

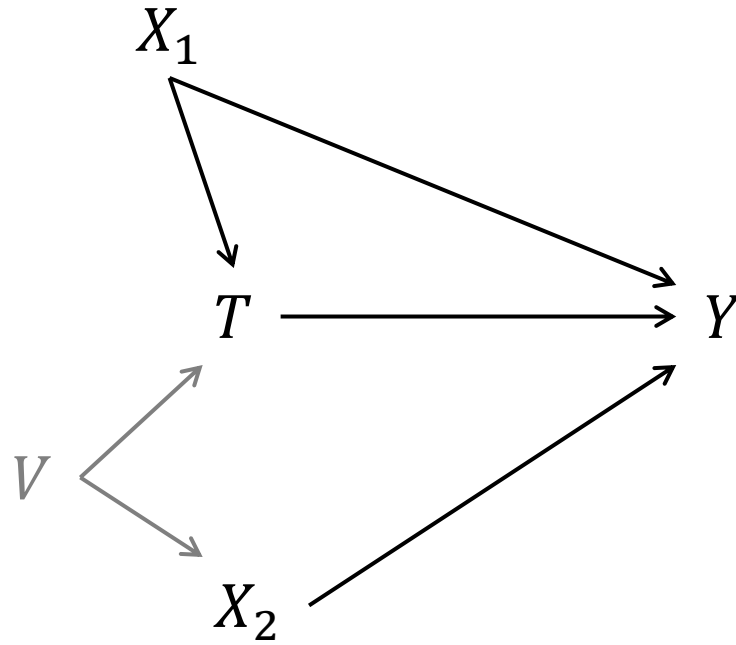
# Sensitivity Analysis for an Unobserved Confounder

Trang Quynh Nguyen (special thanks to Elizabeth Stuart)

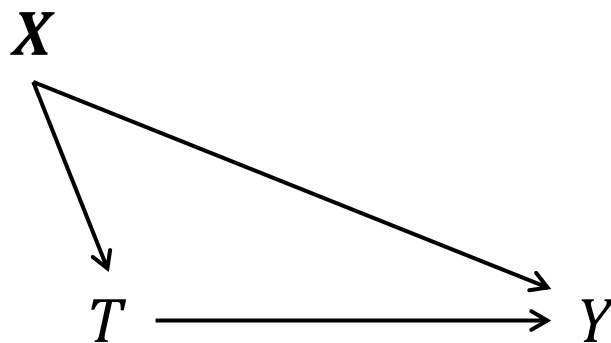
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PHQR772: Advanced Topics in Pharmacoepidemiology  
University of Maryland, 17 Nov 2015

# Confounding

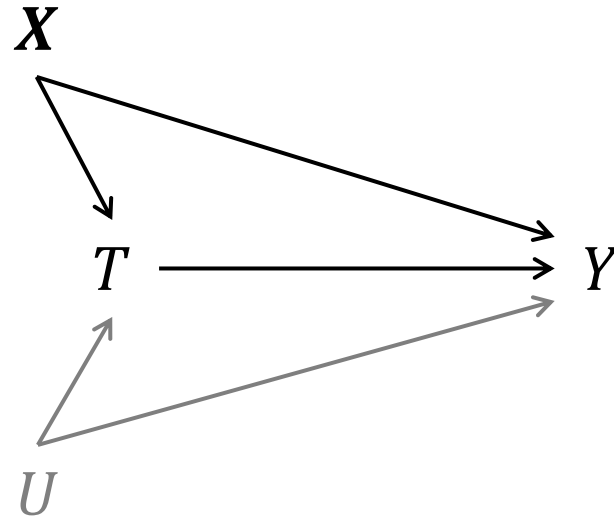


# Adjustment for Observed Confounding

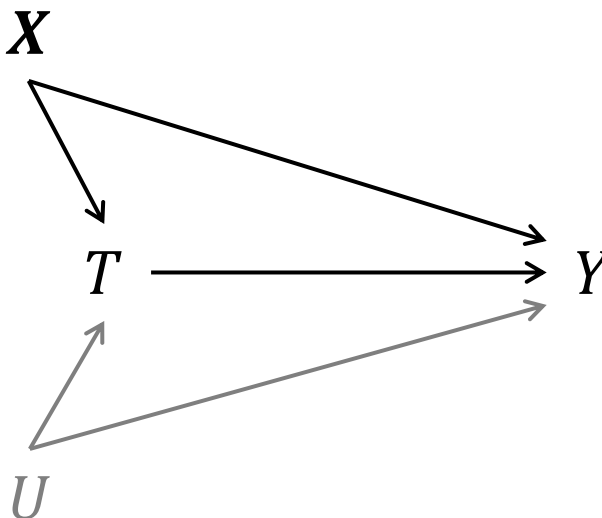


- Adjust for  $X$  via multiple regression (non-causal analysis) or propensity score methods (causal analysis)
- Assumption: No unobserved confounders (no “hidden” bias)

# Unobserved Confounding



# Sensitivity Analysis for an Unobserved Confounder



Questions:

- Given a certain (range of)  $U$ , what is the bias of the  $TY$  effect?
- flip: What would the true  $TY$  effect be?
  - corrected point estimate (and confidence interval?)
- With what  $U$  would the  $TY$  effect go away?
  - statistically non-significant
  - zero point estimate
- related: Could there be a  $U$  that makes the  $TY$  effect go away?

# Main message

- Many flavors
- Depends on specific situation (data, main analysis)
- Depends on question asked

Caveat: Only several methods will be covered to get you started.  
Far from exhaustive.

# Methods to be presented

- Cornfield et al. (1959) smoking and lung cancer sensitivity analysis
- Rosenbaum's approach
  - Sensitivity analysis for subclasses (Rosenbaum & Rubin 1983)
  - Sensitivity analysis for match pairs (Rosenbaum 1987; Gastwirth, Krieger, Rosenbaum 1998)
- 2x2 tables and a binary  $U$  (Greenland 1996; Harding 2003)
- Regression-based methods
  - Simple linear system & omitted variable bias (Harding 2009)
  - Complex non-linear systems (Lin, Psaty & Kronmal 1998)
- VanderWeele & Arah's (2011) bias formulas for general  $Y, T, U$

# Original example: Smoking and Lung Cancer

- R. A. Fisher (1958) thought that the observed relationship between smoking and lung cancer was due to some unobserved genetic factor that made people more susceptible to both.
- Cornfield et al. (1959) analysis apparently changed his mind: that genetic factor would have to be more strongly related to smoking and to lung cancer than anything already observed.

Fisher RA. Cigarettes, cancer and statistics. *Centennial Rev Arts and Sciences*. 2:151, Michigan State University, 1958.

Cornfield, J., Haenszel, W., Hammond, E. C., Lilienfeld, A. M., Shimkin, M. B., & Wynder, E. L. (1959). Smoking and lung cancer: Recent evidence and a discussion of some questions. *Journal of the National Cancer Institute*, 22:173–203.



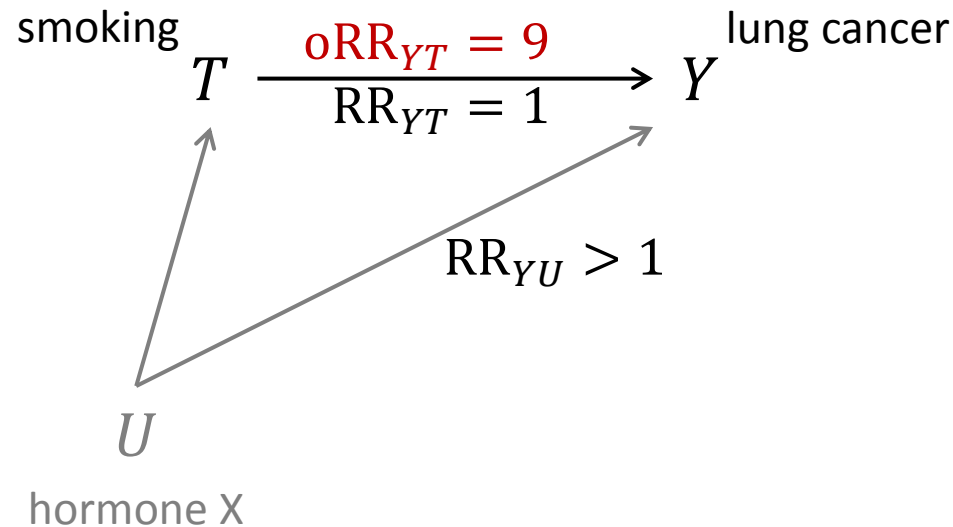
“Thus, if cigarette smokers have 9 times the risk of nonsmokers for developing lung cancer, and this is not because cigarette smoke is a causal agent, but only because cigarette smokers produce hormone X, then the proportion of hormone X-producers among cigarette smokers must be at least 9 times greater than that of nonsmokers. If the relative prevalence of hormone X-producers is considerably less than ninefold, then hormone X cannot account for the magnitude of the apparent effect.” (Cornfield et al., 1959)

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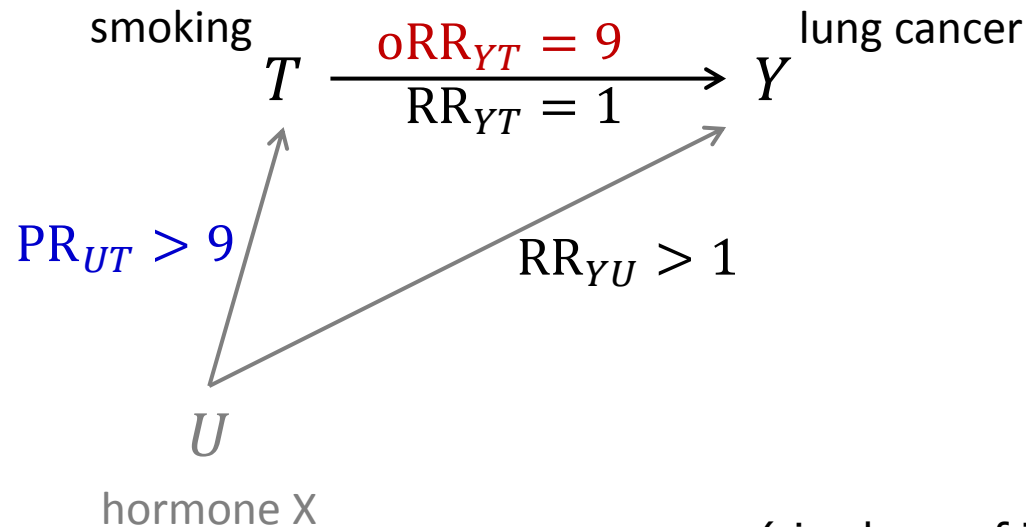
$$\text{smoking } T \xrightarrow{\text{oRR}_{YT} = 9} Y \text{ lung cancer}$$

subscript  $_{YT}$  means  $T$  predicting  $Y$

“Thus, if cigarette smokers have 9 times the risk of nonsmokers for developing lung cancer, and this is not because cigarette smoke is a causal agent, but only because cigarette smokers produce hormone X, then the proportion of hormone X-producers among cigarette smokers must be at least 9 times greater than that of nonsmokers. If the relative prevalence of hormone X-producers is considerably less than ninefold, then hormone X cannot account for the magnitude of the apparent effect.” (Cornfield et al., 1959)



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(simple proof in appendix A)

Cornfield et al. answered which of the following questions?

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Also, need methods that

accommodate both observed confounders and unobserved confounding!

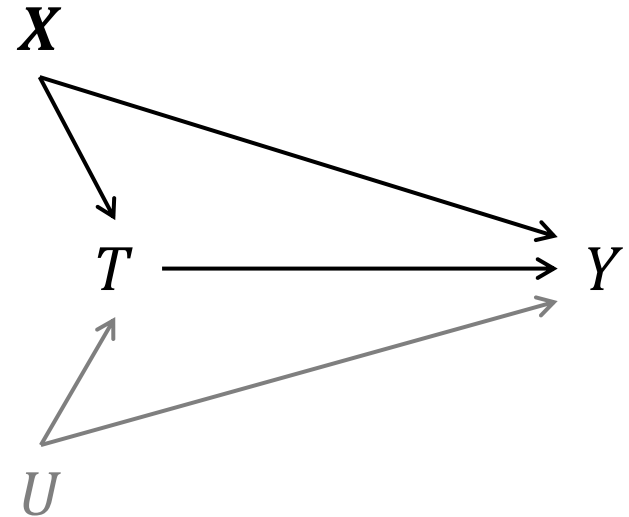
Treatment is not unconfounded given observed  $X$ , but is unconfounded given observed  $X$  and unobserved  $U$ .

# Rosenbaum's approach

use propensity score methods  
to get balance on observed  
confounders  $X$

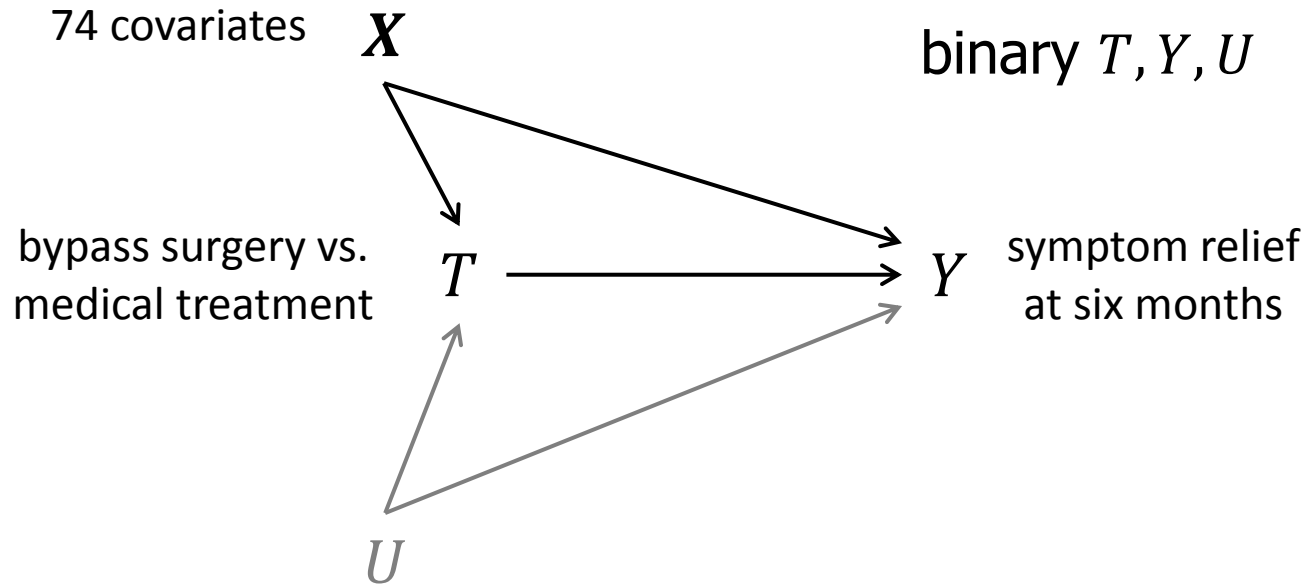
and then

conduct sensitivity analysis on  
an unobserved confounder  $U$



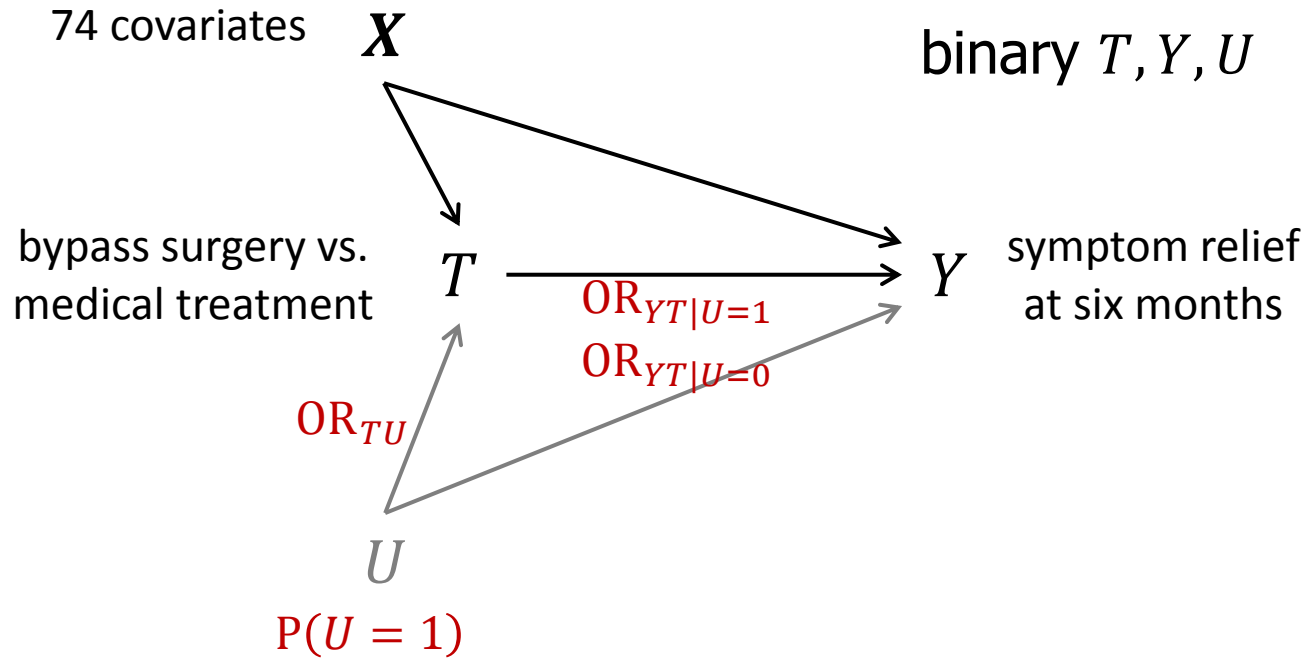


# Rosenbaum & Rubin (1983) with subclassification



Usual analysis: propensity score subclassification to balance  $X$  and estimate the average treatment effect (ATE),  $E[Y_1] - E[Y_0]$  (risk difference of symptom relief at six months)

# Rosenbaum & Rubin (1983) with subclassification



## Sensitivity analysis:

- propensity score subclassification to balance  $X$
- within each subclass, sensitivity analysis on how  $U$  affects the ATE
- average over the subclasses

subclass-specific SA similar in spirit to SA for 2x2 table in Greenland (1996), Harding (2003) & Schneeweiss (2006)

Rosenbaum & Rubin's method answers which of the following questions?

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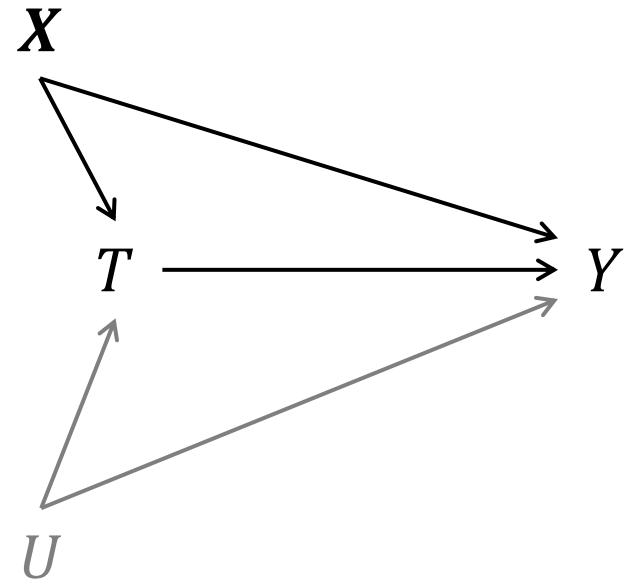
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# Rosenbaum & colleagues with matched pair data

Similar idea:

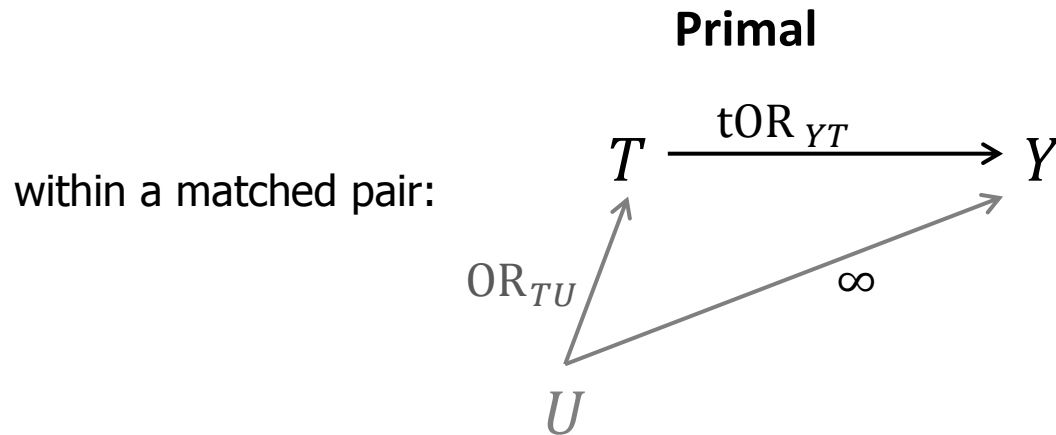
- Matching to balance  $X$  in each pair
- Find values of sensitivity parameters concerning an unobserved  $U$  where the true  $TY$  effect may be no longer statistically significant



Rosenbaum, P. R. (1987). Sensitivity analysis for certain permutational inferences in matched observational studies. *Biometrika*, 74, 13–26.

Gastwirth, J. L., Krieger, A. M., & Rosenbaum, P. R. (1998). Dual and simultaneous sensitivity analysis for matched pairs. *Biometrika*, 85(4), 907–920.

## Three methods for a binary $Y$ : primal, dual and simultaneous

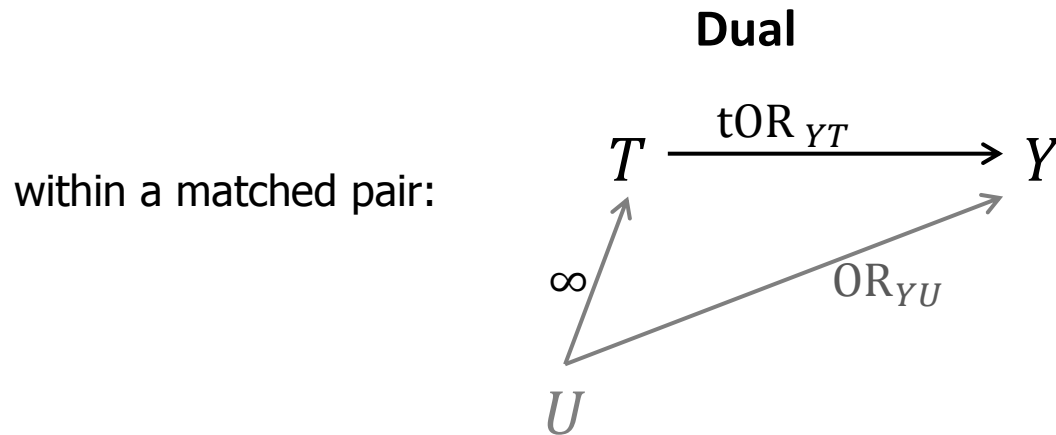


- If no unobserved confounding, the two individuals in a matched pair would have equal probability of treatment assignment
- Due to confounding by some unobserved  $U$  that is extremely predictive of the outcome, their odds of treatment assignment are different,  $OR_{TU} \neq 1$
- Say they are different by at most a factor of  $\Gamma > 1$

$$\frac{1}{\Gamma} \leq OR_{TU} \leq \Gamma$$

- $tOR_{YT}$  is different from  $oOR_{YT}$ , and the true p-value for treatment effect is different from the observed p-value.
- What is the value of  $\Gamma$  where  $tOR_{YT}$  may become statistically non-sig?

## Three methods for a binary $Y$ : primal, dual and simultaneous



- If no unobserved confounding, the two individuals in a matched pair would have equal odds of outcome (for the same treatment)
- Due to confounding by some unobserved  $U$  that is extremely correlated with treatment assignment, their odds of outcome are different,  $OR_{YU} \neq 1$
- Say they are different by at most a factor of  $\Delta > 1$

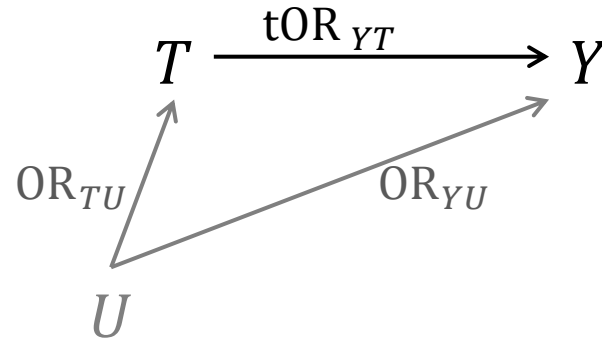
$$\frac{1}{\Delta} \leq OR_{YU} \leq \Delta$$

- $tOR_{YT}$  is different from  $oOR_{YT}$ , and the true p-value for treatment effect is different from the observed p-value.
- What is the value of  $\Delta$  where  $tOR_{YT}$  may become statistically non-sig?

# Three methods for a binary $Y$ : primal, dual and simultaneous

## Simultaneous

within a matched pair:



- If no unobserved confounding, the two individuals in a matched pair would have equal odds of treatment and equal odds of outcome (for the same treatment)
- Due to confounding by some unobserved  $U$ , their odds of treatment are different,  $OR_{TU} \neq 1$ , and their odds of outcome are different,  $OR_{YU} \neq 1$
- Say these differences are bounded by factors of  $\Gamma$  and  $\Delta$  (both  $> 1$ )

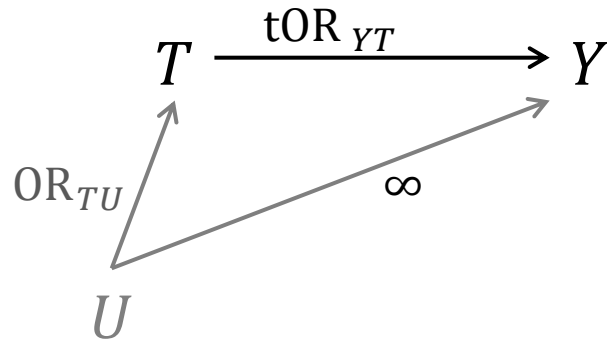
$$\frac{1}{\Gamma} \leq OR_{TU} \leq \Gamma, \quad \frac{1}{\Delta} \leq OR_{YU} \leq \Delta$$

- $tOR_{YT}$  is different from  $oOR_{YT}$ , and the true p-value for treatment effect is different from the observed p-value.
- What are the values of  $\Gamma$  and  $\Delta$  where  $tOR_{YT}$  may be statistically non-sig?

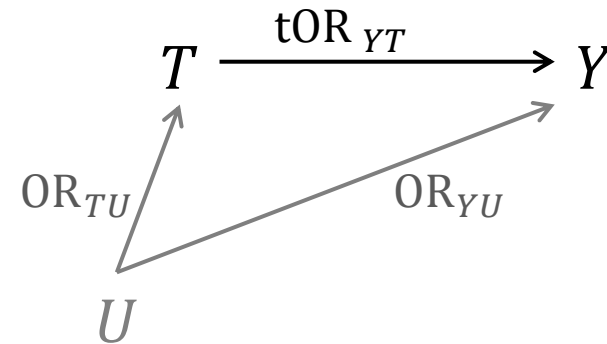


# Three methods for a binary $Y$ : primal, dual and simultaneous

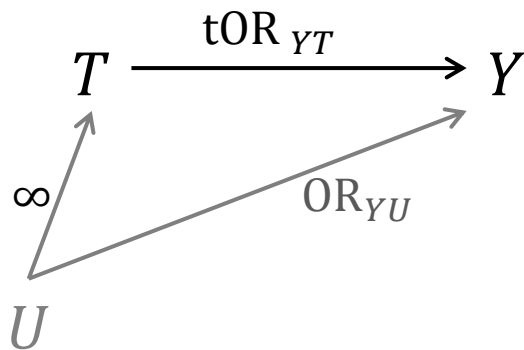
## Primal



## Simultaneous



## Dual



$$\frac{1}{\Gamma} \leq OR_{TU} \leq \Gamma$$

$$\frac{1}{\Delta} \leq OR_{YU} \leq \Delta$$

$$\Gamma > 1, \Delta > 1$$

What are the values of  $\Gamma$  and/or  $\Delta$  where  $tOR_{YT}$  is statistically non-sig?

using a modified McNemar's exact test for paired data

		$T = 0$		
		$Y = 1$	$Y = 0$	
$T = 1$	$Y = 1$	$a$	$b$	$b > c$
	$Y = 0$	$c$	$d$	

use a modified McNemar's exact binomial test for paired data

		$T = 0$		
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$T = 1$	$Y = 1$	$a$	$b$	$b > c$
	$Y = 0$	$c$	$d$	

Liu, Kuramoto & Stuart (2013) example:

		Mother death by accident		
		Child suicide hospitalization	Child no suicide hospitalization	
Mother death by suicide	Child suicide hospitalization	7	226	233
	Child no suicide hospitalization	121	5246	5367
		128	5472	5600

use a modified McNemar's exact binomial test for paired data

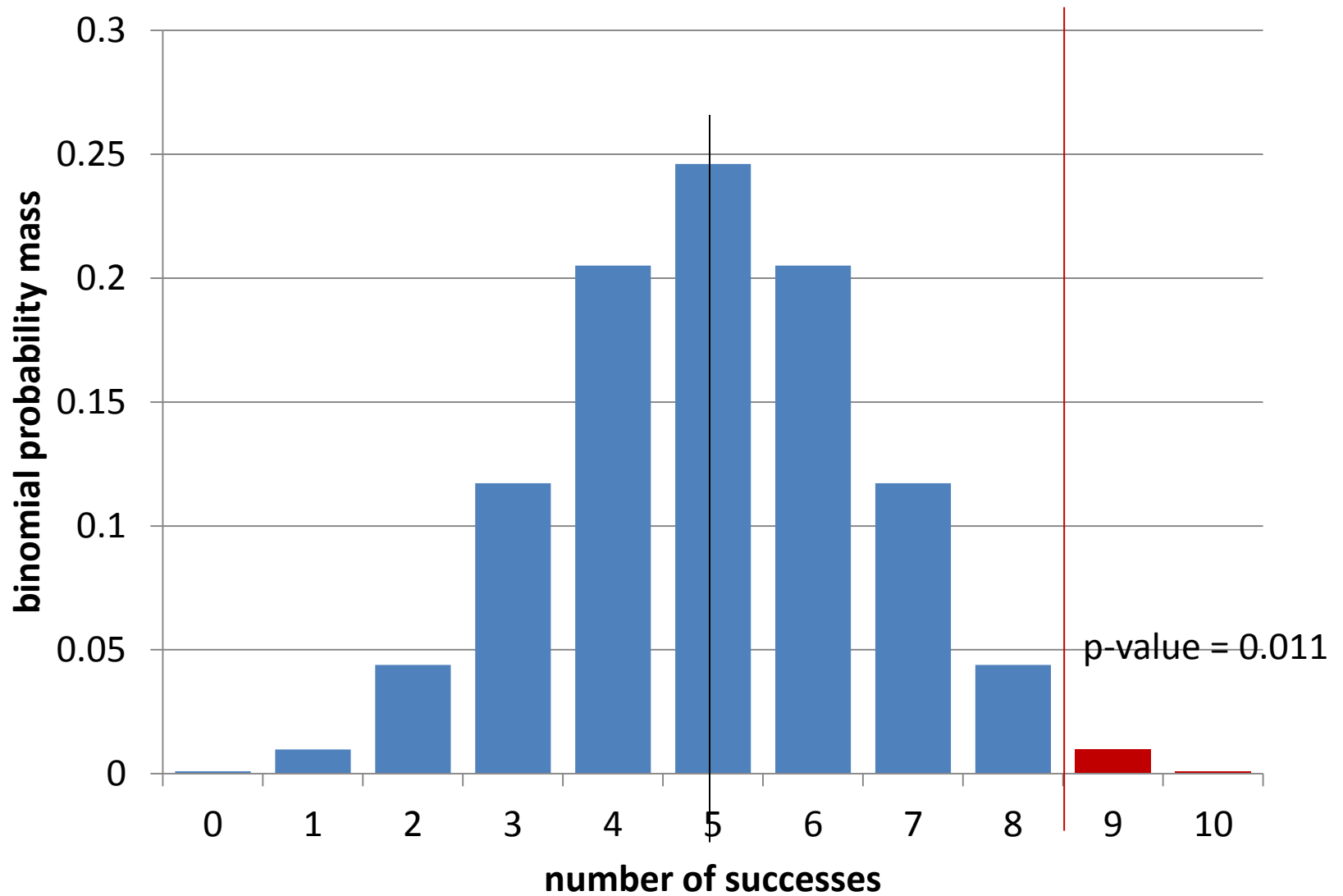
		$T = 0$		
		$Y = 1$	$Y = 0$	
$T = 1$	$Y = 1$	$a$	$b$ <sup>[10]</sup>	$b > c$
	$Y = 0$	$c$ <sub>[01]</sub>	$d$	

Original test:

- $H_0$ : for discordant pair, equal probability (0.5) of each type
- one-sided p-value = probability of observing  $b$  or more pairs of type [10] among  $m = b + c$  discordant pairs

$$p = \sum_{i=b}^m \binom{m}{i} (0.5)^i (0.5)^{m-i}$$

**m=10, b=9, pi=0.5**



Excel function `BINOM.DIST(b,m,pi,0)` (each column); or Stata function `bitest`, R function `binom.test`

use a modified McNemar's exact binomial test for paired data

		$T = 0$			
		$Y = 1$	$Y = 0$		
$T = 1$	$Y = 1$	$a$	$b$ <sup>[10]</sup>	$b > c$	
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Modified test:

- H0: for discordant pairs, probability  $\pi$  of type [10],  $(1 - \pi)$  of type [01]

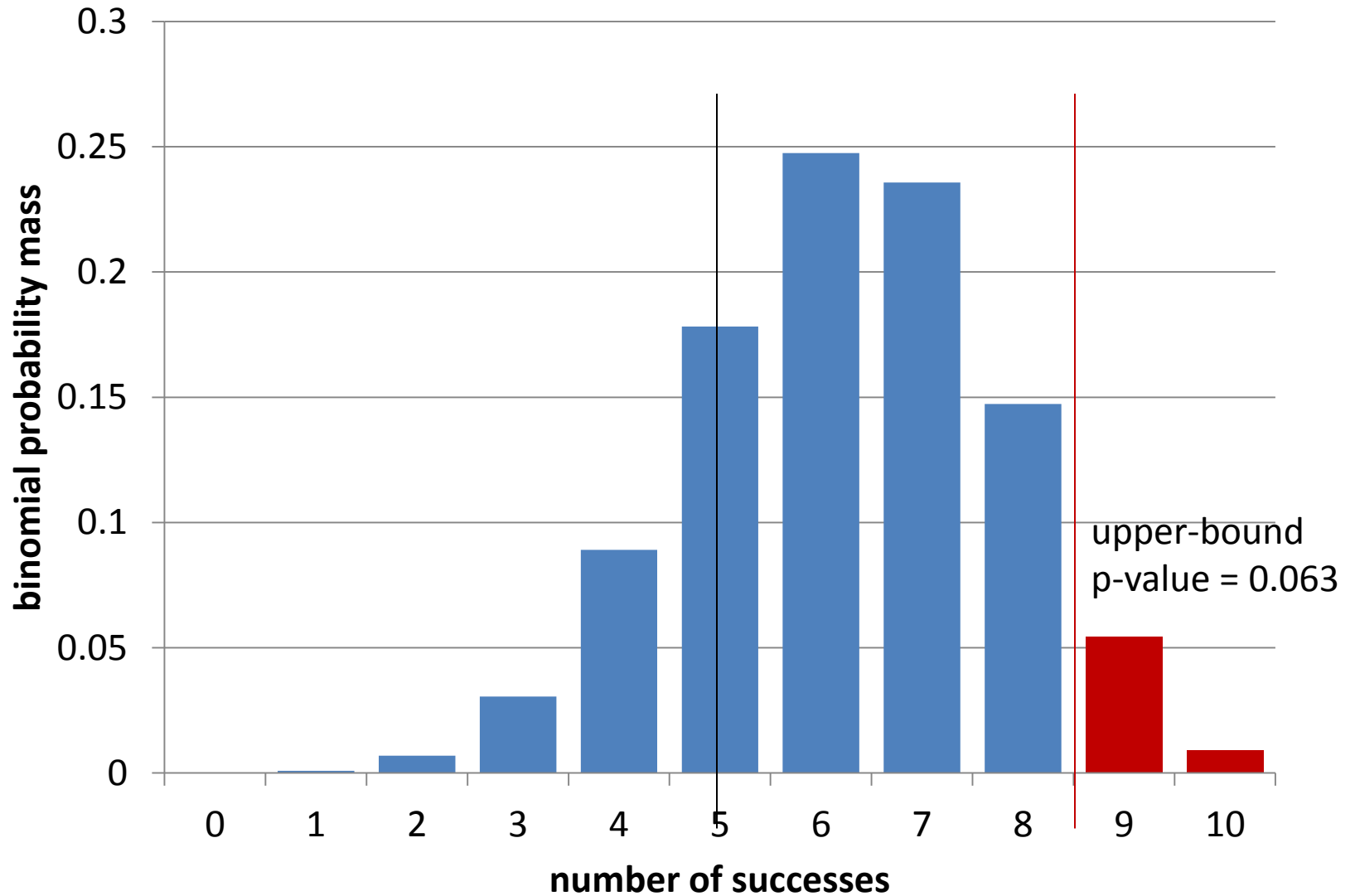
primal:  $\frac{1}{1+\Gamma} \leq \pi \leq \frac{\Gamma}{1+\Gamma}$ ; dual:  $\frac{1}{1+\Delta} \leq \pi \leq \frac{\Delta}{1+\Delta}$ ; simultaneous:  $0.5 \leq \pi \leq \frac{\Gamma}{1+\Gamma} \cdot \frac{\Delta}{1+\Delta} + \frac{1}{1+\Gamma} \cdot \frac{1}{1+\Delta}$

- plugging in the bounds of  $\pi$  gives bounds of p-value:

$$p = \sum_{i=b}^m \binom{m}{i} \pi^i (1 - \pi)^{m-i}$$

- which are the values of  $\Gamma$  and/or  $\Delta$  where p-value **upper-bound**  $\geq 0.05$

**m=10, b=9, upper-bound  $\pi=0.625$  ( $\Gamma=\Delta=3$ )**



Excel function `BINOM.DIST(b,m, $\pi$ , $\theta$ )` (each column); or Stata function `bitest`, R function `binom.test`

# Application to Liu et al. (2013)

Upper-bound of one-sided p-value associated with  $\Gamma$  and  $\Delta$  using Rosenbaum's simultaneous sensitivity analysis

		$\Delta$					
		1.0	2.0	3.0	4.0	5.0	infinity
$\Gamma$	1.0	<.001	<.001	<.001	<.001	<.001	<.001
	2.0	<.001	<.001	.006	.03	<b>.07</b>	<b>.75</b>
	3.0	<.001	.006	<b>.17</b>	<b>.50</b>	<b>.75</b>	<b>1</b>
	4.0	<.001	.03	<b>.50</b>	<b>.89</b>	<b>.98</b>	<b>1</b>
	5.0	<.001	<b>.07</b>	<b>.75</b>	<b>.98</b>	<b>.99</b>	<b>1</b>
	infinity	<.001	<b>.75</b>	<b>1</b>	<b>1</b>	<b>1</b>	<b>1</b>



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$\Gamma$	1.0	<.001	<.001	<.001	<.001	<.001	<.001
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	3.0	<.001	.006	<b>.05</b>	.17	<b>.50</b>	<b>.75</b>
	4.0	<.001	.03	<b>.50</b>	<b>.89</b>	<b>.98</b>	<b>1</b>
	5.0	<.001	<b>.07</b>	<b>.75</b>	<b>.98</b>	<b>.99</b>	<b>1</b>
	infinity	<.001	<b>.75</b>	<b>1</b>	<b>1</b>	<b>1</b>	<b>1</b>

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		1.0	2.0	3.0	4.0	5.0	infinity
$\Gamma$	1.0	<.001	<.001	<.001	<.001	<.001	<.001
	2.0	<.001	<.001	.006	.03	<b>.05</b>	<b>.07</b>
	3.0	<.001	.006	<b>.05</b>	.17	.50	.75
	4.0	<.001	.03	.50	.89	.98	1
	5.0	<.001	<b>.05</b>	.75	.98	.99	1
	infinity	<.001	<b>.05</b>	.75	1	1	1
	infinity	<.001	<b>.05</b>	.75	1	1	1

Rosenbaum's primal, dual and simultaneous methods answer which of the following questions?

- Given a certain (range of)  $U$ , what is the bias of the  $TY$  effect?
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If due to unobserved confounding, between the treated and control units in matched pairs, the odds of treatment differ by a factor of up to 2.8 and the odds of outcome (net of treatment) also differ by a factor of up to 2.8, then the true treatment effect may be statistically non-sig.

## Other comments:

- Brilliant idea!
  - Only two sensitivity parameters
  - Directly relevant when main analysis is matched analysis
- In practice, matching might be done only to obtain balance, with analysis then ignoring that data are matched. Often regression analysis is used to adjust for any remaining imbalance in (observed) confounders – double robustness.
- Need to know the two numbers of discordant pairs
  - Conservative because considers things at the edge:
    - When effect becomes non-sig, not when effect becomes zero
    - Upper-bound of p-value, not simply p-value
    - McNemar's exact test tends to be conservative for small  $m$
  - Can also be interpreted as sensitivity analysis for a binary  $U$
  - The question of one-sided or two-sided test

## Other methods in this genre:

- Matched data, continuous outcome: use a modified Wilcoxon signed rank test (Rosenbaum 1987)
- Sensitivity analysis in the context of matching with multiple controls (Gastwirth, Krieger & Rosenbaum 2000)
- Sensitivity analysis in the context of propensity score weighting (McCaffrey et al. 2004; Ridgeway 2006)

Gastwirth, J. L., Krieger, a M., & Rosenbaum, P. R. (2000). Asymptotic Separability in Sensitivity Analysis. *Journal of the Royal Statistical Society*, 62, 545–555.

McCaffrey, D. F., Ridgeway, G., & Morral, A. (2004). Propensity score estimation with boosted regression for evaluating causal effects in observational studies. *Psychological Methods*, 9(4), 403–425. Retrieved from <http://psycnet.apa.org/journals/met/9/4/403/>

Ridgeway, G. (2006). Assessing the effect of race bias in post-traffic stop outcomes using propensity scores. *Journal of Quantitative Criminology*, 22(1), 1029. Retrieved from <http://www.jstor.org/stable/23367478>

# Methods covered

- Cornfield et al. (1959) smoking and lung cancer sensitivity analysis
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- VanderWeele & Arah's (2011) bias formulas for general  $Y, T, U$

# Greenland's (1996) and Harding's (2003) methods

- Data as 2x2 table, either case-control or cohort

	$Y = 1$ (child suicide hospitalization)	$Y = 0$ (child no suicide hospitalization)
$T = 1$ (mother suicide)	$A$	$B$
$T = 0$ (mother accident)	$C$	$D$

Greenland, S. (1996). Basic methods for sensitivity analysis of biases. *International Journal of Epidemiology*, 25(6), 1107–1116. doi:10.1093/ije/25.6.1107

Harding, D. J. (2003). Counterfactual Models of Neighborhood Effects: The Effect of Neighborhood Poverty on Dropping Out and Teenage Pregnancy. *American Journal of Sociology*, 109(3), 676–719. doi:10.1086/379217



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$T = 1$ (mother suicide)	$A$	$B$
$T = 0$ (mother accident)	$C$	$D$

- For specified plausible binary unobserved  $U$ , unpack into two tables

	$U = 1$		$U = 0$	
	$Y = 1$	$Y = 0$	$Y = 1$	$Y = 0$
$T = 1$	$a_1$	$b_1$	$a_0$	$b_0$
$T = 0$	$c_1$	$d_1$	$c_0$	$d_0$

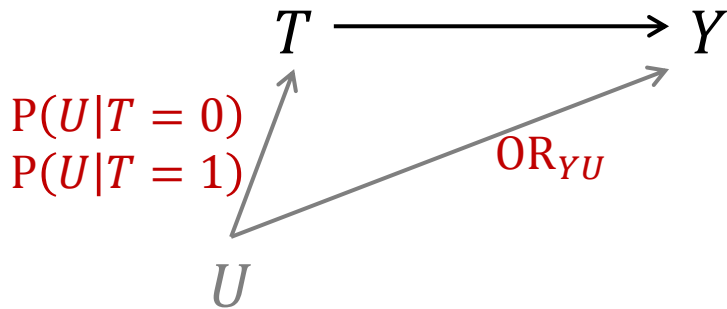
$$a_1 + a_2 = A; \quad b_1 + b_2 = B; \quad c_1 + c_2 = C; \quad d_1 + d_2 = D$$

- and conduct analysis using the two tables or a constructed dataset with  $T, Y, U$  to obtain  $OR_{YT|U}$

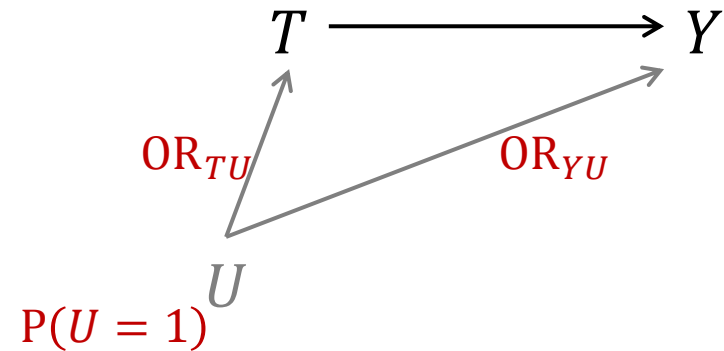
How to specify a plausible range of  $U$ ?

3 sensitivity parameters (4 if allow  $TU$  interaction):

Greenland



Harding



For details on table cells calculation, see Liu et al., which does an excellent job of explaining it for the case without  $TU$  interaction.

Greenland's and Harding's methods can answer which of the following questions?

- Given a certain (range of)  $U$ , what is the bias of the  $TY$  effect?
- flip: What would the true  $TY$  effect be?
  - corrected point estimate (and confidence interval?)
- With what  $U$  would the  $TY$  effect go away?
  - statistically non-significant
  - zero point estimate
- related: Could there be a  $U$  that makes the  $TY$  effect go away?

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- related: Could there be a  $U$  that makes the  $TY$  effect go away?

- Easy to understand
- Relatively easy to implement
- Corrected point estimate and confidence interval! 😊
  
- How to deal with observed confounders  $X$ ?  
Balance  $X$  using propensity score methods and then conduct sensitivity analysis for  $X$ -balanced samples (or subsamples)
  - Suclassification and then sensitivity analysis within subclasses (Rosenbaum & Rubin 1983)
  - Matching (or weighting) and then use the matched/weighted sample as an  $X$ -balanced sample (ignoring matched) for sensitivity analysis (Harding 2003; Liu et al. 2013)

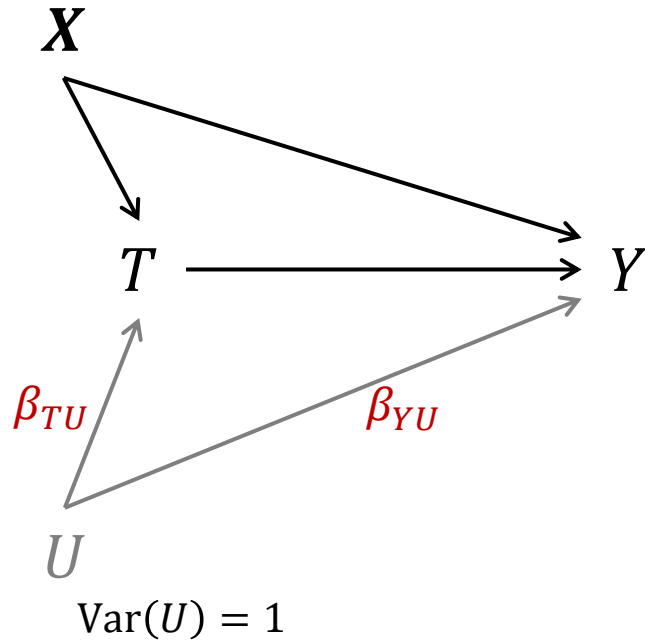
# Schneeweiss (2006)

- class critique

# Methods covered

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# A regression-based approach: sensitivity analysis based on omitted variable bias (Harding 2009)



- $T$  is binary (smoking) – my example, not Harding's.
- $Y$  is binary or continuous (obesity/weight).
- $U$  is continuous (depressive symptom severity), variance fixed at 1, independent of  $X$  (think  $X$  have been "regressed out" of  $U$ ).
- Rely on linear models

$$E[Y] = \alpha_Y + \beta_{YX}X + \beta_{YT}T + \beta_{YU}U$$

$$E[T] = \alpha_T + \beta_{TX}X + \beta_{TU}U$$

- Need to standardize  $T$ , get bias  $\beta_{TU}\beta_{YU}$

$$t\beta_{YT} = o\beta_{YT} - \beta_{TU}\beta_{YU}$$



## Comments:

- Would like to not standardize  $T$

Simple fix: Shift the representation of the  $UT$  relationship from  $\beta_{TU}$  (RD of treatment associated with one SD difference in  $U$ ) to  $\beta_{UT}$  (the difference in mean  $U$  comparing  $T = 1$  and  $T = 0$ ).

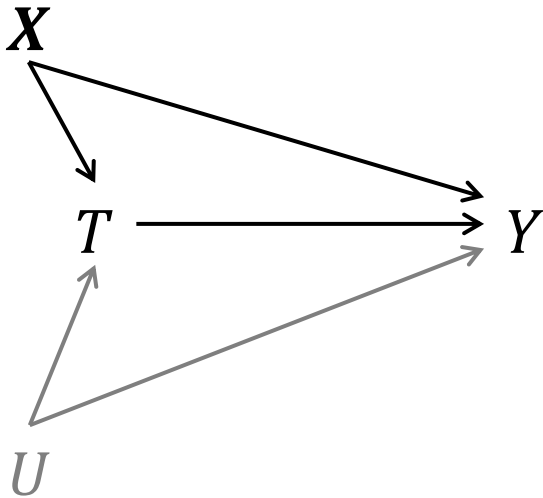
Then

$$t\beta_{YT} = o\beta_{YT} - \beta_{UT}\beta_{YU}$$

Note that this difference in means is not a causal effect (causation is assumed to be the opposite direction).

- Need to be explicit about the assumptions of the linear system

# More regression based: Lin, Psaty & Kronmal (1998)



Very interesting paper!

- $T$  binary
- $Y$  binary (log-linear or logistic) or survival time
- $U$  binary or normal
- allowing  $TU$  interaction

Complicated equations are simplified based on the assumption that  $U$  and  $X$  are independent conditional on  $T$ , which is violated because  $T$  is a collider (Hernan & Robins 1999).

If no  $X$ , reduce to simpler results.

VanderWeele & Arah note that this paper offers an alternative assumption that the conditional mean of  $U$  is additive in  $X$  and  $T$  which is helpful for deriving the bias.

Lin, D. Y., Psaty, B. M., & Kronmal, R. A. (1998). Assessing the sensitivity of regression results to unmeasured confounders in observational studies. *Biometrics*, 54(3), 948–963. doi:10.2307/2533848

Hernan, M. A., & Robins, J. M. (1999). Letter to the Editor: Assessing the sensitivity of regression results to unmeasured confounders in observational studies. *Biometrics*, 55, 1316–1317.

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# VanderWeele & Arah's (2011) general bias formulas

Very general!

For simplicity, let  $U$  be binary, and consider ATE on the additive scale.

- Each individual has a potential outcome under treatment,  $Y_1$ , and a potential outcome under control,  $Y_0$ .
- Treatment effect is:  $ATE = E[Y_1] - E[Y_0]$
- Treatment assignment is unconfounded (as good as random) given observed  $X$  and unobserved  $U$ .

$$ATE = \sum_{\mathbf{x}} \sum_u \{E[Y|T = 1, \mathbf{x}, u] - E[Y|T = 0, \mathbf{x}, u]\} P(u|\mathbf{x}) P(\mathbf{x}).$$

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$$ATE = \sum_{\mathbf{x}} \sum_u \{E[Y|T = 1, \mathbf{x}, u] - E[Y|T = 0, \mathbf{x}, u]\}P(u|\mathbf{x})P(\mathbf{x}).$$

- Adjusting for  $X$  but not  $U$  gives

$$\sum_{\mathbf{x}} \{E[Y|T = 1, \mathbf{x}] - E[Y|T = 0, \mathbf{x}]\}P(\mathbf{x}).$$

- Bias is the difference between these two quantities.

General formula:

$$\begin{aligned} \text{bias} = & \\ & \sum_{\mathbf{x}} \{E[Y|T = 1, U = 1, \mathbf{x}] - E[Y|T = 1, U = 0, \mathbf{x}]\} [P(U = 1|T = 1, \mathbf{x}) - P(U = 1|\mathbf{x})] P(\mathbf{x}) - \\ & \sum_{\mathbf{x}} \{E[Y|T = 0, U = 1, \mathbf{x}] - E[Y|T = 0, U = 0, \mathbf{x}]\} [P(U = 1|T = 1, \mathbf{x}) - P(U = 1|\mathbf{x})] P(\mathbf{x}) \end{aligned}$$

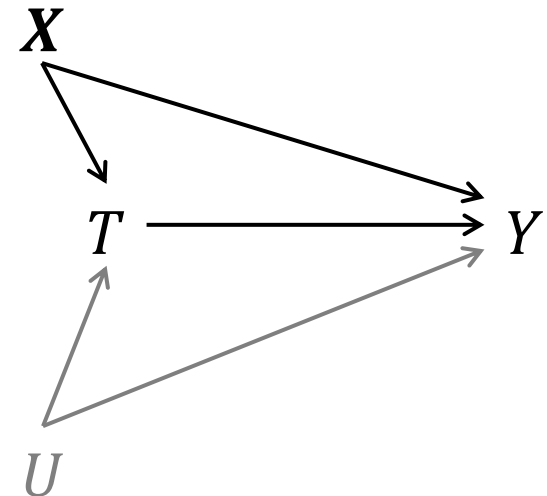
General formula:

$$\text{bias} = \sum_{\mathbf{x}} \{E[Y|T = 1, U = 1, \mathbf{x}] - E[Y|T = 1, U = 0, \mathbf{x}]\} [P(U = 1|T = 1, \mathbf{x}) - P(U = 1|\mathbf{x})] P(\mathbf{x}) - \sum_{\mathbf{x}} \{E[Y|T = 0, U = 1, \mathbf{x}] - E[Y|T = 0, U = 0, \mathbf{x}]\} [P(U = 1|T = 1, \mathbf{x}) - P(U = 1|\mathbf{x})] P(\mathbf{x})$$

$\uparrow$   
*UY* given *T* within *X* stratum
 $\uparrow$   
*UT* within *X* stratum

Strata could be strata of *X* (eg female & college)  
or strata (subclasses) of propensity score.

Complicated, but simplifies in some cases.



If simplification 1: within  $X$  stratum, no  $UT$  interaction

$$\text{bias} = \sum_x \{E[Y|U = 1, T, \mathbf{x}] - E[Y|U = 0, T, \mathbf{x}]\} [P(U = 1|T = 1, \mathbf{x}) - P(U = 1|T = 0, \mathbf{x})] P(\mathbf{x})$$

plus simplification 2: the  $UY$  relationship given  $T$  does not vary across  $X$  strata

$$\text{bias} = \{E[Y|U = 1, T, \mathbf{X}] - E[Y|U = 0, T, \mathbf{X}]\} \sum_x [P(U = 1|T = 1, \mathbf{x}) - P(U = 1|T = 0, \mathbf{x})] P(\mathbf{x})$$

or plus simplification 3: the  $UT$  relationship does not vary across  $X$  strata

$$\text{bias} = [P(U = 1|T = 1, \mathbf{X}) - P(U = 1|T = 0, \mathbf{X})] \sum_x \{E[Y|U = 1, T, \mathbf{x}] - E[Y|U = 0, T, \mathbf{x}]\} P(\mathbf{x})$$

or plus both simplifications 2 and 3

$$\text{bias} = \{E[Y|U = 1, T, \mathbf{X}] - E[Y|U = 0, T, \mathbf{X}]\} [P(U = 1|T = 1, \mathbf{X}) - P(U = 1|T = 0, \mathbf{X})]$$



How does this translate to sensitivity parameters?

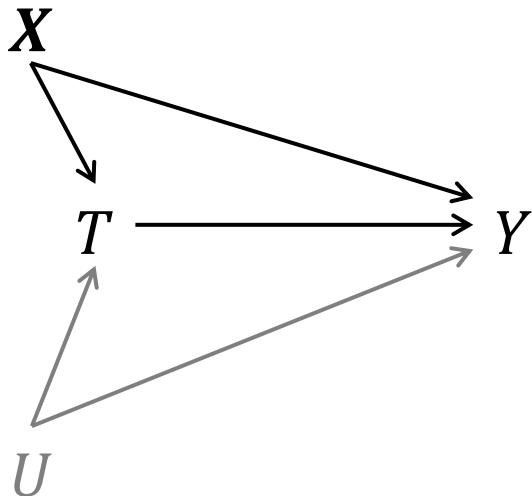
How does it relate to prior methods?

Consider the simplest formula, with all three simplifications,

$$\text{bias} = \{E[Y|U = 1, T, \mathbf{X}] - E[Y|U = 0, T, \mathbf{X}]\} [P(U = 1|T = 1, \mathbf{X}) - P(U = 1|T = 0, \mathbf{X})]$$

$RD_{YU|T, \mathbf{X}}$

$PD_{UT|\mathbf{X}}$



How does this translate to sensitivity parameters?

How does it relate to prior methods?

Consider the simplest formula, with all three simplifications,

$$\text{bias} = \{E[Y|U = 1, T, \mathbf{X}] - E[Y|U = 0, T, \mathbf{X}]\} [P(U = 1|T = 1, \mathbf{X}) - P(U = 1|T = 0, \mathbf{X})]$$

$RD_{YU|T, \mathbf{X}}$

$PD_{UT|\mathbf{X}}$

In the  $X$  stratum specific case (or no  $X$  case), alternatives to specifying  $PD_{UT|x}$ :

- To combine a relative measure of association

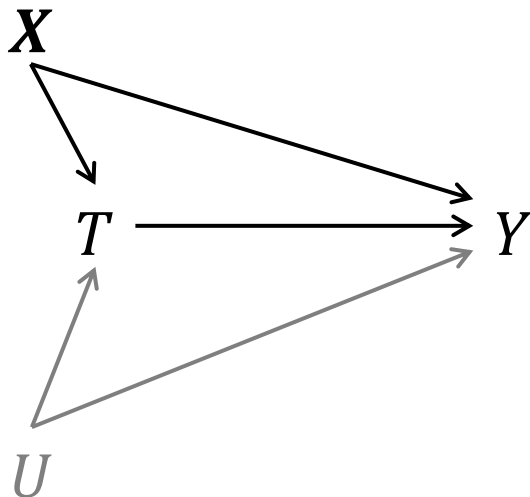
$PR_{UT|x}$  or  $RR_{TU|x}$  or  $OR_{TU|x}$

and a prevalence

$P(U = 1|T = 0, \mathbf{x})$  or  $P(U = 1|\mathbf{x})$

- To specify two prevalences

$P(U = 1|T = 0, \mathbf{x})$  or  $P(U = 1|T = 1, \mathbf{x})$



With fewer simplifications, more parameters!

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# Other approaches

- Simulation
  - Arah, O., Chiba, Y., & Greenland, S. (2008). Bias formulas for external adjustment and sensitivity analysis of unmeasured confounders. *Annals of Epidemiology*, *18*(8), 637–46. doi:10.1016/j.annepidem.2008.04.003
  - Steenland, K., & Greenland, S. (2004). Monte Carlo Sensitivity Analysis and Bayesian Analysis of Smoking as an Unmeasured Confounder in a Study of Silica and Lung Cancer. *American Journal of Epidemiology*, *160*(4), 384–392. doi:10.1093/aje/kwh211
- Bayesian methods
  - Steenland & Greenland (2004)
  - McCandless, L. C., Gustafson, P., & Levy, A. (2007). Bayesian sensitivity analysis for unmeasured confounding in observational studies. *Statistics in Medicine*, *26*, 2331–2347. doi:10.1002/sim
- Using external data to adjust results
  - Stürmer, T., Schneeweiss, S., Avorn, J., & Glynn, R. J. (2005). Adjusting effect estimates for unmeasured confounding with validation data using propensity score calibration. *American Journal of Epidemiology*, *162*(3), 279–89. doi:10.1093/aje/kwi192
- Design sensitivity
  - Zubizarreta, J. R., Cerdá, M., & Rosenbaum, P. R. (2013). Effect of the 2010 Chilean earthquake on posttraumatic stress: reducing sensitivity to unmeasured bias through study design. *Epidemiology*, *24*(1), 79–87. doi:10.1097/EDE.0b013e318277367e