

## Introduction to Marginal Structural Models

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### Recommended reading

Hernán MA, Robins JM (2016). *Causal Inference*. Boca Raton: Chapman & Hall/CRC, forthcoming. Preprint available at <http://www.hsph.harvard.edu/miguel-hernan/causal-inference-book/>. Last accessed Aug. 20, 2015

Daniel RM, Cousens SN, De Stavola BL, Kenward MG, Sterne JAC. Methods for dealing with time-dependent confounding. *Stat Med*. 2013; **32**(9):1584-618.

### Copies of these slides are posted at

<http://biostat.mc.vanderbilt.edu/WilliamDupontJan202016>

### Topics

- ❖ Counterfactual events and the definition of causality
- ❖ What is a marginal structural model (MSM)?
- ❖ Simple simulation study with a confounding variable
  - Classical analysis vs MSM
- ❖ Simple longitudinal example
  - Classical analysis vs MSM
- ❖ More complex longitudinal example where classical methods do not work.
- ❖ Applying MSMs to real longitudinal studies

### Counterfactual Events

Let  $Y_i$  be the outcome for the  $i^{th}$  patient,  
 $a$  be some exposure, and  
 $a_i$  be the actual exposure of the  $i^{th}$  patient.

Then  $Y_i^a$  is the potential outcome that would have been observed for the  $i^{th}$  patient if, possibly contrary to fact, she received exposure  $a$ .

### For example:

$Y_i$  denotes whether or not the  $i^{th}$  patient develops bronchiolitis

$a = \begin{cases} 1 & \text{patient receives immunoprophylactic treatment} \\ 0 & \text{patient does not receive this treatment} \end{cases}$

$a_i = \begin{cases} 1 & \text{if } i^{th} \text{ patient receives treatment} \\ 0 & \text{if } i^{th} \text{ patient does not receive treatment} \end{cases}$

$Y_i = Y_i^{a_i}$  is the bronchiolitis outcome that actually happened to the  $i^{th}$  patient given her true treatment.

$Y_i^1$  is the bronchiolitis outcome that would have happened to the  $i^{th}$  patient if she had been treated

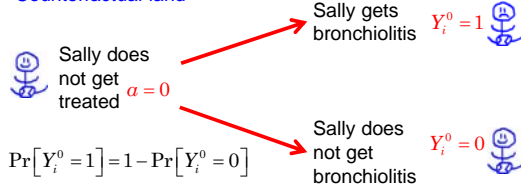
$Y_i^0$  is the bronchiolitis outcome that would have happened to the  $i^{th}$  patient if she had not been treated.

$Y_i^a$  is the counterfactual event that would have occurred given an exposure that may or may not have happened

### Reality



### Counterfactual land



### Causal Relationships

A treatment A has a causal effect on outcome Y if

$$\Pr[Y_i^a] \neq \Pr[Y_i^{a'}]$$

for exposures  $a$  and  $a'$

For example, treatment has a **causal** effect on disease if

$$\Pr[\text{Sally gets disease given she is treated}] \neq \Pr[\text{Sally gets disease given she is not treated}]$$

Note that this is very different from

$$\Pr[\text{Sally gets disease after being treated}] \neq \Pr[\text{Johnny gets disease after not being treated}]$$

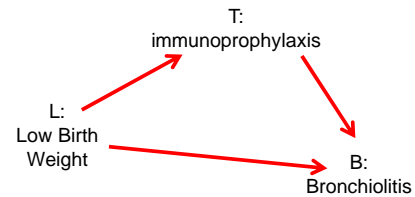
due to possible **confounding by indication**.

Causal inference gurus think of observational data as having **massive amounts of missing data**

If we are lucky and/or diligent we could observe all of the data that actually occurred.

In addition there is at least twice as much missing data from **counterfactual outcomes that did not occur as a consequence of exposures not received**.

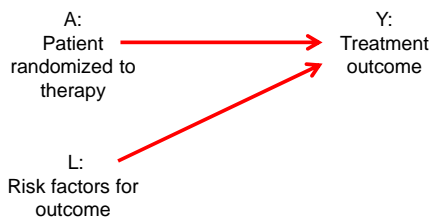
### Directed Acyclic Graphs (DAGs)



Causally related events are indicated by arrows

Here L confounds the causal effect of T on B

### Randomized Clinical Trials



Since treatment assignment is controlled by randomization there can be no causal path between L and A

Hence, any association between A and Y must be causal.

### Exchangeability

A model with treatment  $A$ , covariate  $L$  and outcome  $Y$  is **exchangeable** if the counterfactual outcome  $Y_i^a | L_i$  is independent of  $A_i$ .

$$\text{That is } \Pr[Y_i^a | L_i, A_i] = \Pr[Y_i^a | L_i]$$

In other words, knowing the actual treatment that a patient receives tells me nothing about her counterfactual outcome.

In a RCT treatment is assigned at random, and it does not matter which group receives which treatment. Hence, the treatment groups are exchangeable and

$$\Pr[Y_i^a = 1] = \Pr[Y_i^a = 1 | A_i]$$

### Example of Non-Exchangeability



Here, immunoprophylaxis is much more likely to be given to babies at high risk of bronchiolitis.

Hence, if I know Johnny did not receive T then I know that he is less likely to develop B than if I did not know his actual treatment.

In other words, babies who did, and did not, receive T are **not exchangeable**, since treated babies are inherently at greater risk than untreated babies.

### Marginal Structural Models

A model is

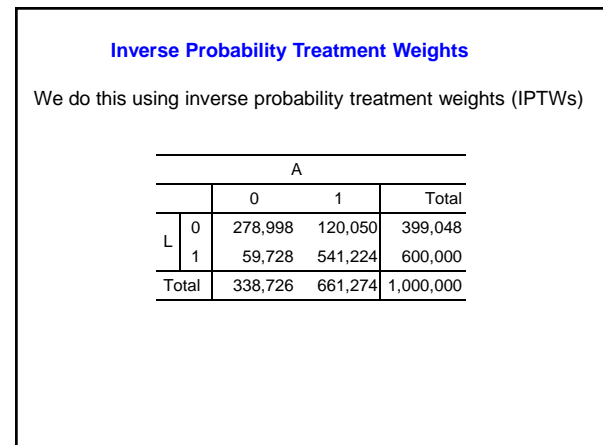
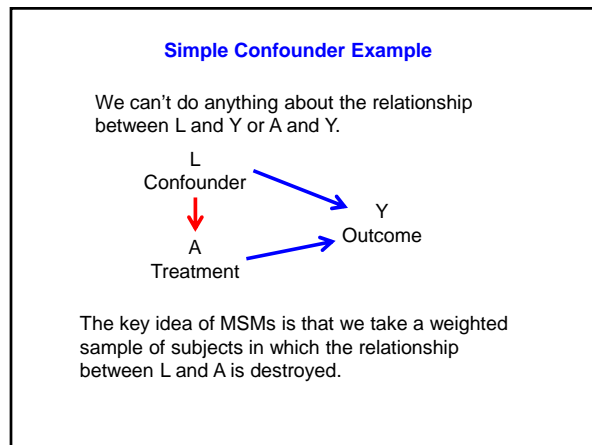
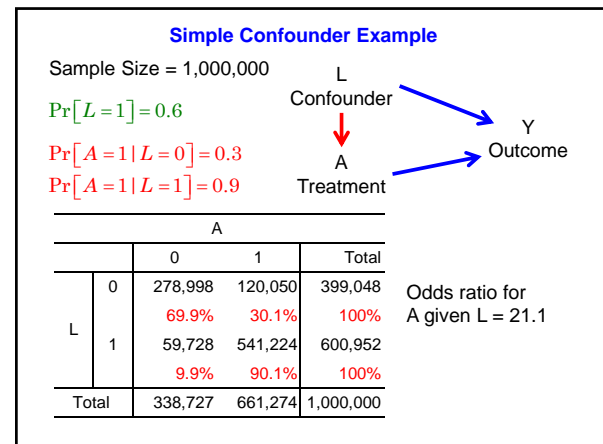
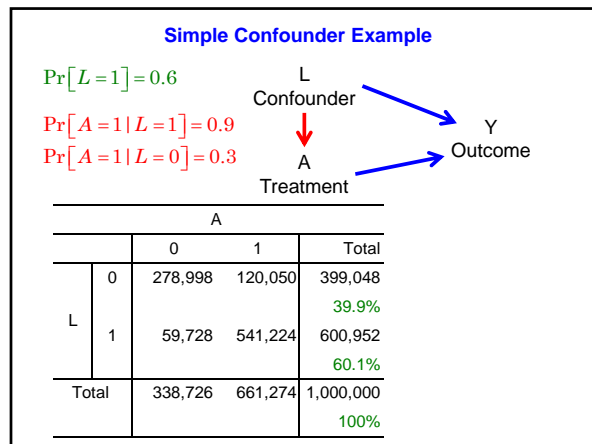
**structural** if it regresses a counterfactual outcome  $Y_i^a$  against treatment and other covariates.

**marginal** if it only involves treatment covariates.

In a RCT if I use simple logistic regression to regress outcome  $Y_i$  against treatment  $A_i$  then I have a **marginal structural model (MSM)**

$$\text{logit}[\pi(Y_i^a)] = \text{logit}[\pi(Y_i)] = \alpha + \beta \times A_i$$

True due to exchangeability      No confounders in model



**Inverse Probability Treatment Weights**

L	A	True Sample Size	Pr[A L]		Estimated IPTW	IPTW × N
			True	Estimated		
0	0	278,998	0.7	0.6992	1/0.6992 = 1.430	399,048
0	1	120,050	0.3	0.3008	1/0.3008 = 3.324	399,048
		399,048				
1	0	59,728	0.1	0.0994	1/0.0994 = 10.061	600,952
1	1	541,224	0.9	0.9006	1/0.9006 = 1.110	600,952
		600,952				
Total		1,000,000	2,000,000			

**Inverse Probability Treatment Weights**

L	A	True Sample Size	Pr[A L]		Estimated IPTW	Weighted Sample Size
			True	Estimated		
0	0	278,998	0.7	0.6992	1/0.6992 = 1.430	199,524
0	1	120,050	0.3	0.3008	1/0.3008 = 3.324	199,524
		399,048				
1	0	59,728	0.1	0.0994	1/0.0994 = 10.061	300,476
1	1	541,224	0.9	0.9006	1/0.9006 = 1.110	300,476
		600,952				
Total		1,000,000	1,000,000			

**True Sample**

A				
		0	1	Total
L	0	278,998	120,050	399,048
	1	59,728	541,224	600,952
Total		338,726	661,274	1,000,000

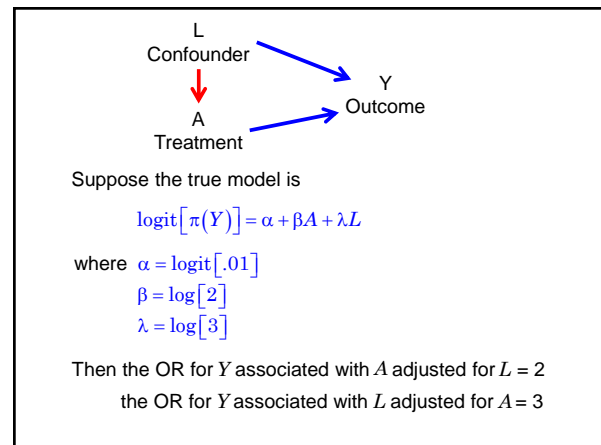
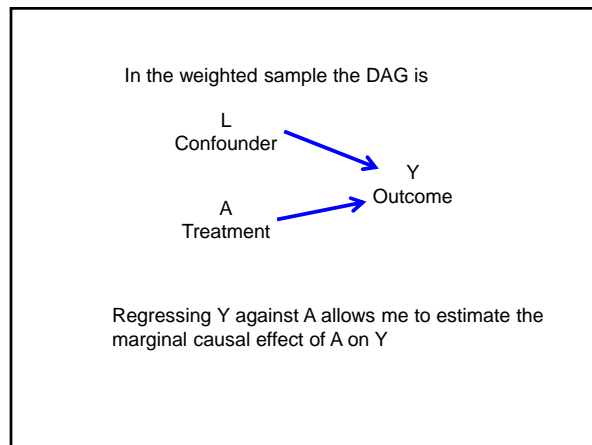
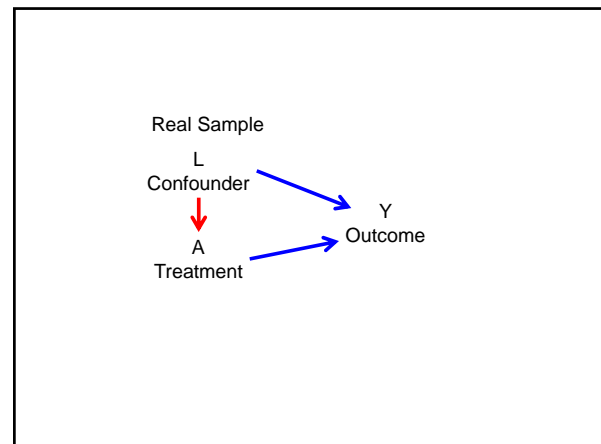
Odds ratio for A given L = 21.1

**Weighted Sample**

A				
		0	1	Total
L	0	199,524	199,524	399,048
	1	300,476	300,476	600,952
Total		500,000	500,000	1,000,000

Odds ratio for A given L = 1



Simulate number of patients with the indicated values of Y given L and A

L	A	Y	Sample Size
0	0	0	276,183
0	0	1	2,815
0	0		278,998
0	1	0	117,656
0	1	1	2,394
0	1		120,050
1	0	0	57,978
1	0	1	1,750
1	0		59,728
1	1	0	510,294
1	1	1	30,930
1	1		541,224
Total			1,000,000

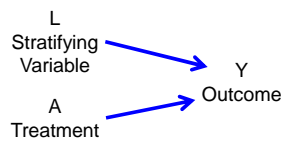
Classical Model			MSM Model	
$\text{logit}[\pi(Y_i)] = \alpha + \beta A_i + \lambda L_i$			$\text{logit}[\pi(Y_i^a)] =$	
applied to real data			$\text{logit}[\pi(Y_i)] = \alpha + \beta A_i$	
			applied to IPTW data with robust standard error estimate	
	Odds Ratio	Standard Error	Odds Ratio	Robust Std. Err.
L	2.973	0.052		
A	2.003	0.037	A	1.998 0.042

Robust standard errors are calculated with the Huber-White sandwich estimator

$\text{logit}[Y_i = 1] = \alpha + \beta A_i$

applied to real data gives OR for Y associated with A = 3.88

### Collapsible Statistics



If the relative risks of Y given A are identical in strata defined by L, then these relative risks also equal the overall (marginal) risk of Y associated with A

Statistics with this property are called **collapsible**

Relative risks are collapsible, odds ratios are not unless they approximate relative risks.

$$L=0 \quad RR = (4/10)/(2/10) = 2$$

		A		
		0	1	Total
Y	0	8	6	14
	1	2	4	6
Total		10	10	20

$$L=1 \quad RR = (6/10)/(3/10) = 2$$

		A		
		0	1	Total
Y	0	7	4	11
	1	3	6	9
Total		10	10	20

$$\text{All Subjects} \quad RR = (10/20)/(5/20) = 2$$

		A		
		0	1	Total
Y	0	15	10	25
	1	5	10	15
Total		20	20	40

Marginal relative risk = within-strata relative risks.

$$L=0 \quad OR = (8/2)/(4/2) = 2$$

		A		
		0	1	Total
Y	0	2	2	4
	1	4	8	12
Total		6	10	16

$$L=1 \quad OR = (2/4)/(2/8) = 2$$

		A		
		0	1	Total
Y	0	8	4	12
	1	2	2	4
Total		10	6	16

$$\text{All Subjects} \quad OR = (10/6)/(6/10) = 2.78$$

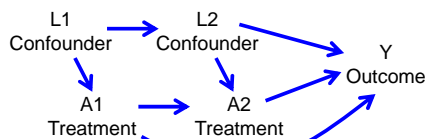
		A		
		0	1	Total
Y	0	10	6	16
	1	6	10	16
Total		16	16	32

Marginal odds ratio  $\neq$  within-strata odds ratios.

In Simulated example I chose the probabilities of Y to be < 6% so that the odds ratios can be interpreted as relative risks

L	A	Y	Sample Size
0	0	0	276,183
0	0	1	2,815
0	0		278,998
0	1	0	117,656
0	1	1	2,394
0	1		120,050
1	0	0	57,978
1	0	1	1,750
1	0		59,728
1	1	0	510,294
1	1	1	30,930
1	1		541,224
Total			1,000,000

### MSMs in Longitudinal Data Analysis



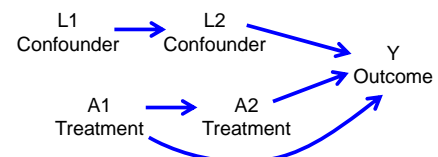
Here A1 affects Y directly and through A2

L1 and L2 confound the effects of A1 and A2 on Y

This model can be analyzed with either classical methods or MSMs.

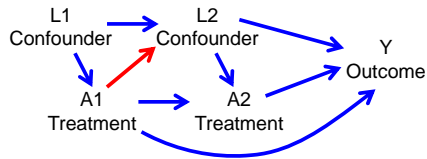
### MSMs in Longitudinal Data Analysis

In the IPT weighted data the model becomes



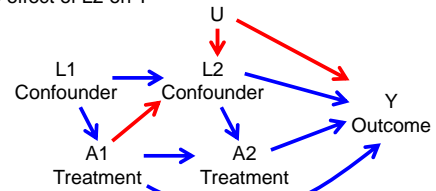
The MSM analysis consists of regressing Y against A1 and A2 in the weighted sample and using robust standard errors.

**Classical methods fail if A1 has a causal effect on L2**



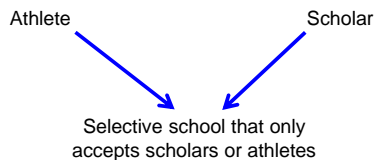
Here L2 confounds the effect of A2 on Y  
I need to adjust for L2 to estimate the causal effect of A2 on Y.  
But L2 is on the causal pathway from A1 to Y  
If I adjust for L2 I block the effect of A1 on Y through L2

Moreover, suppose that unknown variables U confound the effect of L2 on Y



Now the path from A1 to Y through U collides at L2.  
This means that as long as we do not adjust for L2 that the effect of A1 on Y through U is blocked  
If I adjust for L2, I open a path from A1 through U to Y, which means that U confounds the effect of A1 on Y.

Stratifying on (adjusting for) a **collider** induces a relationship (**opens a path**) between its parents



If I attend the school and I am not an athlete then I must be a scholar.

**More Complicated Longitudinal Example**

Sample Size = 1,000,000

$\Pr[A1 = 1] = 0.5$

$\Pr[L = 1 | A1 = 0] = 0.4$

$\Pr[L = 1 | A1 = 1] = 0.8$

$\alpha = \text{logit}[.01]$      $\text{logit}[\pi(Y_i)] = \alpha + \beta A2_i + \lambda L_i$

$\beta = \log[2]$

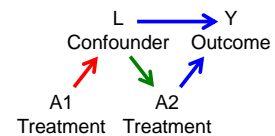
$\lambda = \log[3]$

L is on the causal pathway from A1 to Y

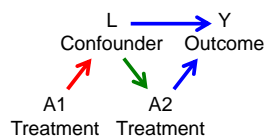
$\Pr[A2 = 1 | L = 0] = 0.3$

$\Pr[A2 = 1 | L = 1] = 0.9$

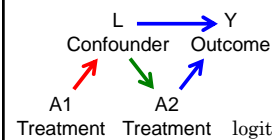
L Strongly confounds the effect of A2 on Y



Simulated number of patients with indicated values of A1, L, A2 and Y



A1	L	A2	Y	Sample Size
0	0	0	0	207,779
0	0	0	1	2,119
0	0	1	0	88,567
0	0	1	1	1,758
0	1	0	0	19,420
0	1	0	1	573
0	1	1	0	169,410
0	1	1	1	10,390
1	0	0	0	69,208
1	0	0	1	694
1	0	1	0	29,216
1	0	1	1	612
1	1	0	0	38,778
1	1	0	1	1147
1	1	1	0	339,600
1	1	1	1	207,229



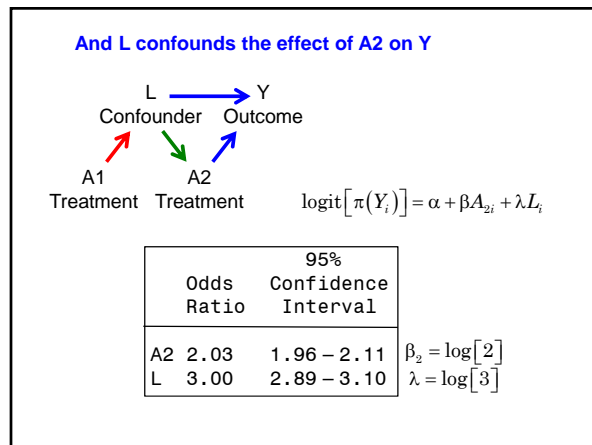
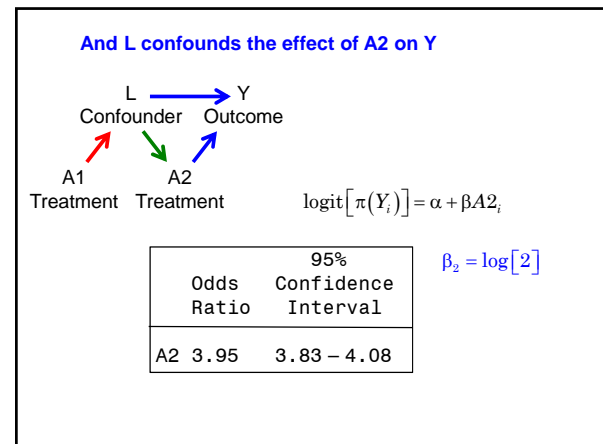
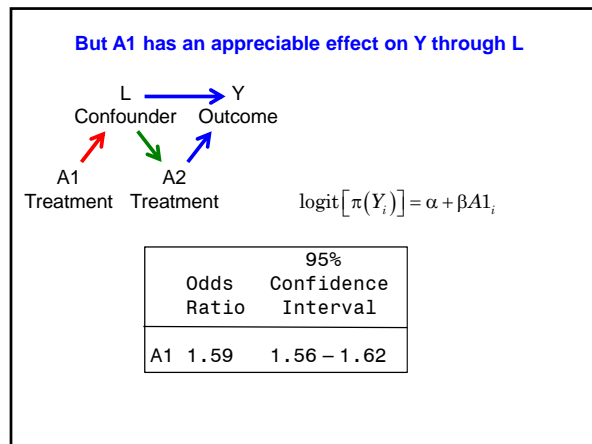
$\text{logit}[\pi(Y_i)] = \alpha + \beta_1 A1_i + \beta_2 A2_i + \lambda L_i$

	Odds Ratio	95% Confidence Interval
A1	1.00	0.98 – 1.02
A2	2.03	1.96 – 2.11
L	3.00	2.89 – 3.11

$\beta_2 = \log[2]$

$\lambda = \log[3]$

The effect of A1 on Y through L is blocked by stratifying on L



**Inverse Probability Treatment Weights**

L	A2	True Sample Size	Estimated Pr[A2 L]	Estimated IPTW	Weighted Sample Size
0	0	279,800	0.6996	1/0.6996 = 1.429	199,976
0	1	120,153	0.3004	1/0.3004 = 3.329	199,976
0		399,953			
1	0	59,918	0.0999	1/0.0999 = 10.014	300,024
1	1	540,129	0.9001	1/0.9001 = 1.111	300,024
1		600,047			
Total		1,000,000			1,000,000

**True Sample**

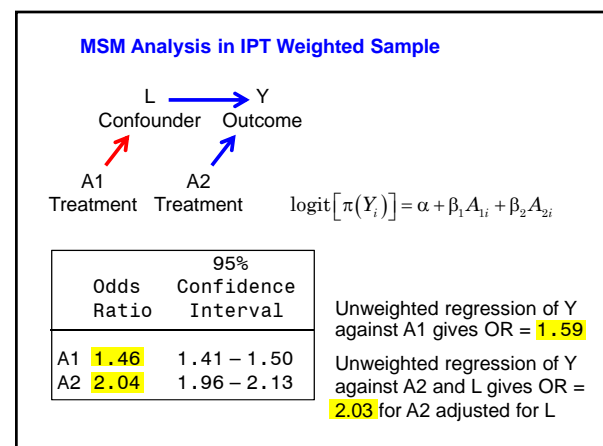
		A2		
		0	1	Total
L	0	279,800	120,153	399,953
	1	59,918	540,129	600,047
Total		339,718	660,282	1,000,000

Odds ratio for A2 given L = 21.0

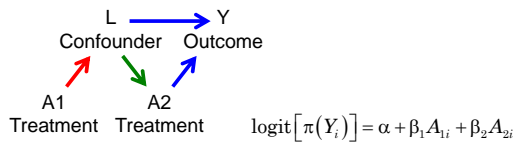
**Weighted Sample**

		A2		
		0	1	Total
L	0	199,976	199,976	399,952
	1	300,024	300,024	600,048
Total		500,000	500,000	1,000,000

Odds ratio for A2 given L = 1



### Classical Regression of Y against A1 and A2 without Adjustment for L



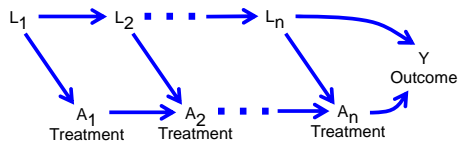
	Odds Ratio	95% Confidence Interval
A1	1.25	1.23 – 1.28
A2	3.72	3.60 – 3.84

This OR is too high

### Applying MSMs to Real Data

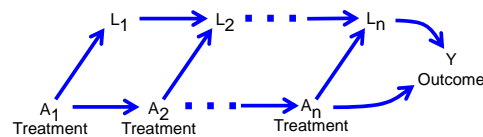
- ❖ For real data we are unlikely to have a model with a perfect fit in which the IPT weighting perfectly blocks the causal link between confounders and treatment.
- ❖ Often use propensity scores to adjust for indication bias.
- ❖ IPT weights are stabilized by multiplying by the unadjusted probability of treatment
- ❖ The results from MSMs are less powerful and are harder to interpret than classical methods. This is particularly true when common events are modeled by odds ratios.
- ❖ My advice is to avoid using them whenever classical methods are viable. They should only be used with some longitudinal studies.

### Longitudinal Data Does not Always Require MSMs



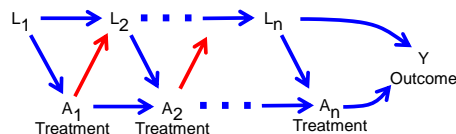
Use classical methods regressing Y against  $A_1, A_2, \dots, A_n$  adjusting for  $L_1, L_2, \dots, L_n$

### Longitudinal Data with Covariates on the Causal Pathway to Y



Use classical methods regressing Y against  $A_1, A_2, \dots, A_n$

### Longitudinal Data with Covariates that are both Confounders and on the Causal Pathway to Y



Use marginal structural models to regress Y against  $A_1, A_2, \dots, A_n$  in an IPT weighted sample

### Thanks

Advice on MSMs  
Bryan Shepherd  
Wayne Ray

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Pingsheng Wu

Programming and data management  
Tan Ding  
Dale Plummer