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# Causal Inference

a summary

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# Contents

<b>1</b>	<b>General</b>	<b>3</b>	2.2 G-Methods . . . . .	5
			2.2.1 Time-varying A . . . . .	6
<b>2</b>	<b>Models</b>	<b>5</b>		
	2.1 Traditional Methods . . . . .	5	<b>3</b>	<b>Longitudinal Data</b>
				<b>8</b>

# 1 General

## Causal Roadmap (Petersen and van der Laan, 2014)

systematic approach linking causality to statistical procedures

**1. Specifying Knowledge.** structural causal model (unifying counterfactual language, structural equations, & causal graphs): a set of possible data-generating processes, expresses background knowledge and its limits

**2. Linking Data.** specifying measured variables and sampling specifics (latter can be incorporated into the model)

**3. Specifying Target.** define hypothetical experiment: decide

1. variables to intervene on: one (point treatment), multiple (longitudinal, censoring/missing, (in)direct effects)
2. intervention scheme: static, dynamic, stochastic
3. counterfactual summary of interest: absolute or relative, marginal structural models, interaction, effect modification
4. population of interest: whole, subset, different population

**4. Assessing Identifiability.** are knowledge and data sufficient to derive estimand and if not, what else is needed?

**5. Select Estimand.** current best answer: knowledge-based assumptions + which minimal convenience-based assumptions (transparency) gets as close as possible

**6. Estimate.** choose estimator by statistical properties, nothing causal here

**7. Interpret.** hierarchy: statistical, counterfactual, feasible intervention, randomized trial

## Average Causal Effect $E[Y^{a=1}] \neq E[Y^{a=0}]$

$$E[Y^a] = \sum_y y p_{Y^a}(y) \quad (\text{discrete})$$

$$= \int y f_{Y^a}(y) dy \quad (\text{continuous})$$

individual causal effect  $Y_i^{a=1} \neq Y_i^{a=0}$  generally unidentifiable

*null hypothesis:* no average causal effect

*sharp null hypothesis:* no causal effect for any individual

**notation**  $A, Y$ : random variables (differ for individuals);  $a, y$ : particular values; counterfactual  $Y^{a=1}$ :  $Y$  under treatment  $a = 1$

**stable unit treatment value assumption (SUTVA)**  $Y_i^a$  is well-defined: no interference between individuals, no multiple versions of treatment (weaker: treatment variation irrelevance)

**causal effect measures** typically based on means

*risk difference:*  $\Pr[Y^{a=1} = 1] - \Pr[Y^{a=0} = 1]$

*risk ratio:*  $\frac{\Pr[Y^{a=1} = 1]}{\Pr[Y^{a=0} = 1]}$

*odds ratio:*  $\frac{\Pr[Y^{a=1} = 1]/\Pr[Y^{a=1} = 0]}{\Pr[Y^{a=0} = 1]/\Pr[Y^{a=0} = 0]}$

*number needed to treat (NNT)* to save 1 life:  $-1/\text{risk difference}$

**sources of random error:** sampling variability (use consistent estimators), nondeterministic counterfactuals

**association** compares  $E[Y|A = 1]$  and  $E[Y|A = 0]$ , **causation** compares  $E[Y^{a=1}]$  and  $E[Y^{a=0}]$  (whole population)

## Target Trial

emulating an ideal randomized experiment explicitly formulate target trial & show how it is emulated → less vague causal question, helps spot issues

**missing data problem** unknown counterfactuals

*randomized experiments:* missing completely at random →

exchangeability (= exogeneity as treatment is exogenous)

*ideal randomized experiment:* no censoring, double-blind,

well-defined treatment, & adherence → association is causation

*pragmatic trial:* no placebo/blindness, realistic monitoring

**PICO** (population, intervention, comparator, outcome): some components of target trial

**three types of causal effects:**

*intention-to-treat effect* (effect of treatment assignment)

*per-protocol effect* (usually dynamic when toxicity arises)

*other intervention effect* (strategy changed during follow-up)

**controlled direct effects:** effect of  $A$  on  $Y$  not through  $B$

*natural direct effect*  $A$  on  $Y$  if  $B^{a=0}$  (cross-world quantity)

*principal stratum effect*  $A$  on  $Y$  for subset with  $B^{a=0} = B^{a=1}$

**crossover experiment:** sequential treatment & outcome  $t=0, 1$  individual causal effect  $Y_{it}^{a_t=1} - Y_{it}^{a_t=0}$  only identifiable if: no carryover effect, effect  $\perp$  time, outcome  $\perp$  time

**time zero** if eligibility at multiple  $t$  (observational data):

earliest, random  $t$ , all  $t$  (adjust variance with bootstrapping)

**grace periods:** usually treatment starts  $x$  months after first eligible, if death before: randomly assign strategy/copy into both

## Identifiability Conditions

hold in ideal experiments

**consistency** counterfactuals correspond to data  $Y = Y^A$ :

if  $A = a$ , then  $Y^a = Y$  for each individual

- precise definition of  $Y^a$  via specifying  $a$  (sufficiently well-defined  $a$  maybe impossible (effect of DNA before it was discovered), relies on expert consensus)
- linkage of counterfactuals to data ( $a$  must be seen in data)

**positivity**  $\Pr[A = a|L = l] > 0 \quad \forall l$  with  $\Pr[L = l] > 0$ ;

$$f_L(l) \neq 0 \Rightarrow f_{A|L}(a|l) > 0 \quad \forall a, l$$

- structural violations (inference not on full population)
- random variability (smooth over with parametric models)

can sometimes be empirically verified (if all is seen in data)

**exchangeability** unverifiable without randomization

- *marginal:*  $Y^a \perp\!\!\!\perp A \triangleq$  randomized experiment, counterfactuals are missing completely at random (MCAR)
- *conditional:*  $Y^a \perp\!\!\!\perp A|L \triangleq$  conditionally randomized, counterfactuals are missing at random (MAR)

alternative definition:  $\Pr[A = 1|Y^{a=0}, L] = \Pr[A = 1|L]$

**additional conditions:**

*correct measurement* mismeasurement of  $A, Y, L$  results in bias

*correct model specification* models  $\xrightarrow{\text{may}}$  misspecification bias

## Effect Modification

$A$  on  $Y$  varies across levels of  $V$

null average causal effect  $\neq$  null causal effect per subgroup

**population characteristics:** causal effect measure is actually

“effect in a population with a particular mix of effect modifiers”

**transportability:** extrapolation of effect to another population

(issues: effect modification, versions of treatment, interference)

effects conditional on  $V$  may be more transportable

**types:** additive/multiplicative scale, qualitative (effect in

opposite directions)/quantitative, surrogate/causal

**calculation:**

- *stratify* by  $V$  then standardize/IP weight for  $L$ ,
- $L$  as *matching* factor (ensures positivity, difficult if high-dimensional  $L$ )

**collapsibility:** causal risk difference and ratio are weighted

averages of stratum-specific risks, can not be done for odds ratio

**interaction** chapter 5

**causal diagrams** chapter 6, include swigs from 7.5 and that  
one technical point  
more on SWIGS p 242ff

**confounding** chapter 7

**selection bias** chapter 8

**measurement bias** chapter 9

**random variabilty** chapter 10

## 2 Models

**Modeling** data are a sample from the target population

*estimand*: quantity of interest, e. g.  $E[Y|A = a]$   
*estimator*: function to use, e. g.  $\hat{E}[Y|A = a]$   
*estimate*: apply function to data, e. g. 4.1

**model**: a priori restriction of joint distribution/dose-response curve; *assumption*: no model misspecification (usually wrong)

**non-parametric estimator**: no restriction (saturated model) = *Fisher consistent estimator* (entire population data  $\rightarrow$  true value)

**parsimonious model**: few parameters estimate many quantities

**bias-variance trade-off**:

wiggleness  $\uparrow \rightarrow$  misspecification bias  $\downarrow$ , CI width  $\uparrow$

**Variable Selection** can induce bias if  $L$  includes:

(descendant of) collider: *selection bias under the null*  
 noncollider effect of  $A$ : *selection bias under the alternative*  
 mediator: *overadjustment for mediators*

temporal ordering is not enough to conclude anything

**bias amplification**: e.g. by adjusting for an instrument  $Z$  (can also reduce bias)

**Machine Learning**  $L$  is high-dimensional

use lasso or ML for IP weighting/standardization

**but**: ML does not guarantee elimination of confounding and has largely unknown statistical properties

$\rightarrow$  **doubly robust estimator**: consistent if bias  $< \frac{1}{\sqrt{n}}$

*sample splitting*: train estimators on training sample, use resulting estimators for doubly robust method on estimation sample (CIs on estimation sample are valid, but  $n$  halved)

*cross-fitting*: do again the other way round, average the two estimates, get CI via bootstrapping

**problems**: unclear choice of algorithm, is bias small enough?

**Super Learning** (Van der Laan et al., 2007, 2011)

**oracle selector**: select best estimator of set of learners  $Z_i$

**discrete super learner**: select algorithm with smallest cross-validated error (converges to oracle for large sample size)

**super learner**: improves asymptotically on discrete version

$\text{logit}(Y = 1|Z) = \sum_i \alpha_i Z_i$ , with  $0 < \alpha_i < 1$  and  $\sum \alpha_i = 1$   
 can be cross-validated itself to check for overfitting (unlikely)

### 2.1 Traditional Methods

**Stratification** calculate risk for each stratum of  $L$   
 only feasible if enough data per stratum

**Outcome regression** chapter 15

**instrumental variable estimation** chapter 16

**causal survival analysis** chapter 17 (and technical point 22.3)

### 2.2 G-Methods

**G-Methods** generalized treatment contrasts: adjust for (surrogate) confounders  $L$

- **standardization** two types of g-formula
- **IP weighting** also g-formula
- **g-estimation**: not needed unless longitudinal

**Standardization** plug-in (or parametric if so) g-formula

$$E[Y^a] = \overbrace{E[E[Y|A=a, L=l]]}^{\text{conditional expectation}} = \overbrace{\int E[Y|L=l, A=a] f_L[l] dl}^{\text{joint density estimator}}$$

weighted average of stratum-specific risks; unknowns can be estimated non-parametrically or modeled

**no need to estimate  $f_L[l]$ /integrate** as empirical distribution can be used: estimate outcome model  $\rightarrow$  predict counterfactuals on whole dataset  $\rightarrow$  average the results ( $\rightarrow$  CI by bootstrapping)

**for discrete  $L$**   $E[Y|A=a] = \sum_l E[Y|L=l, A=a] \Pr[L=l]$

**time-varying** standardize over all possible  $\bar{l}$ -histories  
 simulates joint distribution of counterfactuals  $(Y^{\bar{a}}, \bar{L}^{\bar{a}})$  for  $\bar{a}$   
**joint density estimator (jde)**

$$\text{discrete: } E[Y^{\bar{a}}] = \sum_{\bar{l}} E[Y|\bar{A}=\bar{a}, \bar{L}=\bar{l}] \prod_{k=0}^K f(l_k|\bar{a}_{k-1}, \bar{l}_{k-1})$$

$$\text{continuous: } \int f(y|\bar{a}, \bar{l}) \prod_{k=0}^K f(l_k|\bar{a}_{k-1}, \bar{l}_{k-1}) dl$$

for *stochastic strategies* multiply with  $\prod_{k=0}^K f^{int}(a_k|\bar{a}_{k-1}, \bar{l}_k)$

**estimation** (Young et al., 2011; Schomaker et al., 2019)

1. model  $f(l_k|\bar{a}_{k-1}, \bar{l}_{k-1})$  and  $E[Y|\bar{A}=\bar{a}, \bar{L}=\bar{l}]$
2. simulate data forward in time:  
 at  $k=0$ : use empirical distribution of  $L_0$  (observed data)  
 at  $k>0$ : set  $\bar{A}=\bar{a}$ , draw from models estimated in 1.
3. calculate mean of  $\hat{Y}_{K,i}^{\bar{a}}$  (bootstrap for CI)

**iterated conditional expectation (ice)**

$$E[Y_T^{\bar{a}}] = E[E[E[...E[Y_T|\bar{A}_{T-1}=\bar{a}_{T-1}, \bar{L}_T] ...|\bar{A}_0=a_0, L_1]|L_0]]$$

**estimation** (Schomaker et al., 2019)

1. model inside out:  $Q_T = E[Y_T | \bar{A}_{T-1}, \bar{L}_T]$  to  $Q_0 = E[Q_1 | \bar{L}_0]$ , predict  $Q_t$  with  $\bar{A} = \bar{a}$  in each step
2. calculate mean of  $\hat{Q}_{0,i}^{\bar{a}}$  (bootstrap for CI)

**g-null paradox** even if the sharp null holds, model misspecification can lead to it being falsely rejected

Proof: for  $L_0 \rightarrow A_0 \rightarrow Y_0 \rightarrow L_1 \rightarrow A_1 \rightarrow Y_1$ ,  $\bar{a} = (a_0, a_1)$

$$E[Y_1^{\bar{a}}] \stackrel{CE}{=} E[E[Y_1^{\bar{a}} | A_0=a_0, L_0]]$$

$$(ice) \stackrel{CE^*}{=} E[E[Y_1 | \bar{L}, \bar{A}=\bar{a}, Y_0] | A_0=a_0, L_0]$$

$$\stackrel{LTP}{=} E\left[\sum_{l_1} E[Y_1 | A_0=a_0, \bar{L}, Y_0] \Pr[l_1 | a_0, l_0, y_0]\right]$$

$$\stackrel{LTP}{=} \sum_{l_0} \left[ \sum_{l_1} E[Y_1 | A_0=a_0, \bar{L}, Y_0] \Pr[l_1 | a_0, l_0, y_0] \right] \Pr[l_0]$$

$$(jde) \stackrel{sum}{=} \sum_i E[Y_1 | A_0=a_0, \bar{L}, Y_0] \Pr[l_1 | a_0, l_0] \Pr[l_0]$$

CE: conditional expectation; \*: exchangeability;

LTP: law of total probability

**Marginal Structural Models** association is causation in the IP weighted pseudo-population

associational model  $E[Y|A]$  = causal model  $E[Y^a]$

step 1: estimate/model  $f[A|L]$  (and  $f[A]$ )  $\rightarrow$  get  $(S)W^A$

step 2: estimate regression parameters for pseudo-population

**effect modification** variables  $V$  can be included (e.g.

$\beta_0 + \beta_1 a + \beta_2 V a + \beta_3 V$ ; technically not marginal anymore),

$SW^A(V) = \frac{f[A|V]}{f[A|L]}$  more efficient than  $SW^A$

**Censoring** measuring joint effect of  $A$  and  $C$

$$E[Y^{a,c=0}] \text{ is of interest}$$

**standardization**  $E[Y|A=a] = \int E[Y|L=l, A=a, C=0] dF_L[l]$

**IP weights**  $W^{A,C} = W^A \times W^C$  (uses  $n$ ) or

$$SW^{A,C} = SW^A \times SW^C \text{ (uses } n^{c=0})$$

**g-estimation** can only adjust for confounding, not selection bias

$\rightarrow$  use IP weights

**G-Estimation** (additive) structural nested models

$$\text{logit Pr}[A=1|H(\psi^\dagger), L] = \alpha_0 + \alpha_1 H(\psi^\dagger) + \alpha_2 L$$

$$H(\psi^\dagger) = Y - \psi_1 A$$

find  $\psi^\dagger$  which renders  $\alpha_1 = 0$ ; 95%-CI: all  $\psi^\dagger$  for which  $p > 0.05$

closed-form solution for linear models

**derivation:**  $H(\psi^\dagger) = Y^{a=0}$

$$\text{logit Pr}[A=1|Y^{a=0}, L] = \alpha_0 + \alpha_1 Y^{a=0} + \alpha_2 L$$

$Y^{a=0}$  unknown, but because of exchangeability  $\alpha_1$  should be zero

$$Y^{a=0} = Y^a - \psi_1 a$$

equivalent to  $Y^{a=0} = Y^{a=1} - \psi_1$ , but using no counterfactuals

**structural nested mean model**

$$\text{additive: } E[Y^a - Y^{a=0} | A=a, L] = \beta_1 a (+\beta_2 a L)$$

$$\text{multiplicative: } \log\left(\frac{E[Y^a | A=a, L]}{E[Y^{a=0} | A=a, L]}\right) = \beta_1 a (+\beta_2 a L)$$

multiplicative is preferred if  $Y$  always positive, but does not extend to longitudinal case

semi-parametric: agnostic about  $\beta_0$  and effect of  $L \rightarrow$  robust  $\uparrow$

**no time-varying:** no nesting; model equals marginal structural models with missing  $\beta_0, \beta_3$  (unspecified “no treatment”)

**sensitivity analysis:** unmeasured confounding ( $\alpha_1 \neq 0$ ) can be examined: do procedure for different values of  $\alpha_1 \rightarrow$  plot  $\alpha_1$  vs.  $\psi^\dagger \rightarrow$  how sensitive is estimate to unmeasured confounding?

**effect modification:** add  $V$  in both g-estimation equations

**doubly robust estimators** exist

**IP Weighting** inverse probability of treatment (g-formula)

$$E[Y^a] = E\left[\frac{I(A=a)Y}{f[A|L]}\right]; W^A = \frac{1}{f[A|L]}; SW^A = \frac{f(A)}{f[A|L]}$$

unknowns can be estimated non-parametrically or modeled

**pseudo-population:** everyone is treated & untreated ( $L \not\rightarrow A$ )

**FRCISTG** (fully randomized causally interpreted structured graph): probability tree for  $L \rightarrow A \rightarrow Y$ , can be used to

calculate/visualize simulation of values for  $A$

**for discrete  $A, L$**   $f[a|l] = \Pr[A=a, L=l]$

**estimators:** Horvitz-Thompson; Hajek (modified version)

**stabilized weights  $SW^A$**  should have an average of 1 (check!)

$\rightarrow$  pseudo-population same size  $\rightarrow$  CI width  $\downarrow$

**Standardization and IP Weighting** are equivalent,

**but** if modeled, different “no misspecification” assumptions:

standardization: outcome model

IP weighting: treatment model

**doubly robust estimators:** reduce model misspecification bias, consistent if either model is correct; **e.g.:**

$$R = \begin{cases} +W^A & \text{if } A=1 \\ -W^A & \text{if } A=0 \end{cases}$$

1. fit outcome regression with variable  $R$
2. standardize by averaging

## 2.2.1 Time-varying A

**IP Weighting**

$$W^{\bar{A}} = \prod_{k=0}^K \frac{1}{f(A_k | \bar{A}_{k-1}, \bar{L}_k)}$$

$$SW^{\bar{A}} = \prod_{k=0}^K \frac{f(A_k | \bar{A}_{k-1}, \bar{L}_k)}{\hat{f}(A_k | \bar{A}_{k-1}, \bar{L}_k)}$$

**Doubly Robust Estimator** sequential estimation

1. estimate  $\hat{f}(A_m | \bar{A}_{m-1}, \bar{L}_m)$  (e.g. logistic model), use it to calculate at each time  $m$ :  $\widehat{W}^{\bar{A}_m} = \prod_{k=0}^m \frac{1}{\hat{f}(A_k | \bar{A}_{k-1}, \bar{L}_k)}$  and

$$\text{modified IP weights at } m: \widehat{W}^{\bar{A}_{m-1}, a_m} = \frac{\widehat{W}^{\bar{A}_{m-1}}}{\hat{f}(a_m | \bar{A}_{m-1}, \bar{L}_m)}$$

2. with  $\widehat{T}_{K+1} := Y$ , recursively for  $m = K, K-1, \dots, 0$ :

(a) fit outcome regression on  $\widehat{T}_{m+1}$  with variable  $\widehat{W}^{\bar{A}_m}$

(b) calculate  $\widehat{T}_m$  using the outcome model with  $\widehat{W}^{\bar{A}_{m-1}, a_m}$

3. calculate standardized mean outcome  $\widehat{E}[Y^{\bar{a}}] = E[\widehat{T}_0]$

**valid, if** treatment or outcome model correct, or treatment correct until  $k$  and outcome otherwise ( $k+1$  robustness)

**G-Estimation** nested equations: for each time  $k$

**structural nested mean models** separate effect of each  $a_k$

$$E[Y^{\bar{a}_{k-1}, a_k, \bar{a}_{k+1}} - Y^{\bar{a}_{k-1}, \bar{a}_{k+1}} | \bar{L}^{\bar{a}_{k-1}} = \bar{l}_k, \bar{A}_{k-1} = \bar{a}_{k-1}] = a_k \gamma_k(\bar{a}_{k-1}, \bar{l}_k, \beta)$$

**calculations**

$$H_k(\psi^\dagger) = Y - \sum_{j=k}^K A_j \gamma_j(\bar{A}_{j-1}, \bar{L}_j, \psi^\dagger)$$

function  $\gamma_j$  can be, e.g. constant ( $\psi_1$ ), time-varying only ( $\psi_1 + \psi_2 k$ ), or dependent on treatment/covariate history

$$\text{logit Pr} \left[ A_k = 1 | H_k \left( \psi^\dagger \right), \bar{L}_k, \bar{A}_{k-1} \right] =$$

$$\alpha_0 + \alpha_1 H_k \left( \psi^\dagger \right) + \alpha_2 w_k \left( \bar{L}_k, \bar{A}_{k-1} \right)$$

find  $\alpha_1$  that is closest to zero

closed form estimator exists for the linear case

**Censoring**  $\bar{C}$ : monotonic type of missing data

**standardization:**

$$\int f(y|\bar{a}, \bar{c}=\bar{0}, \bar{l}) \prod_{k=0}^K dF(l_k | \bar{a}_{k-1}, c_{k-1}=0, \bar{l}_{k-1})$$

**IP weighting:**

$$_{SW}^{\bar{C}} = \prod_{k=1}^{K+1} \frac{1 \cdot \text{Pr}(C_k = 0 | \bar{A}_{k-1}, C_{k-1} = 0)}{\text{Pr}(C_k = 0 | \bar{A}_{k-1}, C_{k-1} = 0, \bar{L}_k)}$$

### 3 Longitudinal Data

**Time-Varying Treatments** compare 2 treatments

treatment history up to  $k$ :  $\bar{A}_k = (A_0, A_1, \dots, A_k)$

shorthand: always treated  $\bar{A} = \bar{1}$ , never treated  $\bar{A} = (\bar{0})$

**static strategy:**  $g = [g_0(\bar{a}_{-1}), \dots, g_K(\bar{a}_{K-1})]$

**dynamic strategy:**  $g = [g_0(\bar{l}_0), \dots, g_K(\bar{l}_K)]$

**stochastic strategy:** non-deterministic  $g$

optimal strategy is where  $E[Y^g]$  is maximized (if high is good)

**Sequential Identifiability** sequential versions of

**exchangability:**  $Y^g \perp\!\!\!\perp A_k | \bar{A}_{k-1} \quad \forall g, k = 0, 1, \dots, K$

*conditional exchangeability:*

$$(Y^g, L_{k+1}^g) \perp\!\!\!\perp A_k | \bar{A}_{k-1} = g(\bar{L}_k), \bar{L}^k \quad \forall g, k = 0, 1, \dots, K$$

**positivity:**  $f_{\bar{A}_{k-1}, \bar{L}_k}(\bar{a}_{k-1}, \bar{l}_k) \neq 0 \Rightarrow$

$$f_{A_k | \bar{A}_{k-1}, \bar{L}_k}(a_k | \bar{a}_{k-1}, \bar{l}_k) > 0 \quad \forall (\bar{a}_{k-1}, \bar{l}_k)$$

**consistency:**

$$Y^{\bar{a}} = Y^{\bar{a}^*} \quad \text{if } \bar{a} = \bar{a}^*; \quad Y^{\bar{a}} = Y \quad \text{if } \bar{A} = \bar{a};$$

$$\bar{L}_k^{\bar{a}} = \bar{L}_k^{\bar{a}^*} \quad \text{if } \bar{a}_{k-1} = \bar{a}_{k-1}^*; \quad \bar{L}_k^{\bar{a}} = \bar{L}_k \quad \text{if } \bar{A}_{k-1} = \bar{a}_{k-1}$$

**generalized backdoor criterion** (static strategy): all backdoors into  $A_k$  (except through future treatments) are blocked  $\forall k$

**static sequential exchangeability for  $Y^{\bar{a}}$**

$$Y^{\bar{a}} \perp\!\!\!\perp A_k | \bar{A}_{k-1}, \bar{L}_k \quad \text{for } k = 0, 1, \dots, K$$

use SWIGs to visually check d-separation

**time-varying confounding**  $E[Y^{\bar{a}} | L_0] \neq E[Y | A = \bar{a}, L_0]$

**Treatment-Confounder Feedback**  $A_0 \rightarrow L_1 \rightarrow A_1$ :

an unmeasured  $U$  influencing  $L_1$  and  $Y$  turns  $L_1$  into a collider;

traditional adjustment (e.g. stratification) biased: use g-methods

**g-null test** sequential exchangeability & sharp null true  $\Rightarrow$

$Y^g = Y \quad \forall g \Rightarrow Y \perp\!\!\!\perp A_0 | L_0 \text{ \& } Y \perp\!\!\!\perp A_1 | A_0, L_0, L_1$ ; therefore:

if last two independences don't hold, one assumption is violated

**g-null theorem:**  $E[Y^g] = E[Y]$ , if the two independences hold

( $\Rightarrow$  sharp null: only if strong faithfulness (no effect cancelling))



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*If no citation is given, the information is taken from the book (Hernán and Robins, 2020)*

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