

Sensitivity Analyses

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Sensitivity

What is “sensitivity”?

When inputs change, do outputs change?

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- ▶ With different variables in model, does parameter of interest change?
- ▶ With different assumptions about error structures, does causal mediation estimate change?
- ▶ With different data collected, would causal conclusion change?

Sensitivity to Model Specification

Should we trust our model?

Suppose I present observational results:

	Coefficient
≥ 1000 auto workers	0.87 (0.39)
DW-NOMINATE	-5.04 (0.53)
Ford/Chrysler/GM PAC Contribs (log)	0.15 (0.05)
AFL-CIO PAC Contribs (log)	0.09 (0.04)
Intercept	-0.14 (0.30)
N	406
AIC	258.59
BIC	338.72
$\log L$	-109.30

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What would be your questions?

Multiplicity of Models

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- ▶ So, typically, we show

Multiplicity of Models

	Model 1	Model 2	Model 3	Model 4
Intercept	2.12 (0.24)	0.61 (0.20)	1.26 (0.31)	-0.14 (0.30)
≥ 1000 auto workers	0.98 (0.34)	1.14 (0.37)	0.74 (0.36)	0.87 (0.39)
Republican	(0.33)		(0.42)	
DW-NOMINATE		-4.87 (0.42)		-5.04 (0.53)
Ford/Chrysler/GM PAC Contribs (log)			0.14 (0.05)	0.15 (0.05)
AFL-CIO PAC Contribs (log)			0.14 (0.04)	0.09 (0.04)
N	407	406	407	406
AIC	301.48	268.76	284.11	258.59
BIC	349.58	316.83	364.29	338.72
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Rather than small set of substantively-informed models, just show them all!

Example 1: Congressional Auto Industry Support

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Estimate relationship

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Claim: **Local econ interests** at least on par w/ corporate campaign contributions, lobbying, public positions.

Moore, Powell, and Reeves (2013)

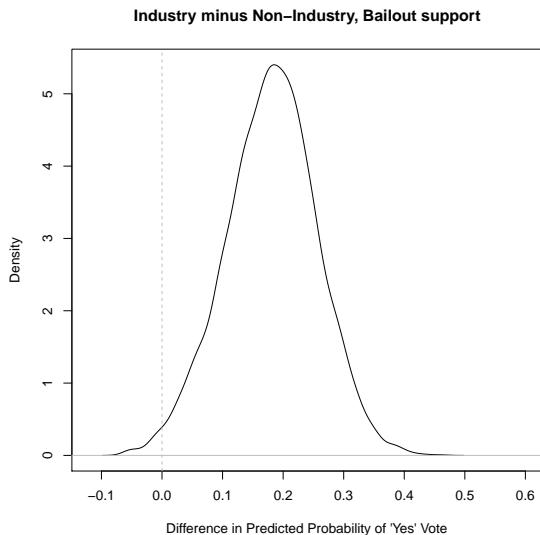


Figure 1: First diffs, predicted prob MoC supports auto bailout, member from industry v. non-industry district, other vars at means.

Moore, Powell, and Reeves (2013)

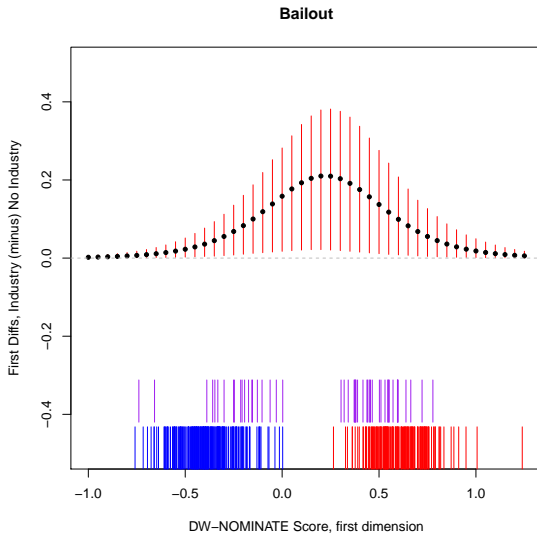


Figure 2: First diffs, industry v. non-industry district member prob of supporting bailout positive at any value of DW-NOMINATE score.

Insensitivity to Specification

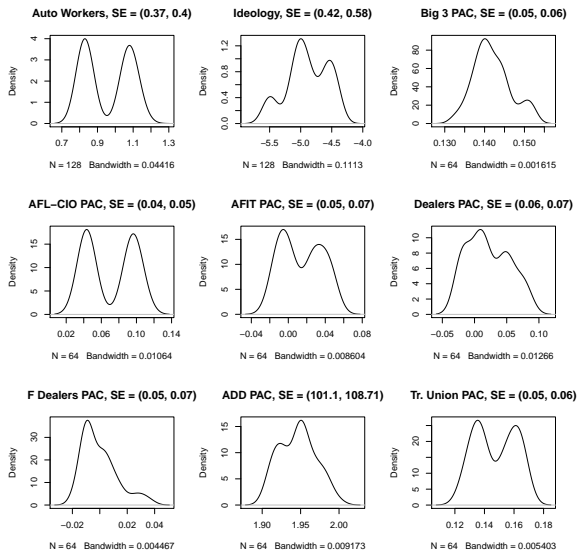


Figure 3: Industry presence coef always positive in Bailout logistic regressions. Coef densities w/ industry presence and

Insensitivity to Specification

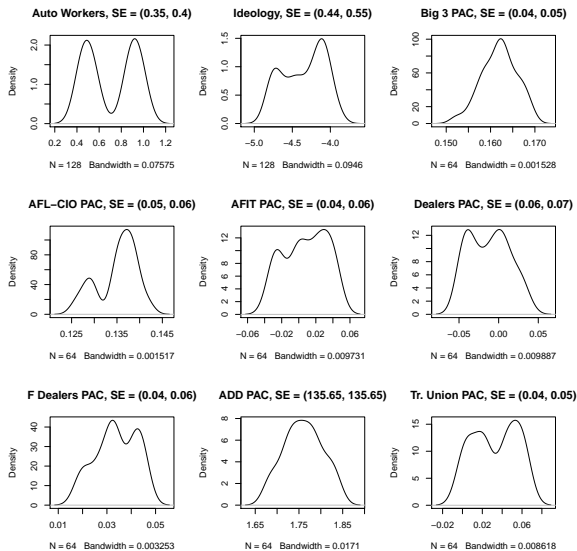


Figure 4: Industry presence coef always positive in Cash for Clunkers logistic regressions. Coef densities w/ industry presence and

Implementation

```
library(olsrr)
```

- ▶ Estimate (all) linear models

Implementation

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- ▶ Estimate (all) linear models
- ▶ Provide model fit stats

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Hebbali (2024)

Example 2: Social Pressure Mailers

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Dear Registered Voter:

WHAT IF YOUR NEIGHBORS KNEW WHETHER YOU VOTED?

Why do so many people fail to vote? We've been talking about the problem for years, but it only seems to get worse. This year, we're taking a new approach. We're sending this mailing to you and your neighbors to publicize who does and does not vote.

The chart shows the names of some of your neighbors, showing which have voted in the past. After the August 8 election, we intend to mail an updated chart. You and your neighbors will all know who voted and who did not.

DO YOUR CIVIC DUTY — VOTE!

MAPLE DR	Aug 04	Nov 04	Aug 06
9995 JOSEPH JAMES SMITH	Voted	Voted	_____
9995 JENNIFER KAY SMITH		Voted	_____
9997 RICHARD B JACKSON		Voted	_____
9999 KATHY MARIE JACKSON		Voted	_____
9999 BRIAN JOSEPH JACKSON		Voted	_____
9991 JENNIFER KAY THOMPSON		Voted	_____
9991 BOB R THOMPSON		Voted	_____
9993 BILL S SMITH			_____
9989 WILLIAM LUKE CASPER		Voted	_____
9989 JENNIFER SUE CASPER		Voted	_____
9987 MARIA S JOHNSON	Voted	Voted	_____

Example 2: Social Pressure Mailers

```
library(qss)
data(social)
```

```
social |> select(-yearofbirth) |> head()
```

	sex	primary2004	messages	primary2006	hhsiz	age
1	male	0	Civic Duty	0	2	65
2	female	0	Civic Duty	0	2	59
3	male	0	Hawthorne	1	3	55
4	female	0	Hawthorne	1	3	56
5	female	0	Hawthorne	1	3	24
6	male	0	Control	0	3	25

Example 2: Social Pressure Mailers

```
lm_out <- lm(primary2006 ~ messages + sex + age +  
             primary2004 + hhsize, data = social)  
  
all_lm_social <- ols_step_all_possible(lm_out)$result  
  
dim(all_lm_social)
```

```
[1] 31 15
```

```
head(all_lm_social)
```

	mindex	n	predictors	rsquare	adjr
4	1	1	primary2004	0.0261502651	0.0261470812 0.457
3	2	1	age	0.0167659386	0.0167627240 0.459
1	3	1	messages	0.0032825640	0.0032727879 0.462
5	4	1	hhsize	0.0025142362	0.0025109749 0.462
2	5	1	sex	0.0001863186	0.0001830498 0.463
13	6	2	age primary2004	0.0409175309	0.0409112596 0.453

Example 2: Social Pressure Mailers

```
all_lm_social_coefs <- ols_step_all_possible_betas(lm_out)
```

```
all_lm_social_coefs
```

	model	predictor	beta
1	1	(Intercept)	0.2966383083
2	1	messagesCivic Duty	0.0178993441
3	1	messagesHawthorne	0.0257363121
4	1	messagesNeighbors	0.0813099129
5	2	(Intercept)	0.3059095493
6	2	sexmale	0.0126509479
7	3	(Intercept)	0.1055564253
8	3	age	0.0041515670
9	4	(Intercept)	0.2508820413
10	4	primary2004	0.1528795252
11	5	(Intercept)	0.3763534949
12	5	hhsizes	-0.0293482475
13	6	(Intercept)	0.2902800648

Example 2: Social Pressure Mailers

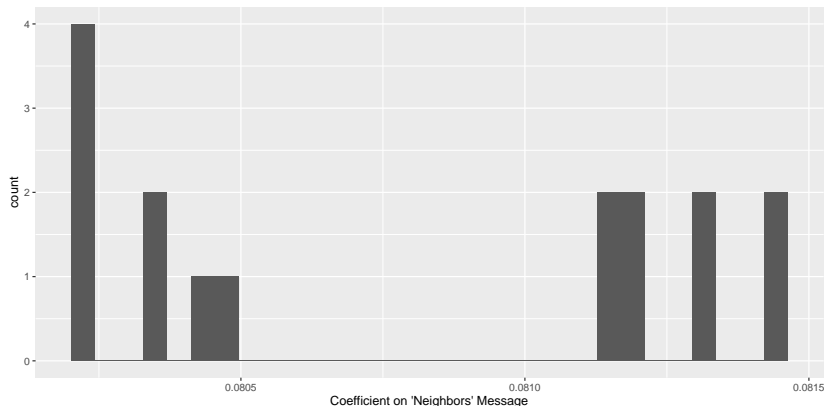


Figure 5: 'Neighbors' Coefs from All Possible Regressions

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
0.08023	0.08032	0.08081	0.08080	0.08122	0.08145

All Coefficients

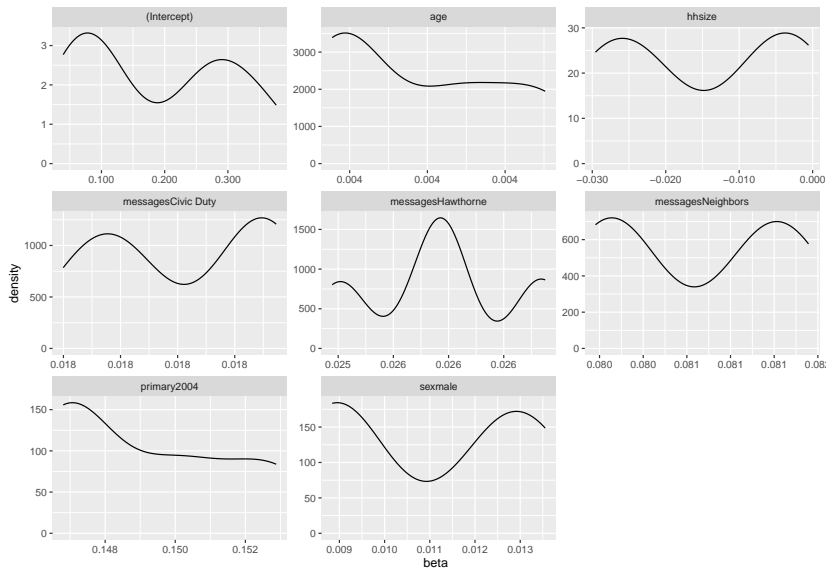


Figure 6: Coefs from All Possible Regressions

Preprocessing to Control Sensitivity

- ▶ So far, “show all the models”

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- ▶ Better: preprocess data to minimize effects of model-based adjustment

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“model-based adjustments ...will give basically the same point estimates”

Matching

X	T	$Y(0)$	$Y(1)$	Y^{obs}
1	1	1	2	2
1	0	1	2	1
1	0	1	2	1
2	1	2	3	3
2	1	2	3	3
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► $\tau_i = 1 \quad \forall i$

► $ATE = \overline{Y(1)} - \overline{Y(0)} = 1$

► $\widehat{ATE} = (\overline{Y(1)}|T=1) - (\overline{Y(0)}|T=0) = \frac{8}{3} - \frac{4}{3} = \frac{4}{3}$

Matching

Suppose we 1:1 exact match on X :

X	T	$Y(0)$	$Y(1)$	Y^{obs}
1	1	1	2	2
1	0	1	2	1
1	0	1	2	1
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$$\widehat{ATE}_m = (\overline{Y_m(1)}|T=1) - (\overline{Y_m(0)}|T=0) = \frac{5}{2} - \frac{3}{2} = 1$$

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$$\widehat{ATE}_m = (\overline{Y_m(1)}|T=1) - (\overline{Y_m(0)}|T=0) = \frac{5}{2} - \frac{3}{2} = 1$$

Not just coincidence; matching removes $X \rightarrow T$.

Ho et al. (2007)

“Matching as Nonparametric Preprocessing for Reducing Model Dependence in Parametric Causal Inference”

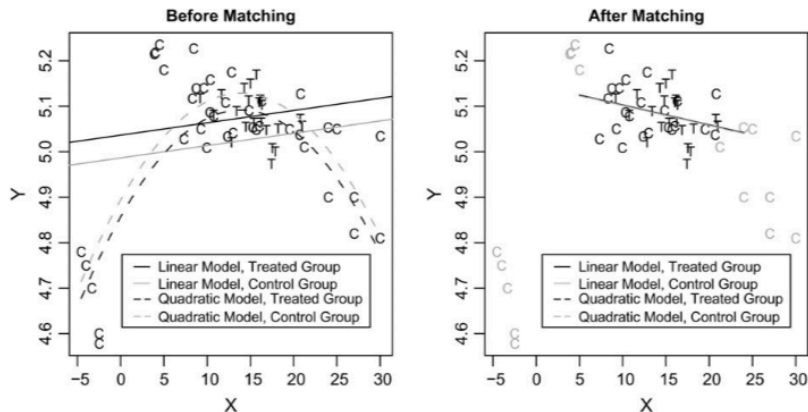


Figure 7: Before: Direction of Effect depends on Model. After: Effect independent of Model.

Reducing Sensitivity in FDA Example

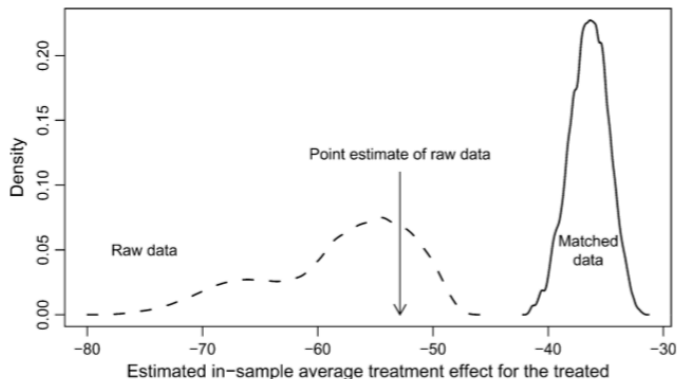


Fig. 2 Kernel density plot (a smoothed histogram) of point estimates of the in-sample ATT of the Democratic Senate majority on FDA drug approval time across 262,143 specifications. The solid line

How to Identify Sensitivity?

Different distributions; non-overlap

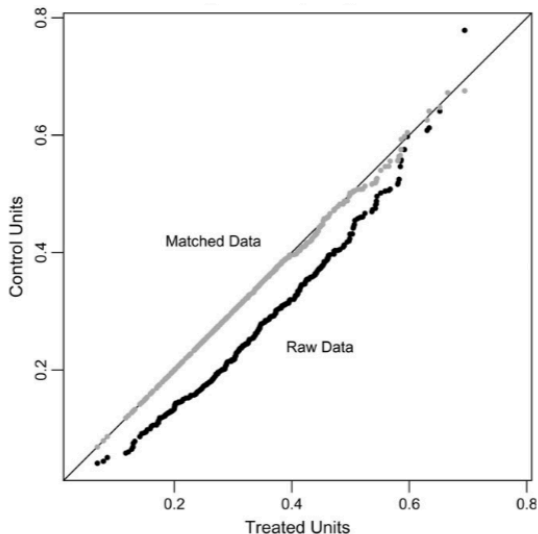


Fig. 3 QQ plot of propensity score for candidate visibility. The black dots represent empirical QQ

Reducing Sensitivity in Candidate Visibility Example

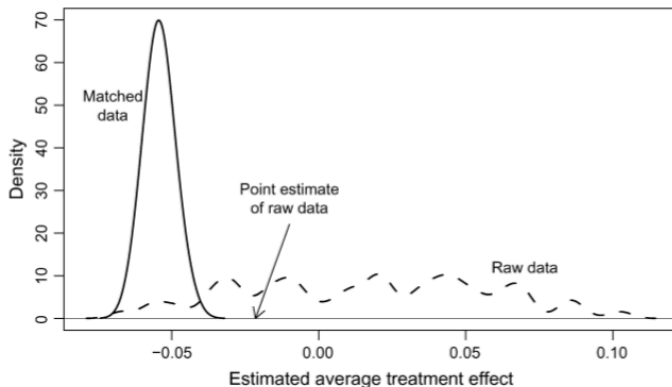


Fig. 4 Kernel density plot of point estimates of the effect of being a less visible male Republican candidate across 63 possible specifications with the Koch data. The dashed line presents estimates for

Paradox of Regression for causal inference?

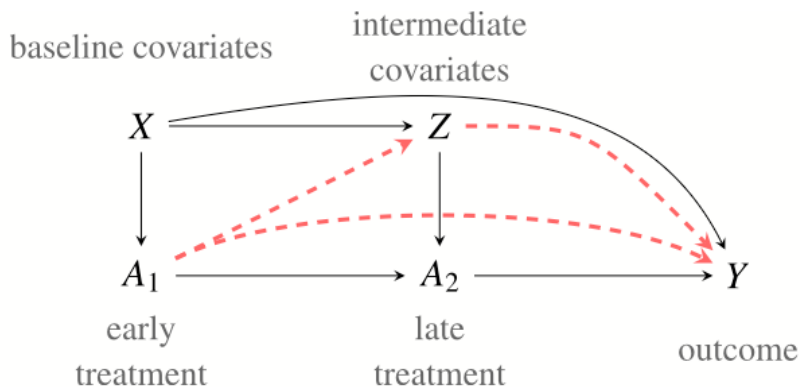
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 \leadsto regression not enough, very sensitive

Paradox of Regression for causal inference?

- ▶ If large diffs in distn's,
 \leadsto regression not enough, very sensitive
- ▶ If small diffs in distn's,
 \leadsto regression won't matter much

Dynamic Treatment Regimes

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Blackwell and Strezhnev (2022)

Telescope Matching

Preprocessing for Dynamic Treatment Regimes:

- ▶ Match across early $\text{Tr}(A_1)$ on baseline covariates X

Telescope Matching

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Telescope Matching

Preprocessing for Dynamic Treatment Regimes:

- ▶ Match across early Tr (A_1) on baseline covariates X
- ▶ Match across late Tr (A_2) on early Tr (exact), baseline + intermediate covariates (A_1, X, Z [or X_1, X_2])
- ▶ Use matches to impute “paths not taken”

Telescope Matching

Preprocessing for Dynamic Treatment Regimes:

Diff-in-means estimator for effect of “early treatment”:

$$\hat{\tau} \equiv \frac{1}{N} \sum_{i=1}^N \left(\hat{Y}_i(1, 0) - \hat{Y}_i(0, 0) \right)$$

Telescope Matching Example

```
library(DirectEffects)  
data(jobcorps)
```

- ▶ Y: self-reported good health (0/1)
- ▶ X1: school/training/job before Job Corps
- ▶ A1: Job Corps program
- ▶ X2: employment in Q4 after assg
- ▶ A2: employment in Q just before outcome

```
# Formula: Y ~ X1 | A1 | X2 | A2
```

```
tm_form <- exhealth30 ~ schobef + trainyrbef + jobeverbef  
  treat | emplq4 + emplq4full | work2year2q
```

```
tm_out <- telescope_match(tm_form, data = jobcorps, verbose = TRUE)
```


Telescope Matching Example

```
tm_out
```

Telescope matching output

Call:

```
telescope_match(formula = tm_form, data = jobcorps, verbose = FALSE)
```

Active treatment: treat

Controlled treatment(s): work2year2q

Estimated controlled direct effects of treat:

	work2year2q	estimate
1	0	-0.006569616
2	1	0.029847086

Telescope Matching Example

```
summary(tm_out)
```

Telescope matching results

Call:

```
telescope_match(formula = tm_form, data = jobcorps, verbose = FALSE)
```

Active treatment: treat

Controlled treatment(s): work2year2q

Matching summary:

	Term	Matching Ratio	L:1	N == 1	N == 0	Matched == 1	Matched == 0
1	treat		5	6034	3991	5800	3989
2	work2year2q		5	6207	3818	3655	3659

Summary of units matching contributions:

	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
treat	0	0.6	0.8	1	1.40	3.80
treat:work2year2q	0	0.0	0.4	1	1.04	90.08
work2year2q	0	0.0	0.4	1	1.20	68.00

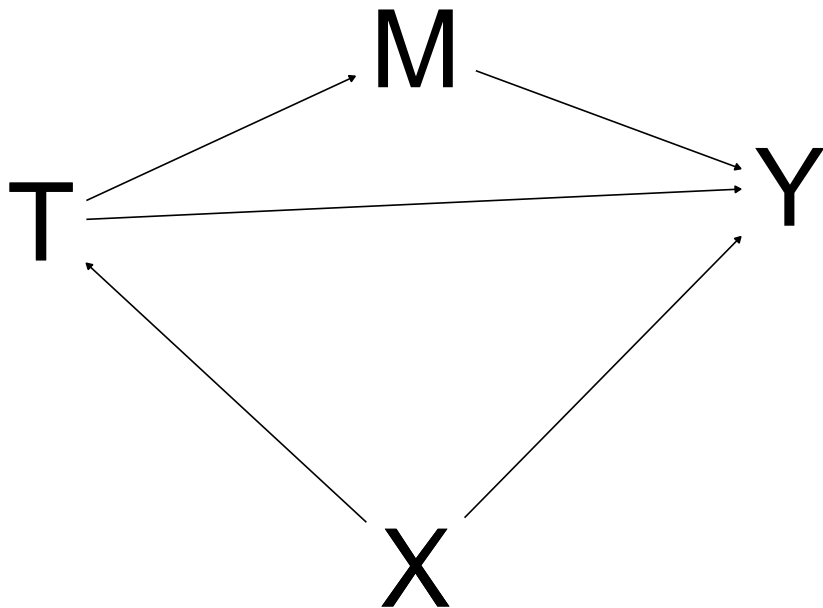
Estimated controlled direct effects of treat:

	work2year2q	Estimate	Estimate (no BC)	Std. Err.
(1, 0) vs. (0, 0)	0	-0.00657	-0.005897	0.03767
(1, 1) vs. (0, 1)	1	0.02985	0.029813	0.01421

Sensitivity to an Unidentifiable Parameter

Mediation Analysis

Confounding in Observational Studies



Mediation Effects

- ▶ If interest is $M \rightarrow Y$, seek experiment-like M

Mediation Effects

- ▶ If interest is $M \rightarrow Y$, seek experiment-like M
 - ▶ random M

Mediation Effects

- ▶ If interest is $M \rightarrow Y$, seek experiment-like M
 - ▶ random M
 - ▶ subclassify/match for M

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 - ▶ RDD, synthetic control for M

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 - ▶ random T
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- ▶ In mediation, interest is $T \rightarrow M \rightarrow Y$

Mediation Effects

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- ▶ If interest is $T \rightarrow Y$, seek experimental T
 - ▶ random T
 - ▶ subclassify/match for T
 - ▶ instrumented T
 - ▶ RDD, synthetic control for T
- ▶ In mediation, interest is $T \rightarrow M \rightarrow Y$
 - ▶ (and maybe $T \rightarrow (\neg M) \rightarrow Y$)

Mediation Effects

Condition on /control for M ?

Mediation Effects

Condition on /control for M ?

► No: how to estimate $M \rightarrow Y$?

Mediation Effects

Condition on /control for M ?

- ▶ No: how to estimate $M \rightarrow Y$?
- ▶ Yes: induces post-treatment bias in estimate of $T \rightarrow Y$

Mediation Effects

Condition on /control for M ?

- ▶ No: how to estimate $M \rightarrow Y$?
- ▶ Yes: induces post-treatment bias in estimate of $T \rightarrow Y$

Mediation Effects

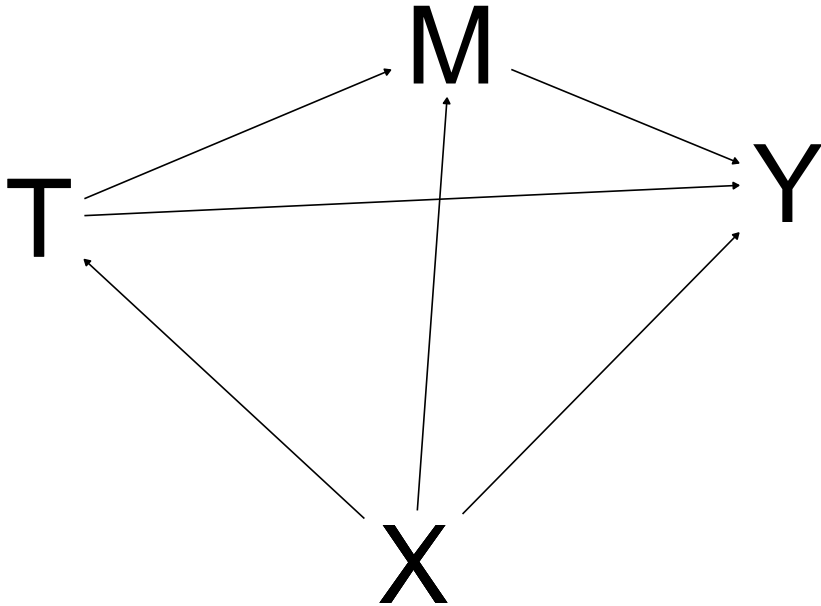
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- ▶ Even worse ...



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To break confounding,

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To break confounding,

- ▶ can't break $X \rightarrow Y$
- ▶ break $X \rightarrow T$
- ▶ but $X \rightarrow M$ may still remain!

Post-Treatment Bias

- ▶ Interest in effect of news on attitude.

Post-Treatment Bias

- Interest in effect of news on attitude. Randomly assign news:

```
n <- 200  
news <- sample(0:1, n, replace = TRUE)
```

Post-Treatment Bias

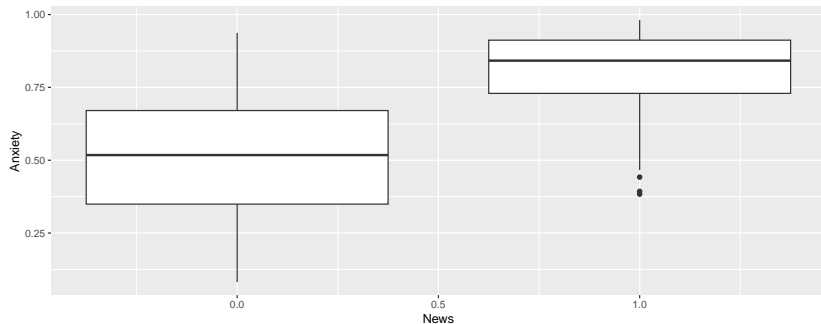
- ▶ News status greatly affects Anxiety:

```
pr.anx <- 1/(1 + exp(-(news * 2 + rnorm(n))))
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Post-Treatment Bias

- ▶ News status greatly affects Anxiety:

```
summary(lm(pr.anx ~ news))$coef |> round(3)
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	0.519	0.018	29.196	0
news	0.294	0.025	11.692	0

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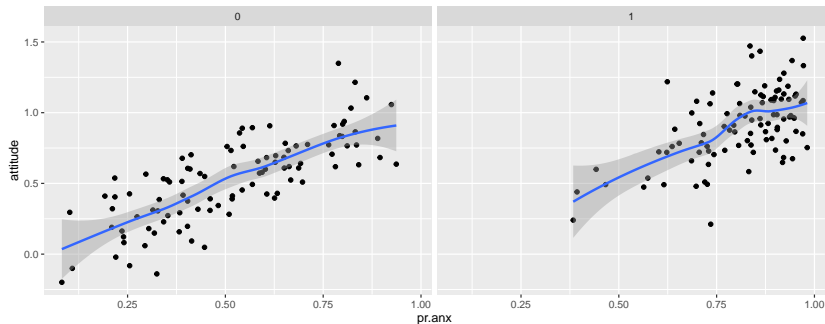
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summary(lm(attitude ~ news + pr.anx))$coef |> round(4)
```

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-0.0438	0.0461	-0.9511	0.3427
news	0.0655	0.0368	1.7807	0.0765
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Mediation

Mediation analysis tries to estimate how much effect of T on Y goes through M .

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- ▶ Quiz: In news/anxiety/attitude example,
 - ▶ what's $Y_i(1, M_i(1))$?
 - ▶ what's $Y_i(0, M_i(0))$?
 - ▶ what's $Y_i(1, M_i(1)) - Y_i(0, M_i(0))$?
 - ▶ what's $Y_i(1, M_i(0))$?

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For individuals:

► $Y_i(1, M_i(1)) - Y_i(0, M_i(0))$: Total effect

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Moderation and Interaction

- ▶ Moderators and mediators are both “third variables”

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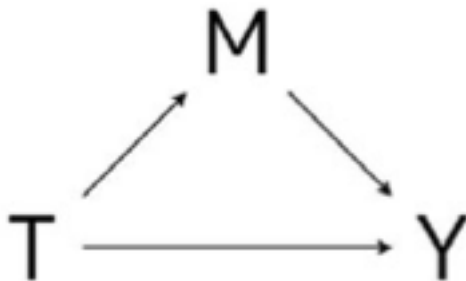
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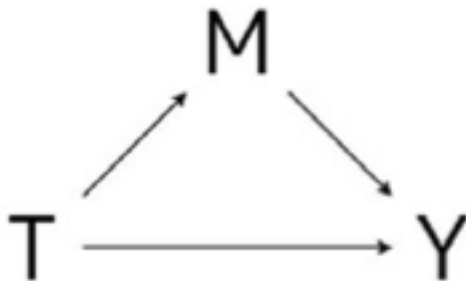
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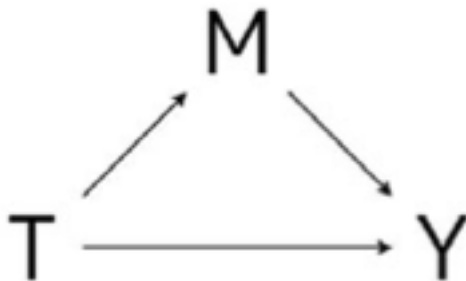
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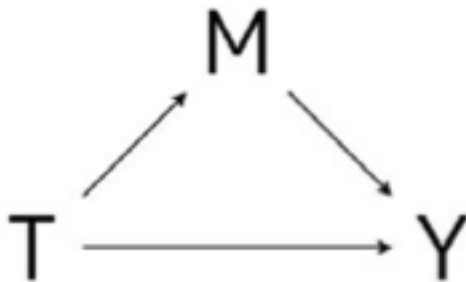
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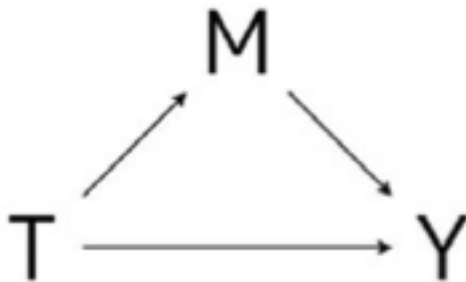
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 - ▶ When there is an “interaction between T and X ”

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 - ▶ Informative bounds for ACME!

Usual (Best-Case?) Way to “Combine Somehow”

“Baron & Kenny Procedure”

$$M_i = \alpha_1 + aT_i + \epsilon_{i1} \quad (1)$$

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Then, call effect of

$$T \rightarrow M = a$$

$$T \rightarrow Y = c \quad (\text{Total})$$

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TABLE 1. The Fallacy of the Causal Chain Approach

Population Proportion	Potential Mediators and Outcomes				Treatment Effect on Mediator $M_i(1) - M_i(0)$	Mediator Effect on Outcome $Y_i(t, 1) - Y_i(t, 0)$	Causal Mediation Effect $Y_i(t, M_i(1)) - Y_i(t, M_i(0))$
	$M_i(1)$	$M_i(0)$	$Y_i(t, 1)$	$Y_i(t, 0)$			
0.3	1	0	0	1	1	-1	-1
0.3	0	0	1	0	0	1	0
0.1	0	1	0	1	-1	-1	1
0.3	1	1	1	0	0	1	0
Average	0.6	0.4	0.6	0.4	0.2	0.2	-0.2

Notes: The left five columns of the table show a hypothetical population proportion of “types” of units defined by the values of potential mediators and outcomes. Note that these values can never be jointly observed. The last row of the table shows the population average value of each column. In this example, the average causal effect of the treatment on the mediator (the sixth column) is positive and equal to 0.2. Moreover, the average causal effect of the mediator on the outcome (the seventh column) is also positive and equals 0.2. And yet the average causal mediation effect (ACME; final column) is negative and equals -0.2.

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0.3	1	0	0	1	1	-1	-1
0.3	0	0	1	0	0	1	0
0.1	0	1	0	1	-1	-1	1
0.3	1	1	1	0	0	1	0
Average	0.6	0.4	0.6	0.4	0.2	0.2	-0.2

Notes: The left five columns of the table show a hypothetical population proportion of “types” of units defined by the values of potential mediators and outcomes. Note that these values can never be jointly observed. The last row of the table shows the population average value of each column. In this example, the average causal effect of the treatment on the mediator (the sixth column) is positive and equal to 0.2. Moreover, the average causal effect of the mediator on the outcome (the seventh column) is also positive and equals 0.2. And yet the average causal mediation effect (ACME; final column) is negative and equals -0.2.

$$T \rightarrow M = a = 0.2$$

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Why Aren't 2 Experiments Enough?

TABLE 1. The Fallacy of the Causal Chain Approach

Population Proportion	Potential Mediators and Outcomes				Treatment Effect on Mediator $M_i(1) - M_i(0)$	Mediator Effect on Outcome $Y_i(t, 1) - Y_i(t, 0)$	Causal Mediation Effect $Y_i(t, M_i(1)) - Y_i(t, M_i(0))$
	$M_i(1)$	$M_i(0)$	$Y_i(t, 1)$	$Y_i(t, 0)$			
0.3	1	0	0	1	1	-1	-1
0.3	0	0	1	0	0	1	0
0.1	0	1	0	1	-1	-1	1
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But, true $\bar{\delta}(t)$, ACME, = -0.2!

What Else Do You Need?

Consistency assumption: $T_i = t$, $M_i = m$ have same effect regardless of how they came to have those values.

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The ACME, e.g., is an estimate of the effect of changes in M due to changing T (but without changing T).

(Other manipulations of M rely on consistency.)

What Else Do You Need?

Big picture: to get more detailed estimates from same data,
need more assumptions

Assumption 1 [Sequential Ignorability (Imai, Keele, and Yamamoto 2010)].

$$\{Y_i(t', m), M_i(t)\} \perp\!\!\!\perp T_i \mid X_i = x, \quad (3)$$

$$Y_i(t', m) \perp\!\!\!\perp M_i(t) \mid T_i = t, X_i = x, \quad (4)$$

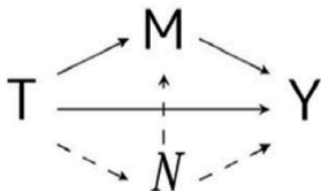
where $0 < \Pr(T_i = t \mid X_i = x)$ and $0 < p(M_i = m \mid T_i = t, X_i = x)$ for $t = 0, 1$, and all x and m in the support of X_i and M_i , respectively.

What Else Do You Need?

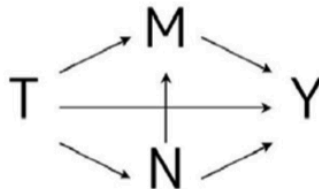
- ▶ Eqn 3: Conditional independence of PotOut's from Tr, given X (pretreatment!)
 - ▶ Ok, for random T , or balanced obs design. T as good as random, exog., etc.
 - ▶ (t' is just saying, for each $t = 0, 1$, must have Y 's from both $t = 0, 1$ must be indep.)
- ▶ Eqn 4: Hard. Mediator is as good as random, given particular Tr status
- ▶ Problem: can't randomize both T and M in same experiment
 - ▶ (if want effect of T through M)
- ▶ You're getting 2 different QoI's if you randomize both: $T \rightarrow M, Y$ and $M \rightarrow Y$.
 - ▶ Showed can't combine those into $T \rightarrow M \rightarrow Y$

When Can You Get It?

FIGURE 8. Second Mediator Causing Serious Problem



(a)



(b)

Estimating Mediation Effects

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- ▶ Practical advice: start there. Then, formal mediation.

Sensitivity for Mediation Effects

Given

$$M_i = \alpha_1 + aT_i + \epsilon_{i1} \quad (4)$$

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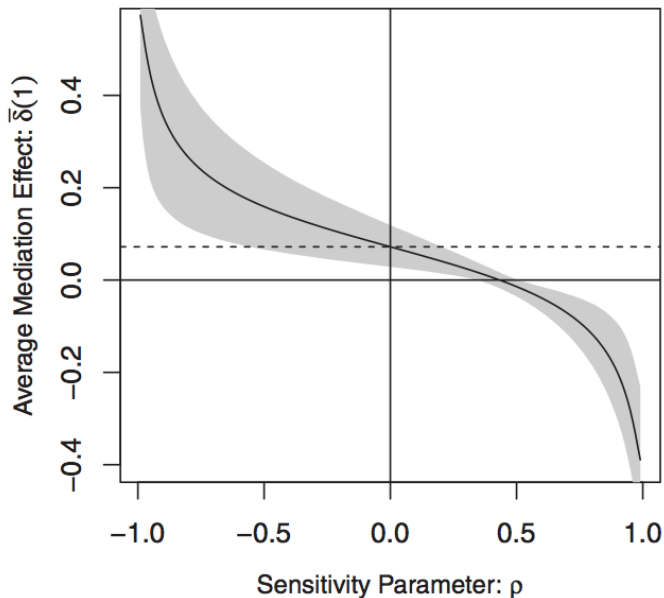
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(If $\neg Q$, then $\neg P$.)
(I.e., From freq. standpoint, you can find “evidence of problem”, or “no evidence of problem”, but not “evidence of no problem”.)

Sensitivity for Mediation Effects



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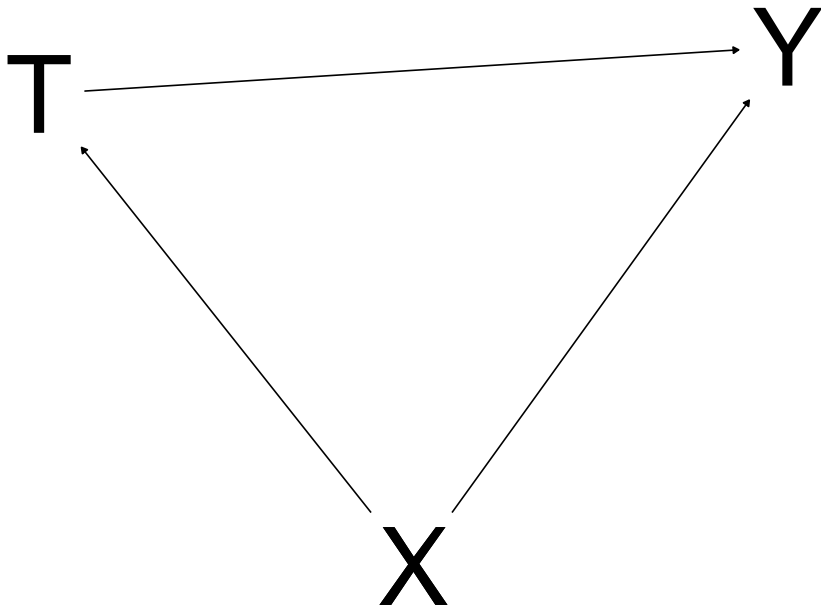
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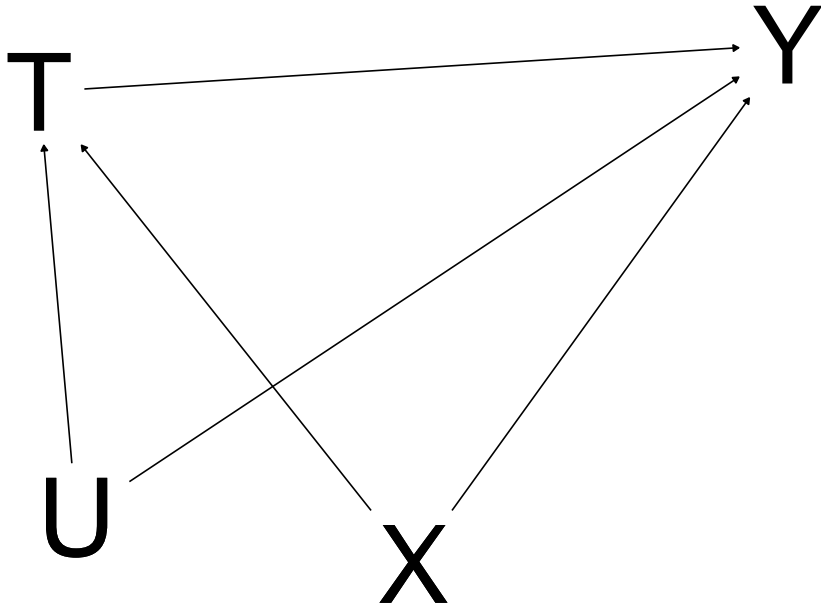
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- ▶ From Bullock, Green, and Ha (2010):

a cumulative enterprise. Persuasive conclusions about mediation are difficult to reach under any circumstances, but they are most likely to be reached when they derive from an experimental research program that addresses the particular challenges of mediation analysis—challenges that we describe here.

Sensitivity to an Unobserved Covariates

Confounding in Observational Studies





Addressing Confounding

To break confounding,

- ▶ can't break $X \rightarrow Y$
- ▶ break $X \rightarrow T$
- ▶ I.e., make $X \perp\!\!\!\perp T$
- ▶ But this doesn't address $U \rightarrow T$ (or $U \rightarrow Y$).

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(Of course, if no causal effect of $U \rightarrow Y$, no problem.)

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Sensitivity: how strong must sociability be to invalidate inference about phone calls?

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$$A_1 : A_2 = \frac{p(A_1)}{p(A_2)}$$

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- ▶ (Diffs betwn groups real, importnt, but how we meas. changes is tricky)

King's Conjecture



Gary King @kinggary

the "odds ratio" is a lame way to communicate statistical results;
I conjecture that there's *always* a better way

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17 October 2012

Modeling Hidden Bias

Odds of treatment for i and j :

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OR of i versus j :

$$\begin{aligned} OR &= \frac{\pi_i}{1 - \pi_i} \div \frac{\pi_j}{1 - \pi_j} \\ &= \frac{\pi_i(1 - \pi_j)}{\pi_j(1 - \pi_i)} \end{aligned}$$

Modeling Hidden Bias

Let Γ be upper bound on OR of treatment.

$$\frac{1}{\Gamma} \leq \frac{\pi_i(1 - \pi_j)}{\pi_j(1 - \pi_i)} \leq \Gamma \quad \forall i, j \text{ s.t. } \mathbf{x}_i = \mathbf{x}_j$$

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By what factor does the odds of treatment differ? (No more than Γ)

Modeling Hidden Bias

Rosenbaum (2020) shows that this is same as

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s.t. $0 \leq u_i \leq 1$.

Interpretation: first rewrite

$$\log\left(\frac{\pi_j}{1-\pi_j}\right) = \kappa(\mathbf{x}_i) + \gamma u_j$$

Exponentiate:

$$\left(\frac{\pi_i}{1-\pi_i}\right) = e^{\kappa(\mathbf{x}_i)+\gamma u_i}$$

$$\left(\frac{\pi_j}{1-\pi_j}\right) = e^{\kappa(\mathbf{x}_j)+\gamma u_j}$$

Exponentiate:

$$\begin{aligned}\left(\frac{\pi_i}{1-\pi_i}\right) &= e^{\kappa(\mathbf{x}_i)+\gamma u_i} \\ \left(\frac{\pi_j}{1-\pi_j}\right) &= e^{\kappa(\mathbf{x}_i)+\gamma u_j}\end{aligned}$$

Calculate OR:

$$\begin{aligned}OR &= \frac{\pi_i(1-\pi_j)}{\pi_j(1-\pi_i)} \\ &= \frac{e^{\kappa(\mathbf{x}_i)+\gamma u_i}}{e^{\kappa(\mathbf{x}_i)+\gamma u_j}} \\ &= e^{(\kappa(\mathbf{x}_i)+\gamma u_i)-(\kappa(\mathbf{x}_i)+\gamma u_j)} \\ &= e^{(\gamma u_i-\gamma u_j)} \\ &= e^{\gamma(u_i-u_j)}\end{aligned}$$

Interpreting Γ

$$OR = e^{\gamma(u_i - u_j)}$$

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Log odds differ by factor of γ times diff in unobs confounder.

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Log odds differ by factor of γ times diff in unobs confounder.

Shows $\Gamma = e^\gamma$.

TABLE 4.1. Sensitivity Analysis for Hammond's Study of Smoking and Lung Cancer: Range of Significance Levels for Hidden Biases of Various Magnitudes.

Γ	Minimum	Maximum
1	< 0.0001	< 0.0001
2	< 0.0001	< 0.0001
3	< 0.0001	< 0.0001
4	< 0.0001	0.0036
5	< 0.0001	0.03
6	< 0.0001	0.1

TABLE 4.1. Sensitivity Analysis for Hammond's Study of Smoking and Lung Cancer: Range of Significance Levels for Hidden Biases of Various Magnitudes.

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3	< 0.0001	< 0.0001
4	< 0.0001	0.0036
5	< 0.0001	0.03
6	< 0.0001	0.1

- ▶ Groups: smokers/nonsmokers
- ▶ Outcome: lung cancer
- ▶ Something must increase smoking by $6\times$ to change inference.
- ▶ If exists, maybe it's that factor, not smoking directly.

(Bias from $U \rightarrow T$; effectively, $U \rightarrow Y$ nearly perfect.)

Γ	Minimum	Maximum
1	≤ 0.0001	≤ 0.0001
2	≤ 0.0001	0.0018
3	≤ 0.0001	0.0136
4	≤ 0.0001	0.0388
4.25	≤ 0.0001	0.0468
5	≤ 0.0001	0.0740

Table 4.2: Signed-Rank Statistic p -value Sensitivity for Lead in Children's Blood

- ▶ Groups: parents occupationally exposed/unexposed
- ▶ Outcome: children's levels
- ▶ Something must increase parents' exposure by $5\times$ to change inference.
- ▶ If exists, maybe it's that, not parental exposure directly.

Γ	Minimum	Maximum
1	≤ 0.0001	≤ 0.0001
2	≤ 0.0001	0.0018
3	≤ 0.0001	0.0136
4	≤ 0.0001	0.0388
4.25	≤ 0.0001	0.0468
5	≤ 0.0001	0.0740

Table 4.2: Signed-Rank Statistic p -value Sensitivity for Lead in Children's Blood

- ▶ Groups: parents occupationally exposed/unexposed
- ▶ Outcome: children's levels
- ▶ Something must increase parents' exposure by $5\times$ to change inference.
- ▶ If exists, maybe it's that, not parental exposure directly.

(one-sided)

Γ	Minimum	Maximum
1	15	15
2	10.25	19.5
3	8	23
4	6.5	25
5	5	26.5

Table 4.3: Point Estimate Sensitivity for Lead in Children's Blood

Γ	Minimum	Maximum
1	15	15
2	10.25	19.5
3	8	23
4	6.5	25
5	5	26.5

Table 4.3: Point Estimate Sensitivity for Lead in Children's Blood

- ▶ HL point estimate: 15 (median of all $m \times n$ possible matched pairs)
- ▶ With confounding, wider range of possible effects.

Γ	95% CI
1	(9.5, 20.5)
2	(4.5, 27.5)
3	(1.0, 32.0)
4	(-1.0, 36.5)
5	(-3.0, 41.5)

Table 4.4: Confidence Interval Sensitivity for Lead in Children's Blood

Γ	95% CI
1	(9.5, 20.5)
2	(4.5, 27.5)
3	(1.0, 32.0)
4	(-1.0, 36.5)
5	(-3.0, 41.5)

Table 4.4: Confidence Interval Sensitivity for Lead in Children's Blood

- ▶ Inverted NHST CI's
- ▶ If something increases parental exposure by $4\times$, negative estimates of parents on children are reasonable.

(two-sided)

Implementation

Packages

- ▶ Frank et al. (2013): `konfound`
- ▶ Keele (2022): `rbounds`
- ▶ `sensitivitymw`
- ▶ `sensitivitymv`

Example

```
anes <- read_csv("../data/anes_pilot_2016.csv")  
dim(anes)
```

```
[1] 1200  594
```

```
anes <- anes |> mutate(age = 2016 - birthyr,  
                      pid_rep = as.numeric(pid3 == 3),  
                      pid_dem = as.numeric(pid3 == 1))
```



```
lm_out <- lm(turnout12 ~ pid_rep, data = anes)
summary(lm_out)
```

Call:

```
lm(formula = turnout12 ~ pid_rep, data = anes)
```

Residuals:

	Min	1Q	Median	3Q	Max
	-0.3395	-0.2451	-0.2451	-0.2451	1.7549

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.24512	0.01868	66.641	< 2e-16 ***
pid_rep	0.09435	0.03320	2.842	0.00456 **

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.535 on 1198 degrees of freedom
Multiple R-squared: 0.006605 Adjusted R-squared: 0.005

```
library(konfound)
konfound(lm_out, pid_rep)
```

```
library(konfound)
konfound(lm_out, pid_rep)
```

Robustness of Inference to Replacement (RIR):

To invalidate an inference, 30.959 % of the estimate would have to be due to bias.

This is based on a threshold of 0.065 for statistical significance ($\alpha = 0.05$).

To invalidate an inference, 372 observations would have to be replaced with cases for which the effect is 0 (RIR = 372).

See Frank et al. (2013) for a description of the method.

Citation: Frank, K.A., Maroulis, S., Duong, M., and Kelcey, B. (2013).

What would it take to change an inference?

Using Rubin's causal model to interpret the robustness of causal inferences

```
lm_out <- lm(turnout12 ~ pid_rep + age, data = anes)
summary(lm_out)
```

Call:

```
lm(formula = turnout12 ~ pid_rep + age, data = anes)
```

Residuals:

Min	1Q	Median	3Q	Max
-0.5825	-0.3388	-0.1711	0.0301	1.9831

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.678649	0.045960	36.524	< 2e-16 ***
pid_rep	0.082685	0.031870	2.594	0.00959 **
age	-0.008943	0.000873	-10.244	< 2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.5122 on 1197 degrees of freedom

```
konfound(lm_out, pid_rep)
```

Robustness of Inference to Replacement (RIR):

To invalidate an inference, 24.379 % of the estimate would have to be due to bias.

This is based on a threshold of 0.063 for statistical significance ($\alpha = 0.05$).

To invalidate an inference, 293 observations would have to be replaced with cases for which the effect is 0 (RIR = 293).

See Frank et al. (2013) for a description of the method.

Citation: Frank, K.A., Maroulis, S., Duong, M., and Kelcey, B. (2013).

What would it take to change an inference?

Using Rubin's causal model to interpret the robustness of causal inferences.

Education, Evaluation and

```
cor(anes[,c("pid_rep", "turnout12", "econnow")])
```

	pid_rep	turnout12	econnow
pid_rep	1.00000000	0.081825966	0.141257803
turnout12	0.08182597	1.000000000	0.008599061
econnow	0.14125780	0.008599061	1.000000000

```
lm_out <- lm(turnout12 ~ pid_rep + age + econnow, data = ar  
summary(lm_out)
```

Call:

```
lm(formula = turnout12 ~ pid_rep + age + econnow, data = ar
```

Residuals:

	Min	1Q	Median	3Q	Max
	-0.60257	-0.33748	-0.17138	0.04458	1.96702

Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	1.6290966	0.0565381	28.814	<2e-16 ***
pid_rep	0.0755031	0.0322095	2.344	0.0192 *
age	-0.0091496	0.0008833	-10.358	<2e-16 ***
econnow	0.0202398	0.0134633	1.503	0.1330

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```
konfound(lm_out, pid_rep)
```

Robustness of Inference to Replacement (RIR):

To invalidate an inference, 16.303 % of the estimate would have to be due to bias.

This is based on a threshold of 0.063 for statistical significance ($\alpha = 0.05$).

To invalidate an inference, 196 observations would have to be replaced with cases for which the effect is 0 (RIR = 196).

See Frank et al. (2013) for a description of the method.

Citation: Frank, K.A., Maroulis, S., Duong, M., and Kelcey, B. (2013).

What would it take to change an inference?

Using Rubin's causal model to interpret the robustness of causal inferences.

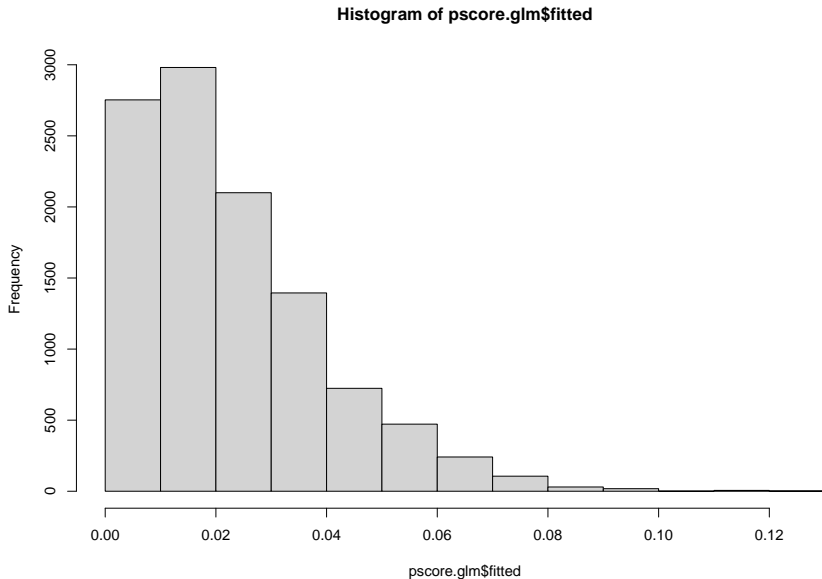
Education, Evaluation and

Implementation in rbounds

```
library(Matching)
data(GerberGreenImai)

# Estimate Propensity Score
pscore.glm <- glm(PHN.C1 ~ PERSONS + VOTE96.1 +
                  NEW + MAJORPTY + AGE + WARD +
                  PERSONS:VOTE96.1 + PERSONS:NEW +
                  AGE2, family = binomial(logit),
                  data = GerberGreenImai)
```

```
hist(pscore.glm$fitted)
```



Implementation in rbounds

```
# Match - without replacement
m.obj <- Match(Y = GerberGreenImai$VOTED98,
              Tr = GerberGreenImai$PHN.C1,
              X = fitted(pscore.glm), M = 1, replace = FALSE)

summary(m.obj)
```

```
Estimate... 0.036437
SE..... 0.041421
T-stat..... 0.87968
p.val..... 0.37903
```

```
Original number of observations..... 10829
Original number of treated obs..... 247
Matched number of observations..... 247
Matched number of observations (unweighted). 247
```

Implementation in rbounds

```
library(rbounds)

# Sensitivity Test
# binarysens(m.obj, Gamma = 2, GammaInc = .1)
```

Implementation in rbounds

```
#hlsens(m.obj, Gamma = 5, GammaInc = 1)
```

Thanks!

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