.3017	22.7422	-9.82	0.000	-267.8755	-178.7278
POmean						
mbSmoke nonsmoker	3360.961	12.75749	263.45	0.000	3335.957	3385.966

RA Probit Outcome ATE

```
. teffects ra (lbweight prenatal1 mmarried mage fbaby, probit) (mbsmoke)
Iteration 0:  EE criterion = 1.018e-18
Iteration 1:  EE criterion = 6.251e-34
Treatment-effects estimation          Number of obs      =      4642
Estimator      : regression adjustment
Outcome model  : probit
Treatment model: none
```

lbweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE						
mbsmoke (smoker vs nonsmoker)	.0500546	.0118733	4.22	0.000	.0267833	.0733259
POmean						
mbsmoke nonsmoker	.0517931	.003734	13.87	0.000	.0444745	.0591116

RA Probit ATET

```
. teffects ra (lbweight prenatal1 mmarried mage fbaby, probit) (mbsmoke), atet
Iteration 0: EE criterion = 1.018e-18
Iteration 1: EE criterion = 2.165e-34
Treatment-effects estimation      Number of obs      =      4642
Estimator      : regression adjustment
Outcome model  : probit
Treatment model: none
```

lbweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATET mbsmoke (smoker vs nonsmoker)	.0458142	.0119394	3.84	0.000	.0224134	.0692149
POmean mbsmoke nonsmoker	.0641478	.0054295	11.81	0.000	.0535063	.0747894

Inverse Probability Weighting (IPW)

- In contrast to RA estimators, IPW estimate models for the treatment
- We fit a model for the treatment and compute the probabilities of treatment
- We then compute a weighted average, using the inverse of the probability of being in each group.

Inverse Probability Weight Calculation

```
. logistic mbsmoke mmarried alcohol mage fedu  
Logistic regression
```

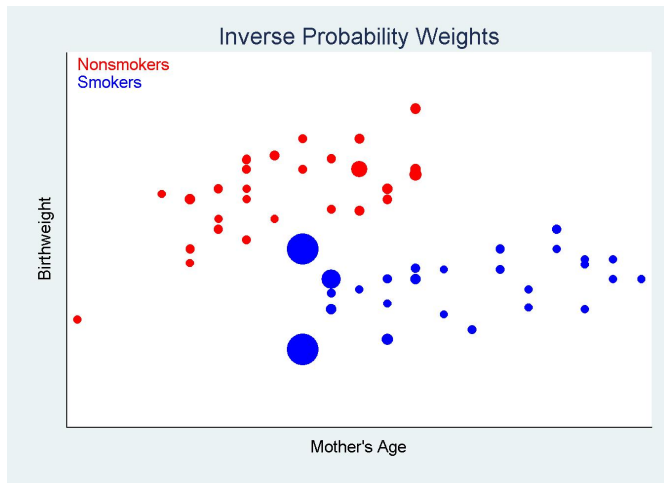
```
Number of obs   =          60  
LR chi2(4)      =         46.50  
Prob > chi2     =         0.0000  
Pseudo R2      =         0.5590
```

```
Log likelihood = -18.339432
```

	Odds Ratio	Std. Err.	z	P> z	[95% Conf. Interval]
mbsmoke					
mmarried	.0785086	.0909212	-2.20	0.028	.0081122 .7597976
alcohol	18.81727	27.98003	1.97	0.048	1.020649 346.9259
mage	2.147569	.459327	3.57	0.000	1.41218 3.265909
fedu	.8189843	.1157528	-1.41	0.158	.6208252 1.080393
_cons	4.46e-07	2.12e-06	-3.07	0.002	3.96e-11 .0050329

```
. predict ps  
(option pr assumed; Pr(mbsmoke))  
. replace ps = 1/ps if mbsmoke==1  
(30 real changes made)  
. replace ps = 1/(1-ps) if mbsmoke==0  
(30 real changes made)
```

Inverse Probability Weighting Graphically



Treatment Models

Treatment Model	$P(\tau w, z)$
logit	$\exp(w\gamma_\tau) / \{1 - \exp(w\gamma_\tau)\}$
probit	$\Phi(w\gamma_\tau)$
poisson	$\exp(w\gamma_\tau)$
hetprobit	$\Phi(w\gamma_\tau/z\theta_\tau)$

- Only the logit model is available for multivalued treatments

IPW ATE

```
. teffects ipw (bweight) (mbsmoke mmarried c.mage##c.mage fbaby medu)
Iteration 0: EE criterion = 1.713e-21
Iteration 1: EE criterion = 4.794e-27
Treatment-effects estimation      Number of obs      =      4642
Estimator      : inverse probability weighted
Outcome model  : weighted mean
Treatment model: logit
```

bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE						
mbsmoke (smoker vs nonsmoker)	-231.7203	25.17975	-9.20	0.000	-281.0717	-182.3689
POmean						
mbsmoke nonsmoker	3403.527	9.576358	355.41	0.000	3384.757	3422.296

IPW ATET

```
. teffects ipw (bweight) (mbsmoke mmarried c.mage##c.mage fbaby medu), atet
Iteration 0: EE criterion = 1.714e-21
Iteration 1: EE criterion = 3.735e-27
Treatment-effects estimation      Number of obs      =      4642
Estimator      : inverse probability weighted
Outcome model  : weighted mean
Treatment model: logit
```

	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATET						
mbsmoke (smoker vs nonsmoker)	-225.6992	23.7133	-9.52	0.000	-272.1764	-179.222
POmean						
mbsmoke nonsmoker	3363.359	14.28989	235.37	0.000	3335.351	3391.367

IPW ATE

```
. teffects ipw (bweight) (mbsmoke mmarried c.mage##c.mage fbaby medu, probit)
Iteration 0:  EE criterion = 4.622e-21
Iteration 1:  EE criterion = 8.622e-26
Treatment-effects estimation      Number of obs      =      4642
Estimator      : inverse probability weighted
Outcome model  : weighted mean
Treatment model: probit
```

bweight		Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE	mbsmoke (smoker vs nonsmoker)	-230.6886	25.81524	-8.94	0.000	-281.2856	-180.0917
POmean	mbsmoke nonsmoker	3403.463	9.571369	355.59	0.000	3384.703	3422.222

IPW ATET

```
. teffects ipw (bweight) (mbsmoke mmarried c.mage##c.mage fbaby medu, probit),  
> ///  
> atet  
Iteration 0: EE criterion = 4.621e-21  
Iteration 1: EE criterion = 7.103e-27  
Treatment-effects estimation      Number of obs      =      4642  
Estimator      : inverse probability weighted  
Outcome model  : weighted mean  
Treatment model: probit
```

	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATET						
mbsmoke (smoker vs nonsmoker)	-225.1773	23.66458	-9.52	0.000	-271.559	-178.7955
POMean						
mbsmoke nonsmoker	3362.837	14.20149	236.79	0.000	3335.003	3390.671

Doubly Robust Estimators

- Doubly robust estimators model both the treatment and the outcome model
- These models are interesting because they are consistent even if one of the models is misspecified
- Augmented Inverse Probability Weighting (AIPW) and Inverse Probability Weighted Regression Adjustment (IPWRA) have this property

Double Robust Estimators AIPW

- Estimate a treatment model and compute inverse-probability weights
- Estimate separate regression model of the outcome for each treatment level
 - ▶ We allow the outcome model to be estimated by nonlinear least squares or weighted nonlinear least squares
- Compute the weighted means of the treatment-specific predicted outcomes, where the weights are the inverse-probability weights computed in step.

ATE for AIPW

```
. teffects aipw (bweight prenatal1 mmarried mage fbaby) ///  
> (mbsmoke mmarried c.mage##c.mage fbaby medu)  
Iteration 0: EE criterion = 1.721e-21  
Iteration 1: EE criterion = 2.247e-26  
Treatment-effects estimation          Number of obs      =      4642  
Estimator      : augmented IPW  
Outcome model  : linear by ML  
Treatment model: logit
```

	bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE	mbsmoke (smoker vs nonsmoker)	-232.0409	25.66973	-9.04	0.000	-282.3527	-181.7292
POmean	mbsmoke nonsmoker	3403.457	9.570043	355.64	0.000	3384.7	3422.214

ATE for AIPW with Nonlinear Least Squares

```
. teffects aipw (bweight prenatal1 mmarried mage fbaby, poisson) ///  
> (mbsmoke mmarried c.mage##c.mage fbaby medu), nls  
Iteration 0: EE criterion = .00018418  
Iteration 1: EE criterion = 1.991e-17  
Treatment-effects estimation      Number of obs      =      4642  
Estimator      : augmented IPW  
Outcome model  : Poisson by NLS  
Treatment model: logit
```

	bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE	mbsmoke (smoker vs nonsmoker)	-232.1593	25.69692	-9.03	0.000	-282.5244	-181.7943
POmean	mbsmoke nonsmoker	3403.444	9.57036	355.62	0.000	3384.687	3422.202

Displaying Treatment and Outcome Equations

```
. teffects aipw (bweight prenatal1 mmarried mage fbaby, poisson) ///
> (mbsmoke mmarried c.mage##c.mage fbaby medu), aequations nolog
Treatment-effects estimation      Number of obs   =      4642
Estimator      : augmented IPW
Outcome model  : Poisson by ML
Treatment model: logit
```

bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE						
mbsmoke (smoker vs nonsmoker)	-232.1369	25.68896	-9.04	0.000	-282.4864	-181.7875
POmean						
mbsmoke nonsmoker	3403.444	9.570363	355.62	0.000	3384.686	3422.202
OME0						
prenatal1	.0191803	.0082502	2.32	0.020	.0030102	.0353503
mmarried	.0480049	.0080048	6.00	0.000	.0323158	.0636939
mage	.0007522	.0006106	1.23	0.218	-.0004447	.001949
fbaby	-.0209166	.0057619	-3.63	0.000	-.0322097	-.0096235
_cons	8.072261	.0159896	504.84	0.000	8.040922	8.1036
OME1						
prenatal1	.0080848	.012943	0.62	0.532	-.0172831	.0334526
mmarried	.0426096	.0130351	3.27	0.001	.0170612	.0681579
mage	-.0023601	.0013552	-1.74	0.082	-.0050163	.0002961
fbaby	.0131662	.0126163	1.04	0.297	-.0115613	.0378937
_cons	8.07972	.0334184	241.77	0.000	8.014221	8.145219
TME1						
mmarried	-1.145706	.0975846	-11.74	0.000	-1.336969	-.9544439
mage	.321518	.0657363	4.89	0.000	.1926773	.4503588
c.mage#c.mage						
	-.0060368	.0012234	-4.93	0.000	-.0084346	-.0036389
fbaby	-.3864258	.0894428	-4.32	0.000	-.5617305	-.2111211
medu	-.1420833	.0179132	-7.93	0.000	-.1771926	-.106974
_cons	-2.950915	.8302955	-3.55	0.000	-4.578264	-1.323565

Double Robust Estimators Inverse Probability Weighted Regression Adjustment (IPWRA)

- Estimate a treatment model and compute inverse-probability weights
- Use the estimated inverse-probability weights and fit weighted regression models of the outcome for each treatment level
- Compute the means of the treatment-specific predicted outcomes

ATET for Inverse Probability Weighted Regression Adjustment

```
. teffects ipwra (bweight prenatal1 mmarried mage fbaby) ///  
> (mbsmoke mmarried c.mage##c.mage fbaby medu), atet  
Iteration 0: EE criterion = 4.620e-21  
Iteration 1: EE criterion = 1.345e-26  
Treatment-effects estimation      Number of obs      =      4642  
Estimator      : IPW regression adjustment  
Outcome model  : linear  
Treatment model: logit
```

	bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATET	mbsmoke (smoker vs nonsmoker)	-224.0108	23.846	-9.39	0.000	-270.7481	-177.2735
POmean	mbsmoke nonsmoker	3361.671	14.54939	231.05	0.000	3333.154	3390.187

Displaying Treatment and Outcome Equations

```
. teffects ipwra (bweight prenatal1 mmarried mage fbaby) ///
> (mbsmoke mmarried c.mage#c.mage fbaby medu), ateq equations
Iteration 0: EE criterion = 4.620e-21
Iteration 1: EE criterion = 1.345e-26
Treatment-effects estimation      Number of obs      =      4642
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Outcome model  : linear
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```

	bweight	Coef.	Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATET	mbsmoke (smoker vs nonsmoker)	-224.0108	23.846	-9.39	0.000	-270.7481	-177.2735
POmean	mbsmoke nonsmoker	3361.671	14.54939	231.05	0.000	3333.154	3390.187
OME0	prenatal1	77.07926	40.4633	1.90	0.057	-2.227341	156.3859
	mmarried	138.9961	29.48776	4.71	0.000	81.20114	196.791
	mage	4.482273	3.033008	1.48	0.139	-1.462313	10.42686
	fbaby	-73.85266	32.55461	-2.27	0.023	-137.6585	-10.0468
	_cons	3157.337	72.75786	43.40	0.000	3014.734	3299.939
OME1	prenatal1	25.11133	40.37541	0.62	0.534	-54.02302	104.2457
	mmarried	133.6617	40.86443	3.27	0.001	53.5689	213.7545
	mage	-7.370881	4.21817	-1.75	0.081	-15.63834	.8965804
	fbaby	41.43991	39.70712	1.04	0.297	-36.38461	119.2644
	_cons	3227.169	104.4059	30.91	0.000	3022.537	3431.801
TME1	mmarried	-1.145706	.0975846	-11.74	0.000	-1.336969	-.9544439
	mage	.321518	.0657363	4.89	0.000	.1926773	.4503588
c.mage#c.mage		-.0060368	.0012234	-4.93	0.000	-.0084346	-.0036389
	fbaby	-.3864258	.0894428	-4.32	0.000	-.5617305	-.2111211
	medu	-.1420833	.0179132	-7.93	0.000	-.1771926	-.106974
	_cons	-2.950915	.8302955	-3.55	0.000	-4.578264	-1.323565

Nearest Neighbor Matching

- Can be understood as an outcome model within our framework
- Matches the closest individuals in terms of covariates
- Is a nonparametric estimate with an asymptotic bias. We provide a bias correction option.
- These estimators are nondifferentiable therefore the bootstrap is not allowed
- These estimators do not allow for multivalued treatments

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ATE with nnmatch

```
. teffects nnmatch (bweight mage prenatal1 mmarried fbaby) (mbSmoke)
Treatment-effects estimation      Number of obs   =      4642
Estimator      : nearest-neighbor matching      Matches: requested =      1
Outcome model  : matching                      min =      1
Distance metric: Mahalanobis                  max =     139
```

	bweight	Coef.	AI Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE							
mbSmoke (smoker vs nonsmoker)		-240.3306	28.43006	-8.45	0.000	-296.0525	-184.6087

Exact Matching and Different Distance

```
. teffects nnmatch (bweight mage) (mb smoke), ///  
> ematch(prenatall mmarried fbaby) metric(euclidean)  
Treatment-effects estimation      Number of obs      =      4642  
Estimator      : nearest-neighbor matching      Matches: requested =      1  
Outcome model  : matching                      min =      1  
Distance metric: Euclidean                      max =     139
```

	bweight	Coef.	AI Robust Std. Err.	z	P> z	[95% Conf. Interval]
ATE	mb smoke (smoker vs nonsmoker)	-240.3306	28.43006	-8.45	0.000	-296.0525 -184.6087

Bias Adjustment

```
. teffects nnmatch (bweight mage fage) (mbsmoke), ///  
> ematch(prenatall mmarried fbaby) biasadj(mage fage)  
Treatment-effects estimation      Number of obs      =      4642  
Estimator      : nearest-neighbor matching      Matches: requested =      1  
Outcome model  : matching                      min =      1  
Distance metric: Mahalanobis                  max =      25
```

	bweight	Coef.	AI Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE	mbsmoke (smoker vs nonsmoker)	-223.8389	26.19973	-8.54	0.000	-275.1894	-172.4883

Propensity Score Matching

- Can be classified within the class of treatment models
- Estimate the treatment probabilities (propensity scores)
- Assign values to unobserved outcomes based on observed ones with similar propensity scores
- Estimate ATE
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Propensity Score Matching controlling matches

```
. teffects psmatch (bweight) (mbsmoke mmarried c.mage##c.mage fbaby medu), ///  
> nneighbor(2)  
Treatment-effects estimation      Number of obs      =      4642  
Estimator      : propensity-score matching      Matches: requested =      2  
Outcome model  : matching                      min =      2  
Treatment model: logit                        max =      74
```

	bweight	Coef.	AI Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE	mbsmoke (smoker vs nonsmoker)	-214.2469	27.47783	-7.80	0.000	-268.1025	-160.3914

Conclusion

- We have presented a host of treatment effects estimators within a unified framework
- The estimators are parametric and nonparametric and in the parametric cases can be consistent under misspecification of the potential outcome or treatment models
- The estimators provide estimates and inference for quantities of interest for researchers, POM, ATE, ATET.

Double Robustness I

- Let $P(\tau|x, z, \hat{\gamma}) =: M_P(\hat{\gamma})$ be our estimated conditional treatment probabilities
- Let $E(y|x, z, \tau, \hat{\beta}) =: M_E(\hat{\beta}_\tau)$ define our estimated conditional means
- We define the following estimators for the POMs

$$\hat{E}(y_1) = \frac{1}{n} \sum_{i=1}^n \left[\frac{\tau_i y_i}{M_P(\hat{\gamma})} - \frac{\{\tau_i - M_P(\hat{\gamma})\}}{M_P(\hat{\gamma})} M_E(\hat{\beta}_1) \right]$$
$$\hat{E}(y_0) = \frac{1}{n} \sum_{i=1}^n \left[\frac{(1 - \tau_i) y_i}{1 - M_P(\hat{\gamma})} - \frac{\{\tau_i - M_P(\hat{\gamma})\}}{1 - M_P(\hat{\gamma})} M_E(\hat{\beta}_0) \right]$$

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Intuition Behind Double Robustness II

- We will focus on $\hat{E}(y_1)$ (a similar argument follows for $\hat{E}(y_0)$)
- By the law of large numbers it follows that $\hat{E}(y_1)$ has the following probability limit:

$$\begin{aligned}\hat{E}(y_1) &\xrightarrow{P} E \left[\frac{\tau y}{M_P(\gamma)} - \frac{\{\tau - M_P(\gamma)\}}{M_P(\gamma)} M_E(\beta_1) \right] \\ &= E \left[\frac{\tau y_1}{M_P(\gamma)} - \frac{\{\tau - M_P(\gamma)\}}{M_P(\gamma)} M_E(\beta_1) + y_1 - y_1 \right] \\ &= E \left[\frac{\tau y_1}{M_P(\gamma)} - \frac{\{\tau - M_P(\gamma)\}}{M_P(\gamma)} M_E(\beta_1) + y_1 - y_1 \frac{M_P(\gamma)}{M_P(\gamma)} \right] \\ &= E \left[\frac{y_1 (\tau - M_P(\gamma))}{M_P(\gamma)} - \frac{\{\tau - M_P(\gamma)\}}{M_P(\gamma)} M_E(\beta_1) + y_1 \right] \\ &= E \left[\frac{\{\tau - M_P(\gamma)\}}{M_P(\gamma)} (y_1 - M_E(\beta_1)) + y_1 \right] \\ &= E(y_1) + E \left[\frac{\{\tau - M_P(\gamma)\}}{M_P(\gamma)} (y_1 - M_E(\beta_1)) \right]\end{aligned}$$

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Intuition Behind Double Robustness III

$$\hat{E}(y_1) \xrightarrow{P} E(y_1) + E \left[\frac{\{\tau - M_P(\gamma)\}}{M_P(\gamma)} (y_1 - M_E(\beta_1)) \right]$$

- Given conditional independence of treatment and outcome conditional on the regressors by the law of iterated expectations:
 - ▶ If the outcome model is correctly specified $E[y_1 - M_E(\beta_1)] = 0$. This implies that even if the treatment model is incorrectly specified, $\hat{E}(y_1) \xrightarrow{P} E(y_1)$
 - ▶ Similarly if the treatment model is correctly specified $E[\tau - M_P(\gamma)] = 0$. Thus, even if $E[y_1 - M_E(\beta_1)] \neq 0$ we have that $\hat{E}(y_1) \xrightarrow{P} E(y_1)$

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Estimation I: An Example of Moment Based Estimation

We define the projection model by:

$$\begin{aligned}y &= X\beta + \varepsilon \\E(X'\varepsilon) &= 0\end{aligned}$$

β is then given by:

$$\begin{aligned}0 &= E(X'\varepsilon) \\0 &= E(X'\{y - X\beta\}) \\ \beta &= E(X'X)^{-1} E(X'y)\end{aligned}$$

A consistent estimator of β is:

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Estimation II: Methodology We Employ

- The different specifications for the outcome generate moment conditions
- We can then use GMM to estimate the parameters of interest
- For the linear model:

$$0 = E [\tau(y - x\beta_1)'x + (1 - \tau)(y - x\beta_0)'x]$$

- For the probit and logit models

$$0 = E \left(\tau \left[\frac{g(x\beta_1) \{y - G(x\beta_1)\}}{G(x\beta_1) \{1 - G(x\beta_1)\}} \right] + (1 - \tau) \left[\frac{g(x\beta_1) \{y - G(x\beta_0)\}}{G(x\beta_0) \{1 - G(x\beta_0)\}} \right] \right)$$

- ▶ $G(\cdot)$ is either the standard normal CDF or the logistic function
- ▶ $g(\cdot)$ is the derivative of $G(\cdot)$

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Distance

The distance function is given by:

$$\|x_i - x_j\|_S = \left\{ (x_i - x_j)' S^{-1} (x_i - x_j) \right\}^{1/2}$$

where S can be:

$$S = \begin{cases} \frac{(X - x1_n)' W (X - x1_n)}{\sum_{i=1}^n w_i - 1} & \text{if metric is mahalanobis} \\ \text{diagonal} \left\{ \frac{(X - x1_n)' W (X - x1_n)}{\sum_{i=1}^n w_i - 1} \right\} & \text{if metric is ivariance} \\ I_k & \text{if metric is euclidean} \end{cases}$$

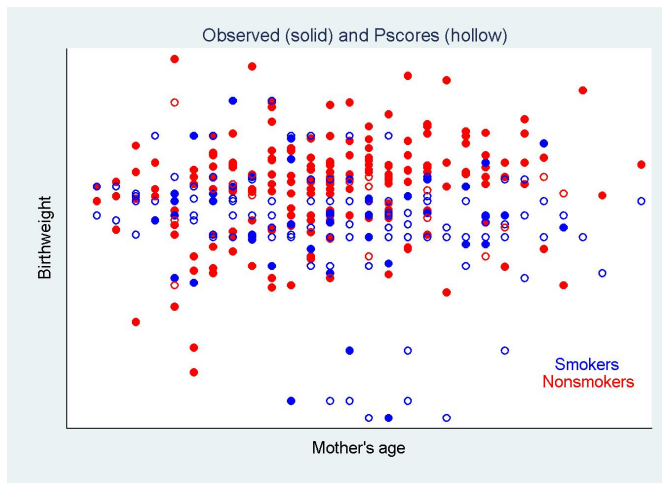
Above 1_n is an n vector of ones, W is a matrix of frequency weights

ATE for Propensity Score Matching

```
. teffects psmatch (bweight) (mbsmoke mmarried c.mage##c.mage fbaby medu), ///  
> generate(ps)  
Treatment-effects estimation      Number of obs      =      4642  
Estimator      : propensity-score matching      Matches: requested =      1  
Outcome model  : matching                      min =      1  
Treatment model: logit                        max =      74
```

	bweight	Coef.	AI Robust Std. Err.	z	P> z	[95% Conf. Interval]	
ATE	mbsmoke (smoker vs nonsmoker)	-210.9683	32.021	-6.59	0.000	-273.7284	-148.2083

Matches Generated by the Estimator



A Nonsequitur

- Imbens and Wooldridge (2009) JEL for a recent survey
- Regression Discontinuity. Lee and Lemieux (2010) JEL
- Nonparametric Multivariate Treatment Effects. See Cattaneo 2010 in the New Palgrave Dictionary and Cattaneo 2010 JOE.
- Stata also offers estimation in the presence of endogeneity