

# Causal Inference Methods and Case Studies

STAT24630

Jingshu Wang

# Lecture 14

Topic: Inverse probability weighting

- IPW
  - Using normalized weights
- Doubly robust estimator
- Bootstrap
- Textbook Chapters 17.8, 19.10.2

# Motivation

- Matching methods can improve covariate balance
- Potential limitations of matching methods:
  - Inefficient: it may throw away many control units
  - Ineffective: it may not be able to balance covariates
  - Biased: not estimating the ATT if a lot of treated units are not matched
- Matching is a special case of weighting

$$\begin{aligned}\hat{\tau}_{\text{match}} &= \frac{1}{N_t} \sum_{i=1}^N W_i \left( Y_i^{\text{obs}} - \frac{1}{|\mathcal{M}_i^c|} \sum_{i' \in \mathcal{M}_i^c} Y_{i'}^{\text{obs}} \right) \\ &= \frac{1}{N_t} \sum_{i:W_i=1} Y_i^{\text{obs}} - \frac{1}{N_c} \sum_{i:W_i=1} \left( \frac{N_c}{N_t} \sum_{i':W_{i'}=1} \frac{1_{i \in \mathcal{M}_{i'}^c}}{|\mathcal{M}_{i'}^c|} \right) Y_i^{\text{obs}}\end{aligned}$$

- Idea: weight each observation in the control group such that it looks like the treatment group

# Inverse probability weighting (IPW)

- Weighting makes use of the following properties to estimate  $\mathbb{E}(Y_i(1))$  and  $\mathbb{E}(Y_i(0))$

$$\mathbb{E} \left[ \frac{Y_i^{\text{obs}} \cdot W_i}{e(X_i)} \right] = \mathbb{E}_{\text{sp}} [Y_i(1)], \quad \text{and} \quad \mathbb{E} \left[ \frac{Y_i^{\text{obs}} \cdot (1 - W_i)}{1 - e(X_i)} \right] = \mathbb{E}_{\text{sp}} [Y_i(0)].$$

- Intuitively, unit that has a smaller  $e(X_i)$  has less chance to appear in the treatment group, so we should give it a higher weight (the less likely a subject is sampled, then the larger population it should represent)

$$\hat{\tau}_{\text{IPW}} = \frac{1}{N} \sum_{i=1}^N \frac{W_i \cdot Y_i^{\text{obs}}}{e(X_i)} - \frac{1}{N} \sum_{i=1}^N \frac{(1 - W_i) \cdot Y_i^{\text{obs}}}{1 - e(X_i)}$$

$$= \frac{1}{N} \sum_{i:W_i=1} \lambda_i \cdot Y_i^{\text{obs}} - \frac{1}{N} \sum_{i:W_i=0} \lambda_i \cdot Y_i^{\text{obs}},$$

where

$$\lambda_i = \frac{1}{e(X_i)^{W_i} \cdot (1 - e(X_i))^{1-W_i}} = \begin{cases} 1/(1 - e(X_i)) & \text{if } W_i = 0, \\ 1/e(X_i) & \text{if } W_i = 1. \end{cases}$$

# IPW for observational studies

- The propensity scores are estimated
- Estimate ATE and ATT
  - ATE

$$\widehat{\text{ATE}} = \frac{1}{N} \sum_{i=1}^N \left\{ \frac{W_i Y_i^{\text{obs}}}{\hat{e}(\mathbf{X}_i)} - \frac{(1 - W_i) Y_i^{\text{obs}}}{1 - \hat{e}(\mathbf{X}_i)} \right\}$$

- ATT

$$\widehat{\text{ATT}} = \frac{1}{N_t} \sum_{i=1}^N \left\{ W_i Y_i^{\text{obs}} - \frac{\hat{e}(\mathbf{X}_i)(1 - W_i) Y_i^{\text{obs}}}{1 - \hat{e}(\mathbf{X}_i)} \right\}$$

- For units that have identical propensity scores  $\rightarrow$  difference-in-means estimator

# Normalizing the weights

- When use any weighting method (e.g. IPW), good practice is to normalize weights – sum of the total of weights within one group should be 1
- Divide each unit's weight ( $\omega_i$ ) by the sum of all weights in that group  $\omega_i / \sum_{i': W_{i'}=w} \omega_{i'}$ , for  $w = 0, 1$ , i.e. the Hajek estimator:
- The new ATE estimator:

$$\widehat{\text{ATE}} = \frac{\sum_{i=1}^N W_i Y_i^{\text{obs}} / \hat{e}(\mathbf{X}_i)}{\sum_{i=1}^N W_i / \hat{e}(\mathbf{X}_i)} - \frac{\sum_{i=1}^N (1 - W_i) Y_i^{\text{obs}} / (1 - \hat{e}(\mathbf{X}_i))}{\sum_{i=1}^N (1 - W_i) / (1 - \hat{e}(\mathbf{X}_i))}$$

- The new ATT estimator:

$$\widehat{\text{ATT}} = \frac{1}{N_t} \sum_{i=1}^N W_i Y_i^{\text{obs}} - \frac{\sum_{i=1}^N (1 - W_i) Y_i^{\text{obs}} \hat{e}(\mathbf{X}_i) / (1 - \hat{e}(\mathbf{X}_i))}{\sum_{i=1}^N (1 - W_i) \hat{e}(\mathbf{X}_i) / (1 - \hat{e}(\mathbf{X}_i))}$$

- Using normalized weights, we can reduce variance and lead to more stable estimate (Hirano, Imbens, Ridder, 2003)

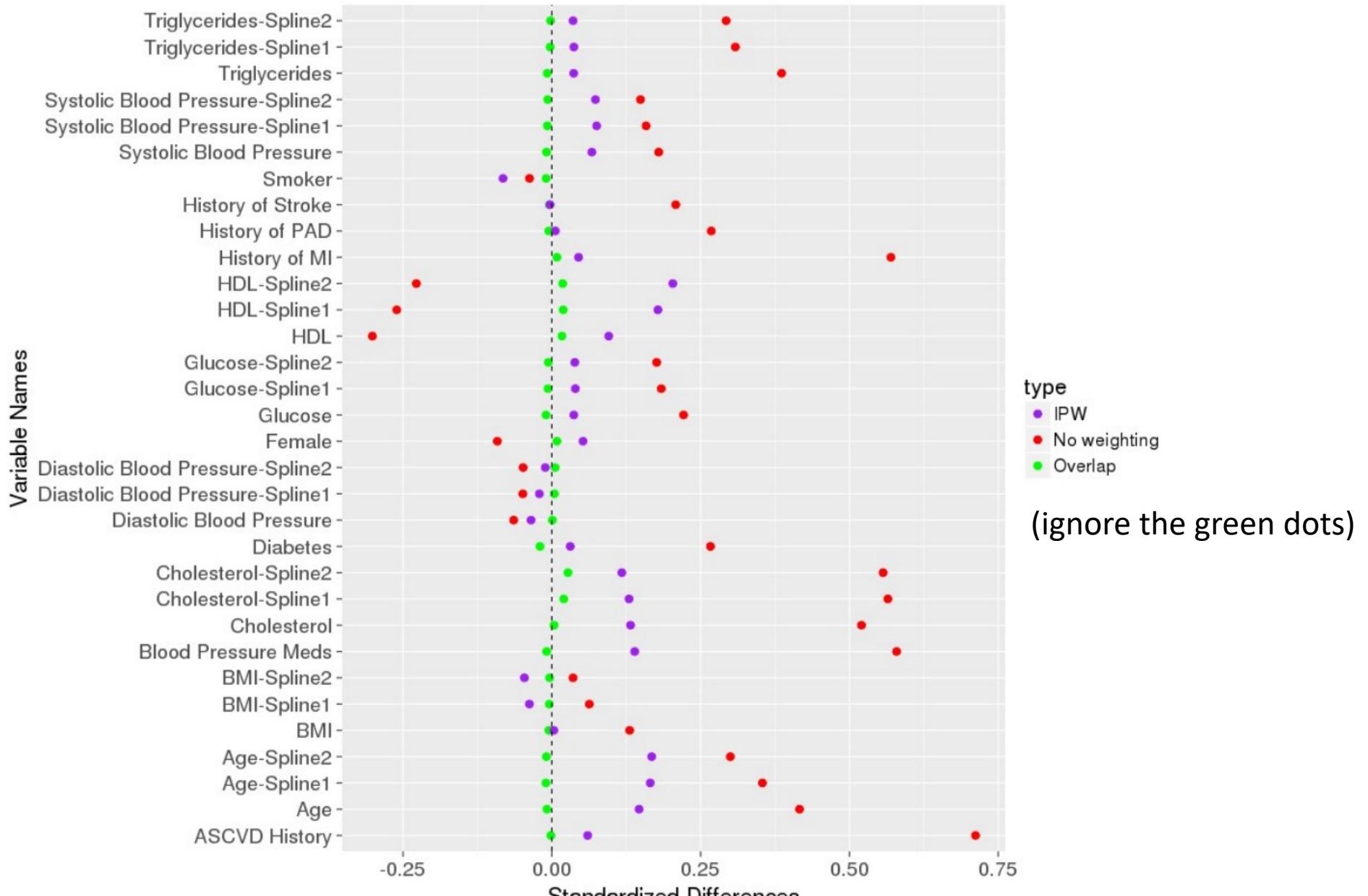
# IPW advantages v.s. disadvantages

- Advantages
  - Simple, with theoretical foundation
  - Global balance
  - Use all data
  - Can be extended to more complex settings
- Disadvantages
  - More sensitive to misspecification of propensity scores than matching
  - Estimated propensity scores near 0 or 1 can yield extreme weights

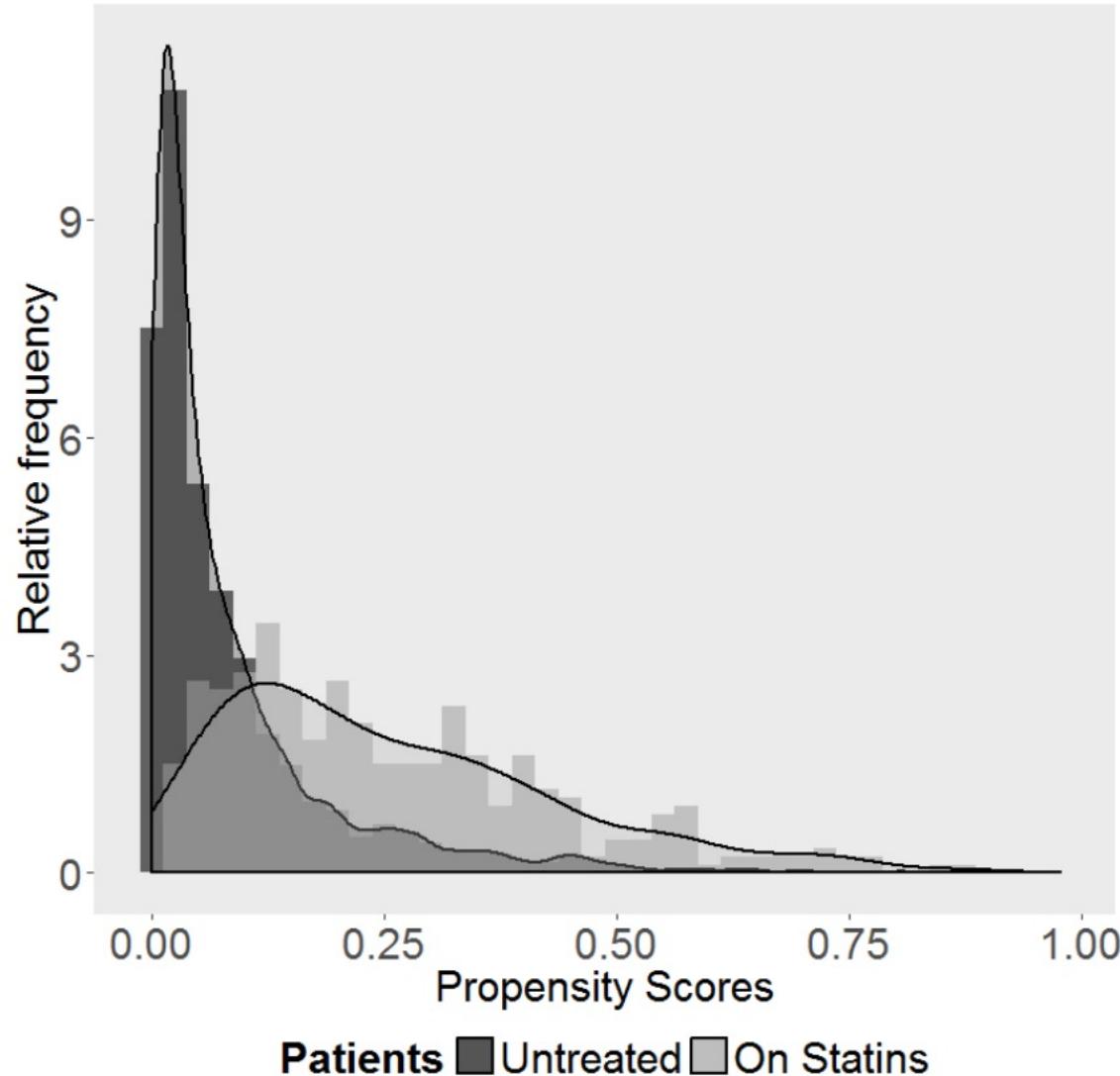
# Example: Framingham Heart Study

- Goal: evaluate the effect of statins on health outcomes
- Patients: cross-sectional population from the offspring cohort with a visit 6 (1995-1998)
- Treatment: statin use at visit 6 vs. no statin use
- Outcomes: CV(cardiovascular) death, myocardial infarction (MI), stroke
- Confounders: sex, age, body mass index, diabetes, history of MI, history of PAD, history of stroke...
- Significant imbalance between treatment and control groups in covariates motivates IPW (or some form of propensity score adjustment)

# Love plot for covariate balancing



# Distribution of estimated propensity scores



- For treated units with  $\hat{e}(\mathbf{X}_i)$  close to 0, then can greatly influence the IPW estimator value
- Will discuss trimming in later lectures

# Doubly robust estimator

- Outcome regression relies on a correctly specified model for the (potential) outcomes depending on  $\mathbf{X}_i$
- IPW / Matching relies on a correctly specified model for the propensity score
- Doubly robust estimator: provide a good estimate of the propensity score when either the outcome or the propensity score model is correct
- Define

$$f(1, \mathbf{X}_i, Y_i^{\text{obs}}) = \frac{Y_i^{\text{obs}} 1_{W_i=1}}{\tilde{e}(\mathbf{X}_i)} - \frac{1_{W_i=1} - \tilde{e}(\mathbf{X}_i)}{\tilde{e}(\mathbf{X}_i)} \tilde{\mu}_1(\mathbf{X}_i)$$

$$f(0, \mathbf{X}_i, Y_i^{\text{obs}}) = \frac{Y_i^{\text{obs}} 1_{W_i=0}}{1 - \tilde{e}(\mathbf{X}_i)} - \frac{1_{W_i=0} - (1 - \tilde{e}(\mathbf{X}_i))}{1 - \tilde{e}(\mathbf{X}_i)} \tilde{\mu}_0(\mathbf{X}_i)$$

- If we correctly specify the **propensity score model**, then  $\tilde{e}(\mathbf{X}_i) = e(\mathbf{X}_i)$
- If we correctly specify the **outcome model**, then  $\tilde{\mu}_w(\mathbf{X}_i) = \mu_w(\mathbf{X}_i)$

# Doubly robust estimator

$$f(1, \mathbf{X}_i, Y_i^{\text{obs}}) = \frac{Y_i^{\text{obs}} 1_{W_i=1}}{\tilde{e}(\mathbf{X}_i)} - \frac{1_{W_i=1} - \tilde{e}(\mathbf{X}_i)}{\tilde{e}(\mathbf{X}_i)} \tilde{\mu}_1(\mathbf{X}_i)$$

$$f(0, \mathbf{X}_i, Y_i^{\text{obs}}) = \frac{Y_i^{\text{obs}} 1_{W_i=0}}{1 - \tilde{e}(\mathbf{X}_i)} - \frac{1_{W_i=0} - (1 - \tilde{e}(\mathbf{X}_i))}{1 - \tilde{e}(\mathbf{X}_i)} \tilde{\mu}_0(\mathbf{X}_i)$$

- $\tilde{e}(\mathbf{X}_i), \tilde{\mu}_w(\mathbf{X}_i)$ : our working models (model under our model assumption)
- $e(\mathbf{X}_i), \mu_w(\mathbf{X}_i)$ : true model that we don't know
- Double robust property

$$\mathbb{E} [f(1, \mathbf{X}_i, Y_i^{\text{obs}}) \mid \mathbf{X}_i] = \frac{(\mu_1(\mathbf{X}_i) - \tilde{\mu}_1(\mathbf{X}_i)) (e(\mathbf{X}_i) - \tilde{e}(\mathbf{X}_i))}{\tilde{e}(\mathbf{X}_i)} + \mu_1(\mathbf{X}_i)$$

$$\mathbb{E} [f(0, \mathbf{X}_i, Y_i^{\text{obs}}) \mid \mathbf{X}_i] = \frac{(\mu_0(\mathbf{X}_i) - \tilde{\mu}_0(\mathbf{X}_i)) (\tilde{e}(\mathbf{X}_i) - e(\mathbf{X}_i))}{1 - \tilde{e}(\mathbf{X}_i)} + \mu_0(\mathbf{X}_i)$$

- If either the outcome or propensity score model is correct, we have

$$\mathbb{E} \left( f(w, \mathbf{X}_i, Y_i^{\text{obs}}) \right) = \mathbb{E}(Y_i(w) \mid \mathbf{X}_i)$$

# Doubly robust estimator

$$f(1, \mathbf{X}_i, Y_i^{\text{obs}}) = \frac{Y_i^{\text{obs}} 1_{W_i=1}}{\tilde{e}(\mathbf{X}_i)} - \frac{1_{W_i=1} - \tilde{e}(\mathbf{X}_i)}{\tilde{e}(\mathbf{X}_i)} \tilde{\mu}_1(\mathbf{X}_i)$$

IPW estimate of  $\mathbb{E}(Y_i(1) | \mathbf{X}_i)$

Adjust for bias if the propensity score model is incorrect  
(if PS model is correct, then this part has expectation 0)

An equivalent expression:

$$f(1, \mathbf{X}_i, Y_i^{\text{obs}}) = \tilde{\mu}_1(\mathbf{X}_i) + \frac{1_{W_i=1}}{\tilde{e}(\mathbf{X}_i)} (Y_i(1) - \tilde{\mu}_1(\mathbf{X}_i))$$

Outcome regression estimate of  $\mathbb{E}(Y_i(1) | \mathbf{X}_i)$

Adjust for bias if the outcome regression model is incorrect  
(if PS model is correct, then this part has expectation 0)

The DR estimator:

$$\hat{\tau} = \frac{1}{N} \sum_i \left[ \frac{Y_i^{\text{obs}} 1_{W_i=1}}{\hat{e}(\mathbf{X}_i)} - \frac{1_{W_i=1} - \hat{e}(\mathbf{X}_i)}{\hat{e}(\mathbf{X}_i)} \hat{\mu}_1(\mathbf{X}_i) \right] - \frac{1}{N} \sum_i \left[ \frac{Y_i^{\text{obs}} 1_{W_i=0}}{1 - \hat{e}(\mathbf{X}_i)} - \frac{1_{W_i=0} - (1 - \hat{e}(\mathbf{X}_i))}{1 - \hat{e}(\mathbf{X}_i)} \hat{\mu}_0(\mathbf{X}_i) \right]$$

# A simulation study (Kang and Schafer. 2007. Statistical Science)

- The deteriorating performance of propensity score weighting methods when the model is misspecified
- Setup:
  - 4 covariates  $X_i^*$ : all are i.i.d. standard normal
  - Outcome model: linear model
  - Propensity score model: logistic model with linear predictors
  - Misspecification induced by measurement error:
    - $X_{i1} = \exp(X_{i1}^*/2)$
    - $X_{i2} = X_{i2}^*/(1 + \exp(X_{1i}^*) + 10)$
    - $X_{i3} = (X_{i1}^* X_{i3}^*/25 + 0.6)^3$
    - $X_{i4} = (X_{i1}^* + X_{i4}^* + 20)^2$
- Weighting estimators to be evaluated:
  - HT: IPW in the original form
  - IPW: IPW with normalized weights
  - Weighted least squares regression with covariates
  - Doubly-robust estimator

# Results: if the propensity score model is correct

Sample size	Estimator	Bias		RMSE	
		logit	True	logit	True
<b>(1) Both models correct</b>					
$n = 200$	HT	0.33	1.19	12.61	23.93
	IPW	-0.13	-0.13	3.98	5.03
	WLS	-0.04	-0.04	2.58	2.58
	DR	-0.04	-0.04	2.58	2.58
$n = 1000$	HT	0.01	-0.18	4.92	10.47
	IPW	0.01	-0.05	1.75	2.22
	WLS	0.01	0.01	1.14	1.14
	DR	0.01	0.01	1.14	1.14
<b>(2) Propensity score model correct</b>					
$n = 200$	HT	-0.05	-0.14	14.39	24.28
	IPW	-0.13	-0.18	4.08	4.97
	WLS	0.04	0.04	2.51	2.51
	DR	0.04	0.04	2.51	2.51
$n = 1000$	HT	-0.02	0.29	4.85	10.62
	IPW	0.02	-0.03	1.75	2.27
	WLS	0.04	0.04	1.14	1.14
	DR	0.04	0.04	1.14	1.14

- Use the true propensity score is worse than using the estimated propensity score when the propensity score model is correct
- Normalizing weights can help a lot in reducing the variance

# Results: if the propensity score model is incorrect

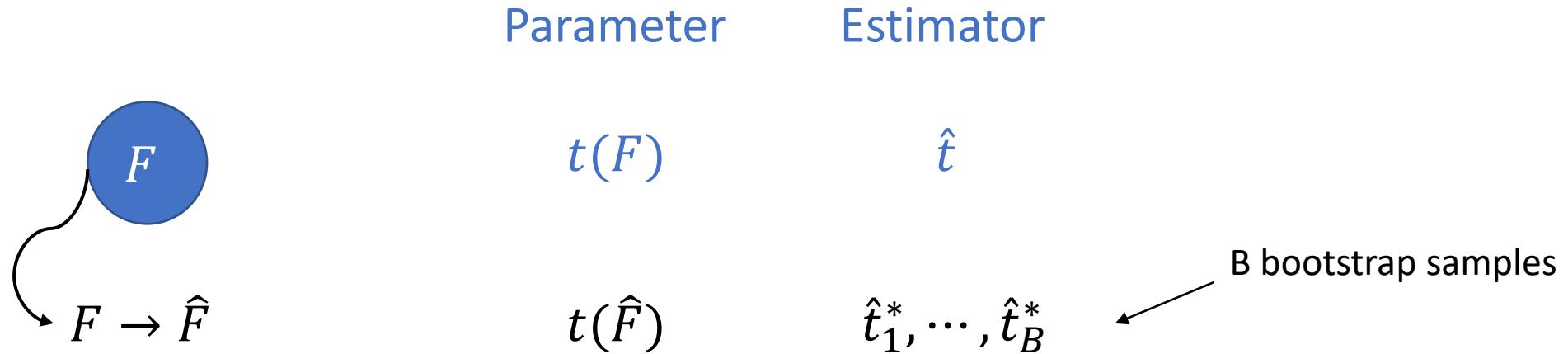
Sample size	Estimator	Bias		RMSE	
		logit	True	logit	True
<b>(3) Outcome model correct</b>					
$n = 200$	HT	24.25	-0.18	194.58	23.24
	IPW	1.70	-0.26	9.75	4.93
	WLS	-2.29	0.41	4.03	3.31
	DR	-0.08	-0.10	2.67	2.58
$n = 1000$	HT	41.14	-0.23	238.14	10.42
	IPW	4.93	-0.02	11.44	2.21
	WLS	-2.94	0.20	3.29	1.47
	DR	0.02	0.01	1.89	1.13
<b>(4) Both models incorrect</b>					
$n = 200$	HT	30.32	-0.38	266.30	23.86
	IPW	1.93	-0.09	10.50	5.08
	WLS	-2.13	0.55	3.87	3.29
	DR	-7.46	0.37	50.30	3.74
$n = 1000$	HT	101.47	0.01	2371.18	10.53
	IPW	5.16	0.02	12.71	2.25
	WLS	-2.95	0.37	3.30	1.47
	DR	-48.66	0.08	1370.91	1.81

- Double robust estimator perform better when outcome model is correct but propensity score model is wrong
- Double robust estimator can perform worse when both models are wrong (maybe we should also normalize the weights in DR)

# Variance of IPW estimator

- Researchers have shown that using the estimated propensity score asymptotically results in smaller variance of the IPW estimator (Hirano, Imbens and Ridder, 2003)
- Closed-form sandwich estimator (M-estimator) of variance that takes into account of the uncertainty in estimating the propensity score (Lunceford and Davidian, 2004)
- Bootstrap: Resample units and refit PS and estimate the causal effects every time – computationally intensive for large sample
- In the R example, we show an approximation of the variance ignoring the uncertainty in estimating the propensity score by regression (not too bad, as the estimation of propensity score only involves pre-treatment covariates)

# Bootstrap



- Nonparametric bootstrap:
  - Repeat B times: for each time  $b$ 
    - sample  $N$  units with replacement (or resample the treated and controls separately)
    - Follow the whole procedure (starting from propensity score estimation to estimate the ATE/ATT using IPW)
    - Obtain an IPW estimator  $\hat{t}_{IPW}^{(b)}$
  - Use the histogram of  $\{\hat{t}_{IPW}^{(1)}, \dots, \hat{t}_{IPW}^{(B)}\}$  as the approximated distribution of  $\hat{t}_{IPW}$ 
    - The standard deviation of these estimates approximates the standard error of  $\hat{t}_{IPW}$