

Confounding adjustment and estimating treatment effects

Without models

Edouard Fu, PhD

Department of Clinical Epidemiology, LUMC

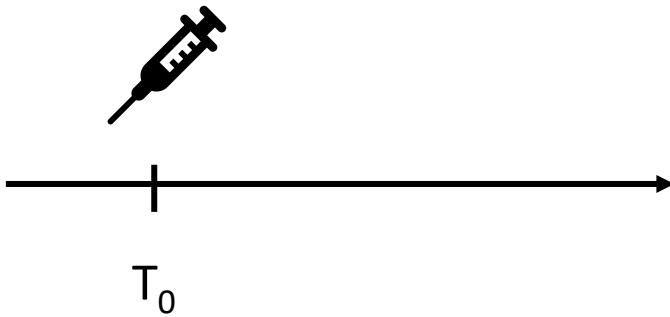


- This is going to be an interactive lecture
- Go to **classpoint.app** and fill in the classcode at the top right corner of this slide

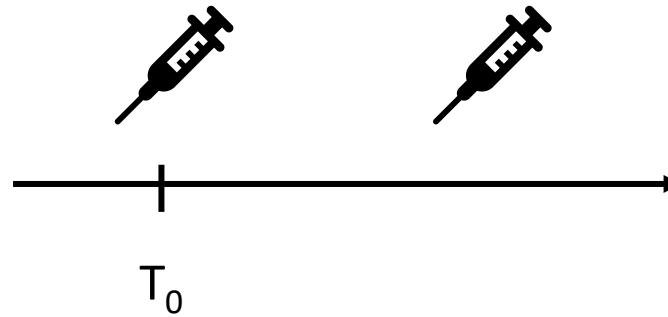
Classification of treatment strategies

Treatment strategies

Point



Sustained



Baseline vs. time-varying confounding

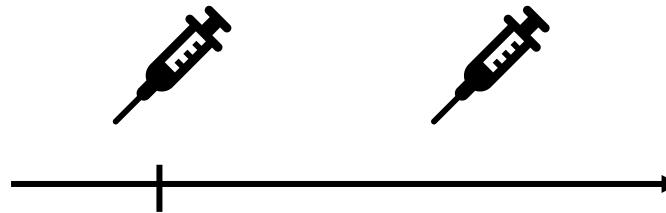
Treatment strategies

Point



- Groups need to be similar at time zero
- Only baseline confounding

Sustained

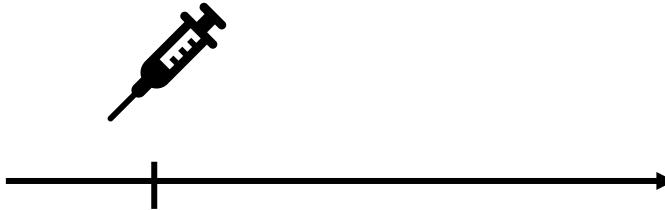


- Groups need to be similar at time zero & during follow-up
- Baseline & time-varying confounding

Baseline vs. time-varying confounding

Treatment strategies

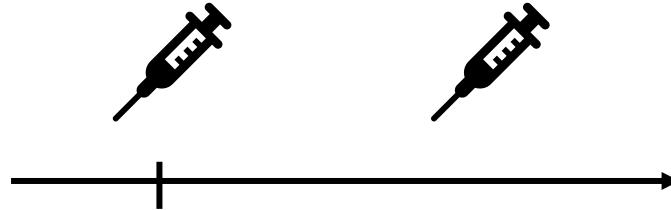
Point



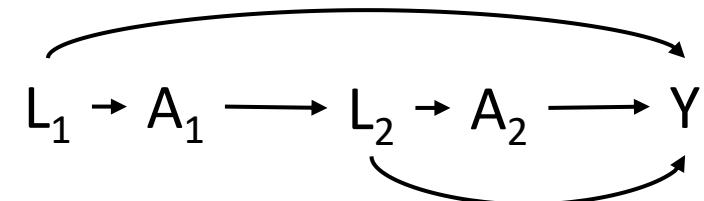
- Groups need to be similar at time zero
- Only baseline confounding



Sustained



- Groups need to be similar at time zero & during follow-up
- Baseline & time-varying confounding



Let's practice with classifying treatment strategies

Go to classpoint.app

Point strategy or sustained treatment strategy?

1. Receive bariatric surgery
2. Receive Pfizer first dose now, and second dose 3 weeks later
3. Start SGLT-2i within 3 months from now
4. Never start SGLT-2i
5. Start GLP-1RA when a cardiovascular event develops

A: point strategy

B: sustained
strategy



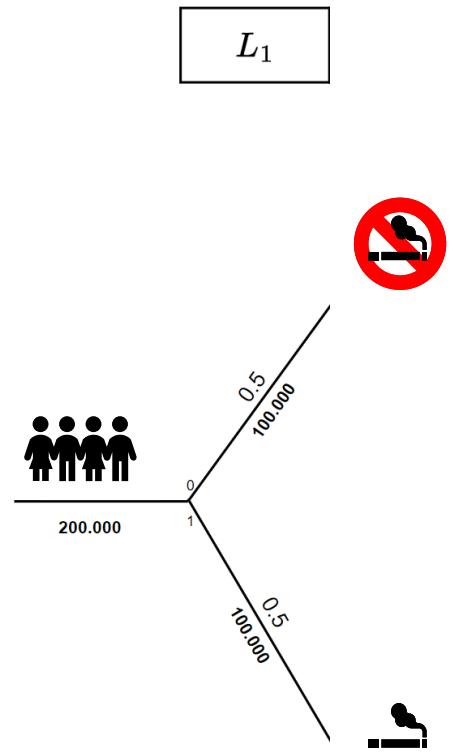
Multiple Choice

Visualizing the history of a population in a tree graph



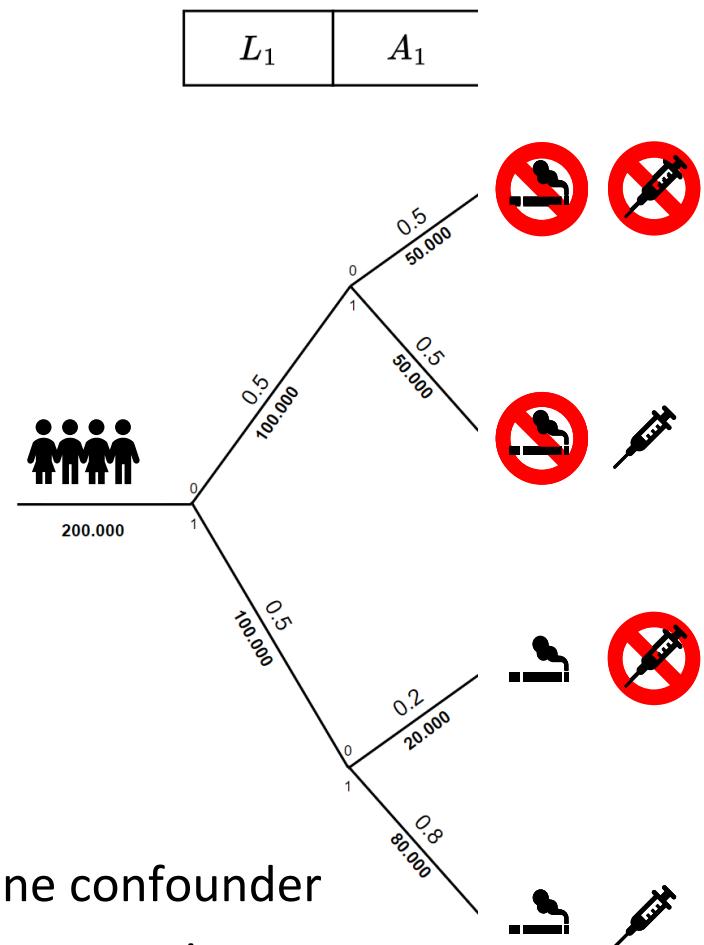
200.000

Visualizing the history of a population in a tree graph



L_1 : Baseline confounder

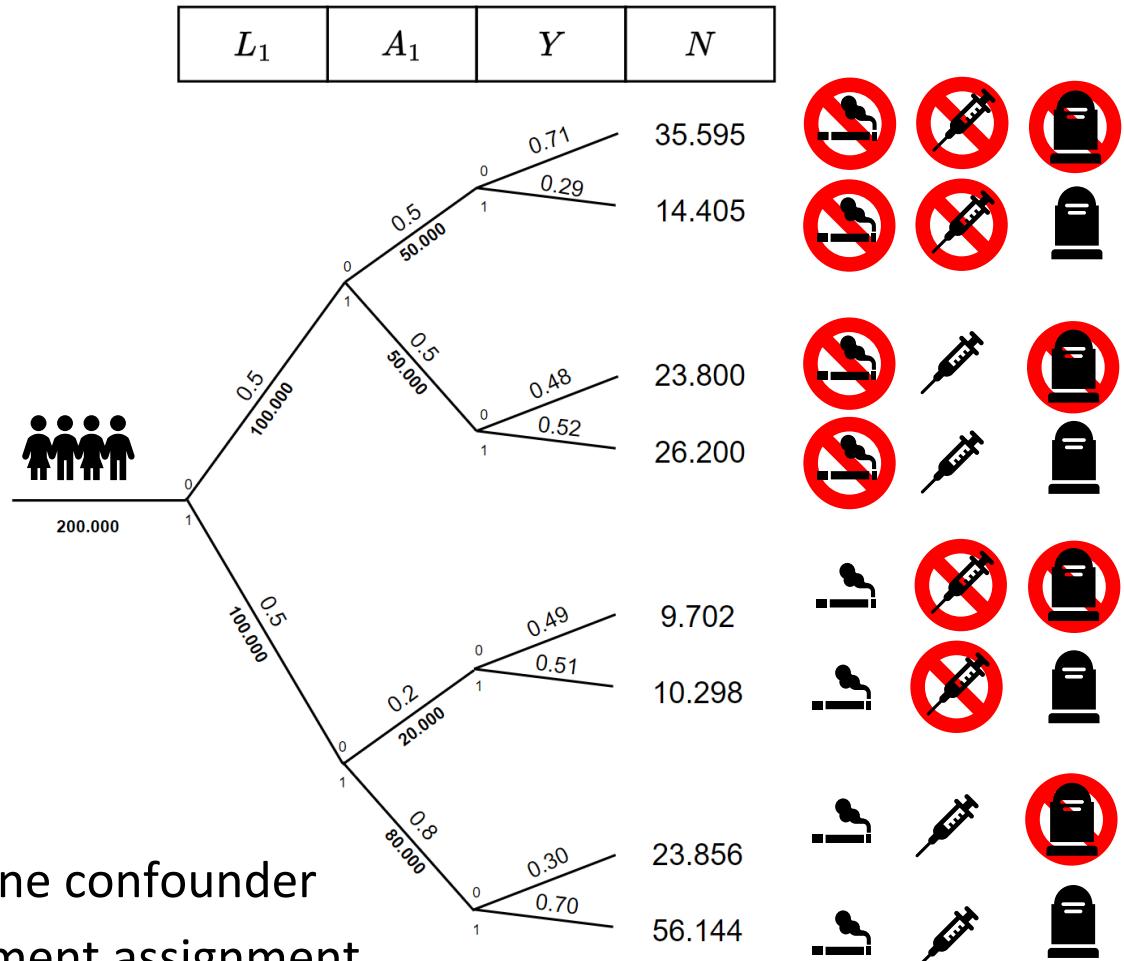
Visualizing the history of a population in a tree graph



L₁: Baseline confounder

A₁: Treatment assignment

Visualizing the history of a population in a tree graph



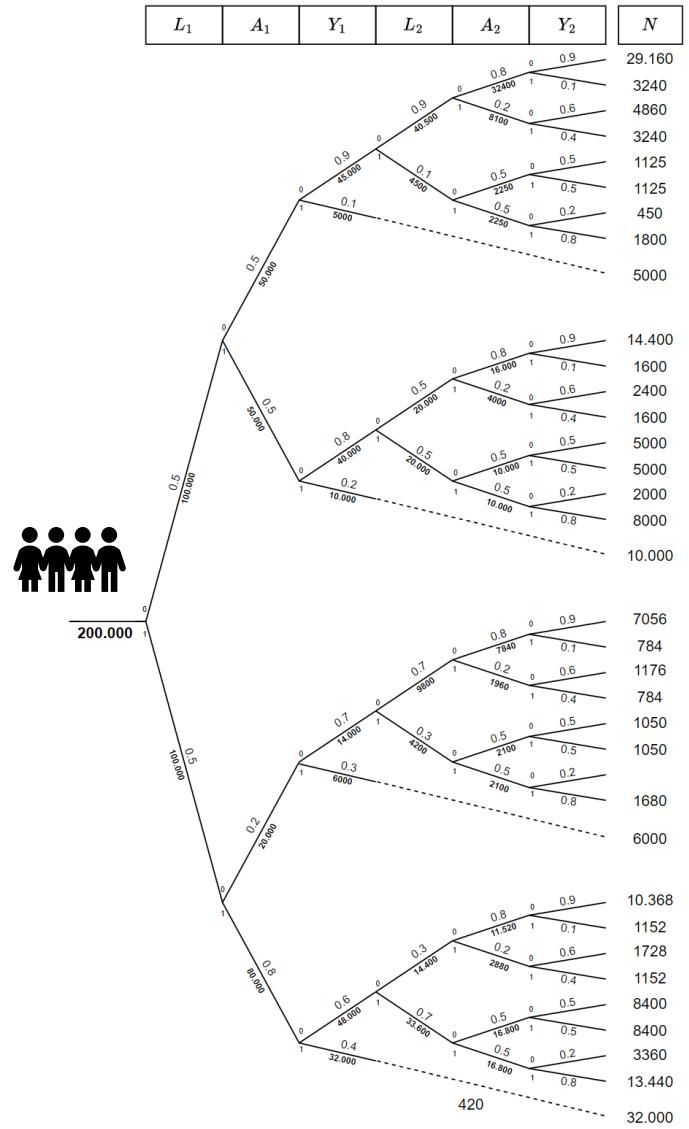
L_1 : Baseline confounder

A_1 : Treatment assignment

Y : Outcome

This is the whole tree of a **point intervention** because we only have treatment at single point in time!

Visualizing the history of a population as a tree



Quickly becomes more complex for
sustained strategies because of multiple A_t

L_1 : Baseline confounder

A_1 : Treatment at time $t=1$

Y_1 : Outcome at time $t=1$

L_2 : Time-varying confounder

A_2 : Treatment at time $t=2$

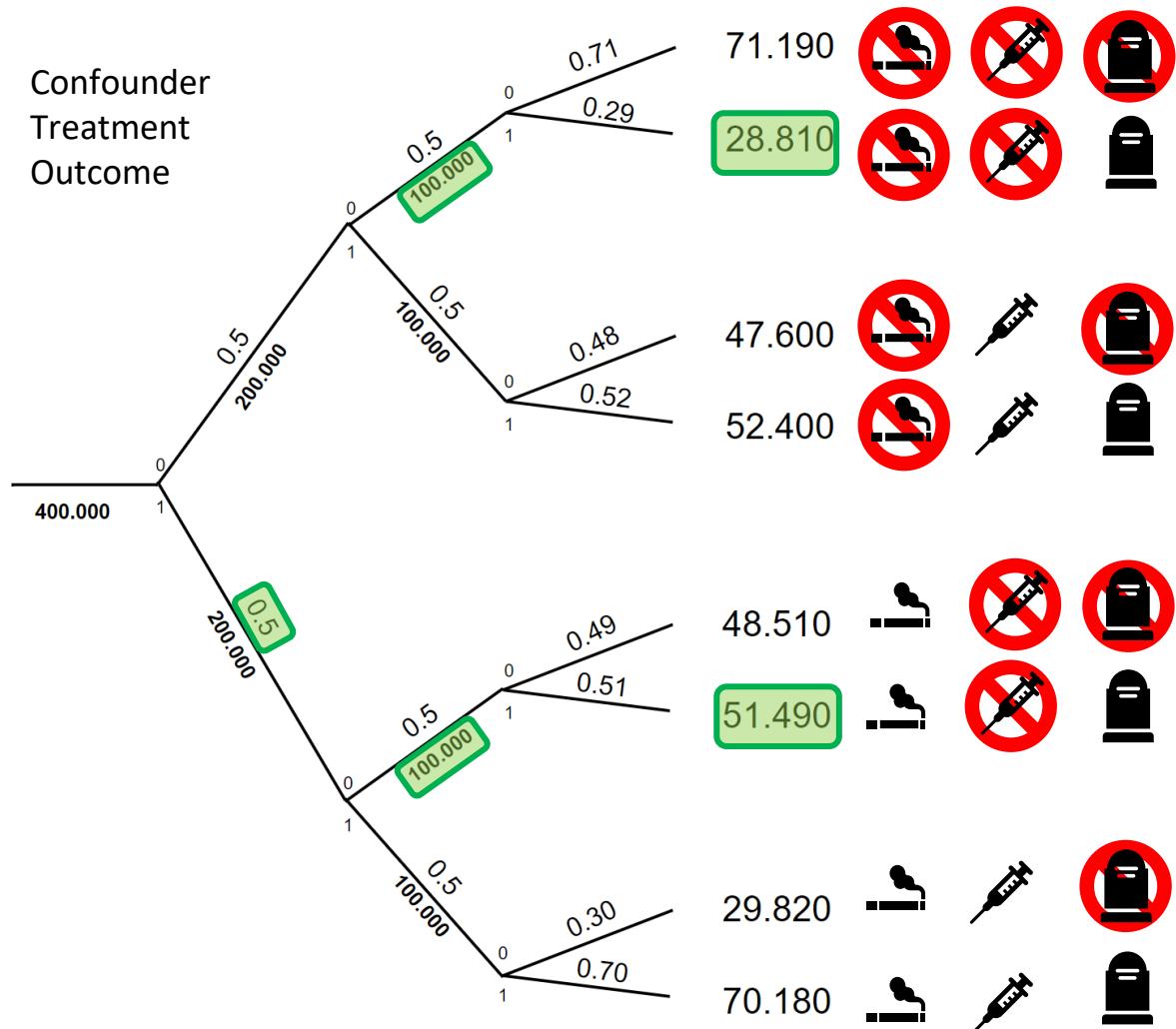
Y_2 : Outcome at time $t=2$

Some exercises

[Go to classpoint.app](https://classpoint.app)

L_1	A_1	Y	N
-------	-------	-----	-----

L: Confounder
A: Treatment
Y: Outcome



Instructions on reading the tree

1 binary confounder L (smoking)

1 binary treatment A (medication)

1 binary outcome Y (death)

Number above the lines represent proportions

Number below the lines represent number of patients

Short Answer

Question 1:

What is the probability that $L_1=1$? **0.5**

Question 2:

How many are untreated? **100.000 + 100.000 = 200.000**

Question 3:

How many die among untreated? **28.810 + 51.490 = 80.300**

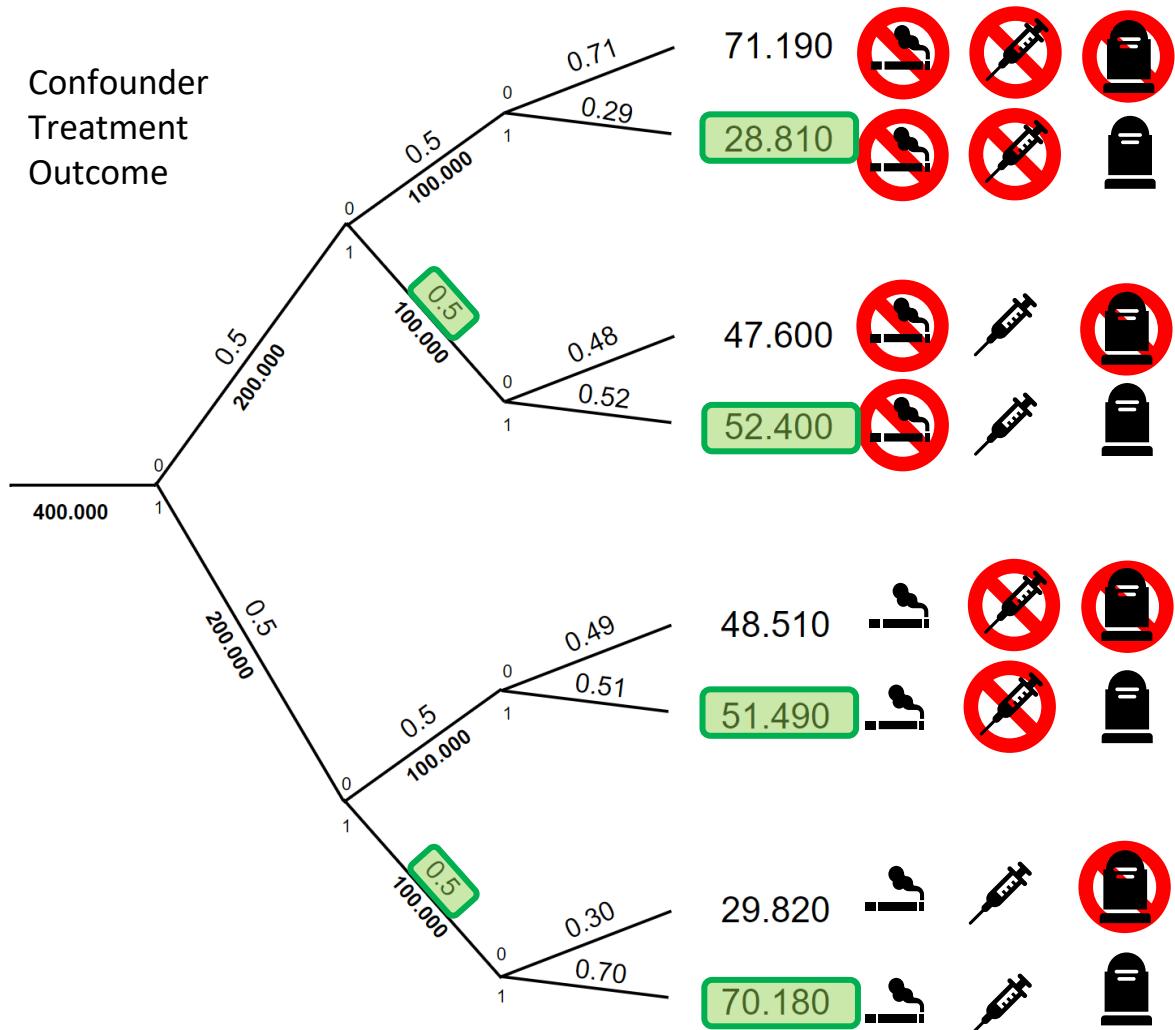
Question 4:

What is risk of death among untreated? **80.300/200.000 = 0.402**

Some exercises

L_1	A_1	Y	N
-------	-------	-----	-----

L: Confounder
A: Treatment
Y: Outcome



Instructions on reading the tree

1 binary confounder L (smoking)

1 binary treatment A (medication)

1 binary outcome Y (death)

Number above the lines represent proportions

Number below the lines represent number of patients

Question 5:

Does L_1 predict A_1 ?

No

$$\Pr[A_1 = 1 | L_1 = 1] = 0.5$$

$$\Pr[A_1 = 1 | L_1 = 0] = 0.5$$

Yes:

$$\Pr[Y = 1 | L_1 = 1] = (51.490 + 70.180) / 200.000 = 0.61$$

$$\Pr[Y = 1 | L_1 = 0] = (28.810 + 52.400) / 200.000 = 0.41$$

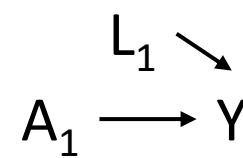
Question 6:

Does L_1 predict Y?

Question 7:

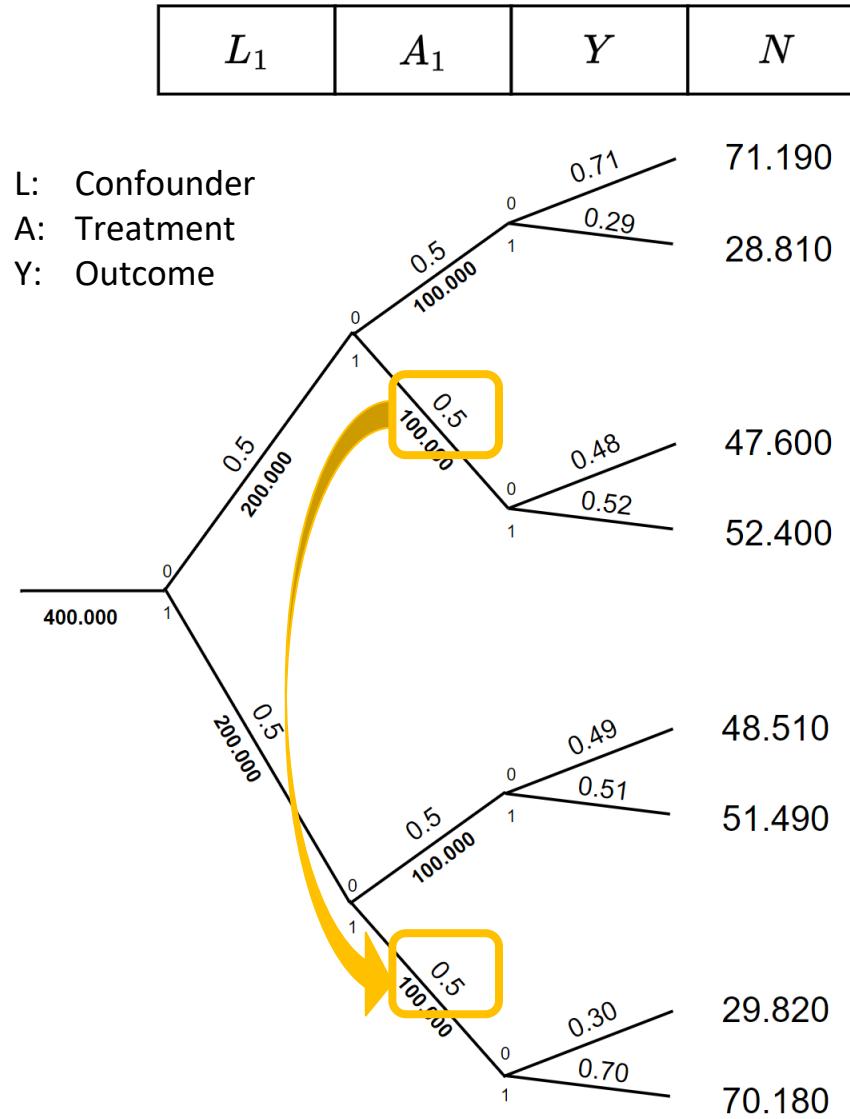
Is L_1 a confounder?

No



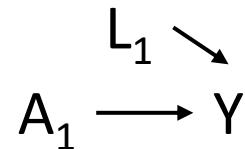
Baseline confounding

Let's check that these data indeed come from a randomized trial



In a randomized trial

- Prognostic factor does not determine whether someone receives treatment or not
- Association is causation in randomized trial



Step 3: Effect estimation

Risk among untreated

$$(28.810 + 51.490) / (100.000 + 100.000) = 0.40$$

Risk among treated

$$(52.400 + 70.180) / (100.000 + 100.000) = 0.61$$

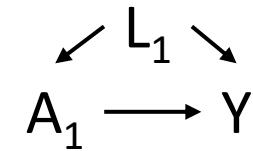
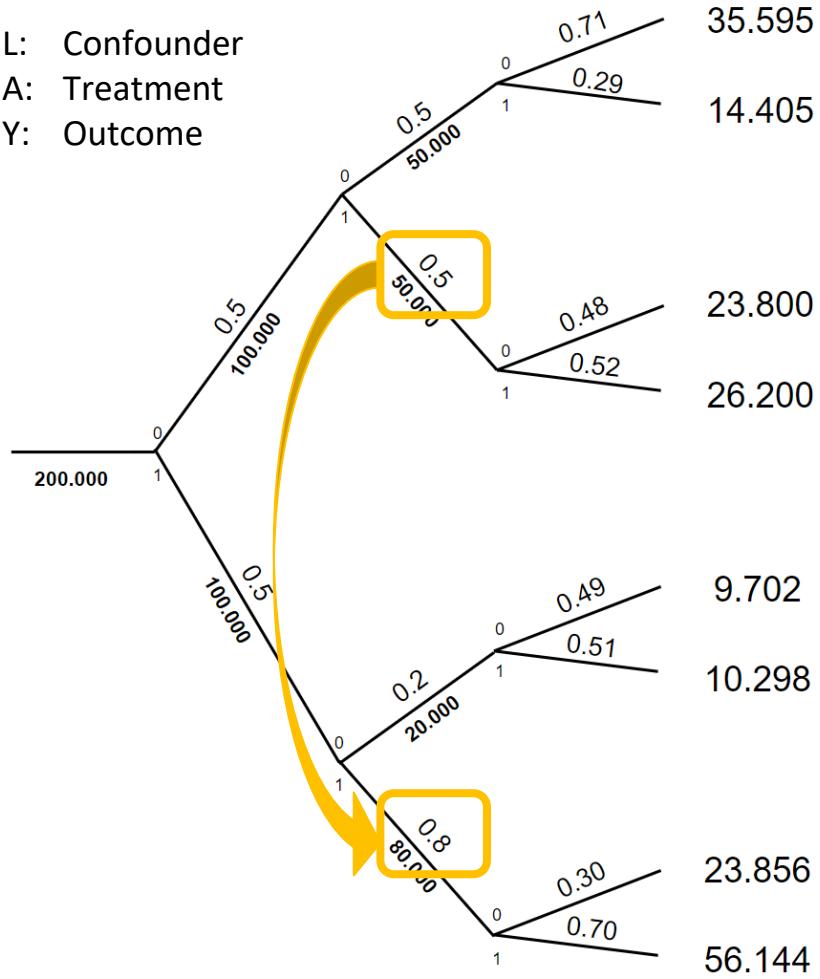
Causal risk difference: $0.61 - 0.40 = 0.21 (= 21\%)$

Causal risk ratio: $0.61 / 0.40 = 1.52$

New tree graph. Do these new data come from a randomized trial?

L_1	A_1	Y	N
-------	-------	-----	-----

L: Confounder
 A: Treatment
 Y: Outcome



In observational studies

- Prognostic factor determines whether someone receives treatment or not (L_1 = confounder)
- Association is NOT causation

Step 3: Effect estimation without adjustment for baseline confounding

Risk among untreated

$$(14.405 + 10.298) / (50.000 + 20.000) = 0.35 \neq 0.40$$

Risk among treated

$$(26.200 + 56.144) / (50.000 + 80.000) = 0.63 \neq 0.61$$

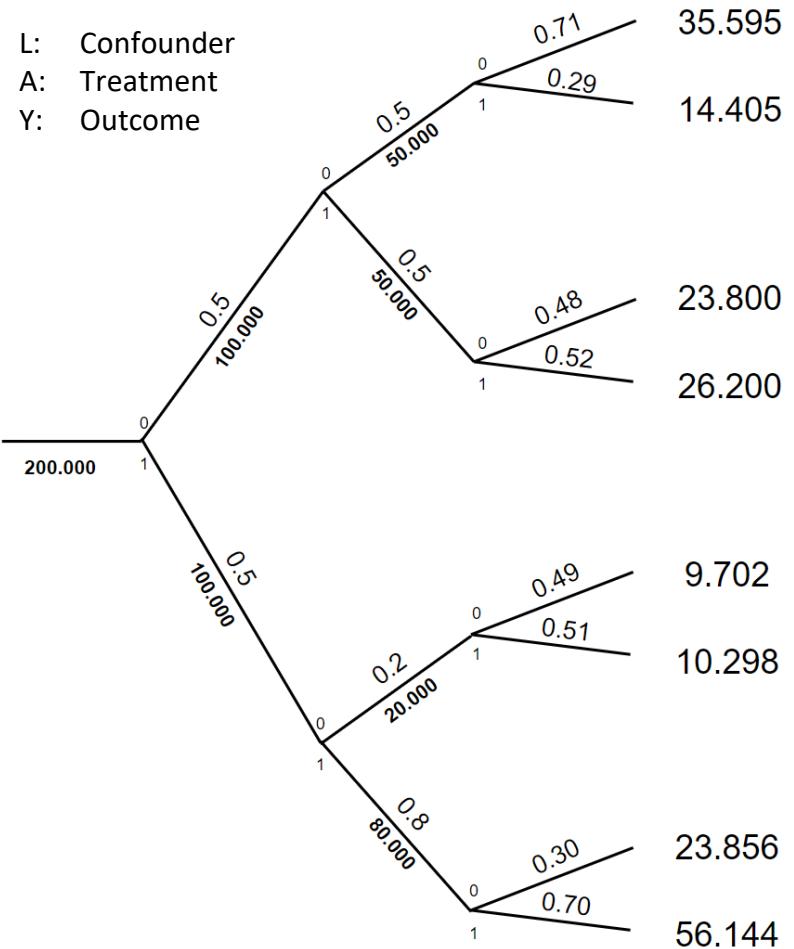
Confounded risk difference: $0.63 - 0.35 = 0.28 (= 28\%) \neq 0.21$

Confounded risk ratio: $0.63 / 0.35 = 1.80 \neq 1.52$

Adjusting for baseline confounding with weighting (IPTW)

L_1	A_1	Y	N
-------	-------	-----	-----

L: Confounder
A: Treatment
Y: Outcome



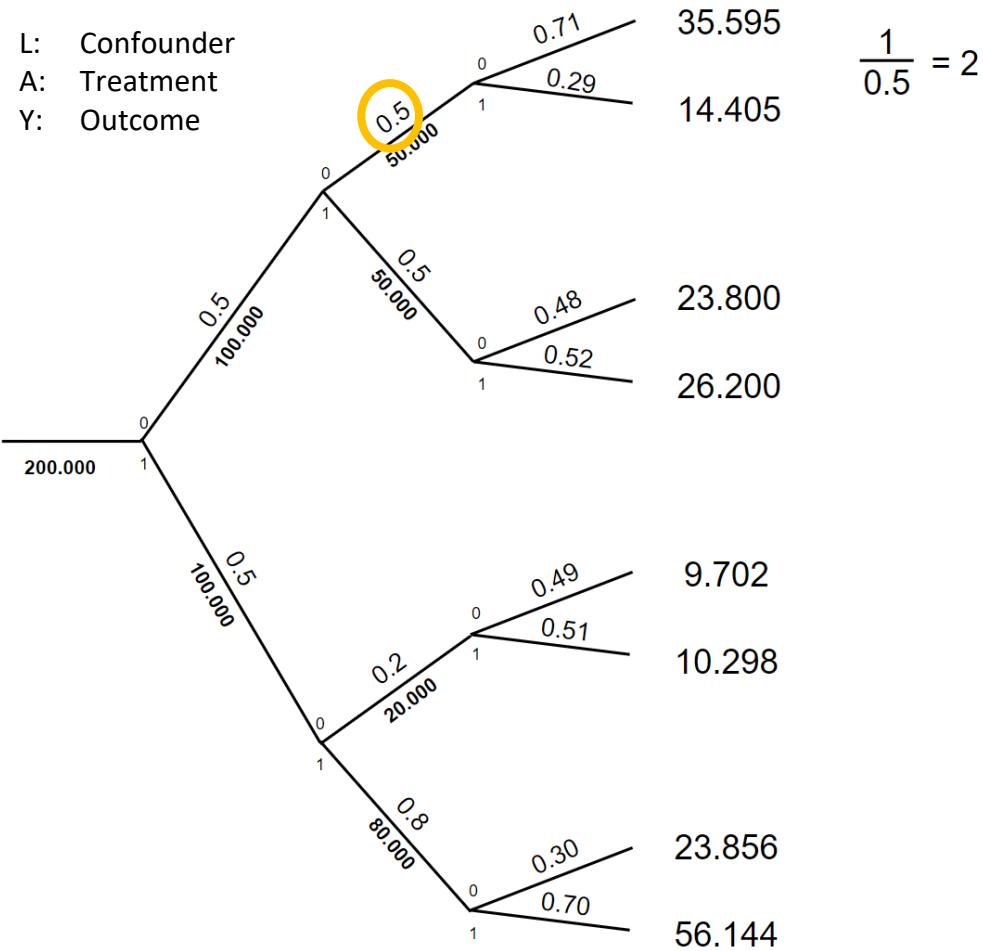
Inverse Probability of Treatment Weights are the inverse of the probability of having received your treatment history given confounders

$$\text{Here: } w_t = \frac{1}{\Pr[A_1|L_1]}$$

Adjusting for baseline confounding with weighting (IPTW)

L_1	A_1	Y	N	w_t	N_w
-------	-------	-----	-----	-------	-------

L: Confounder
 A: Treatment
 Y: Outcome



Inverse Probability of Treatment

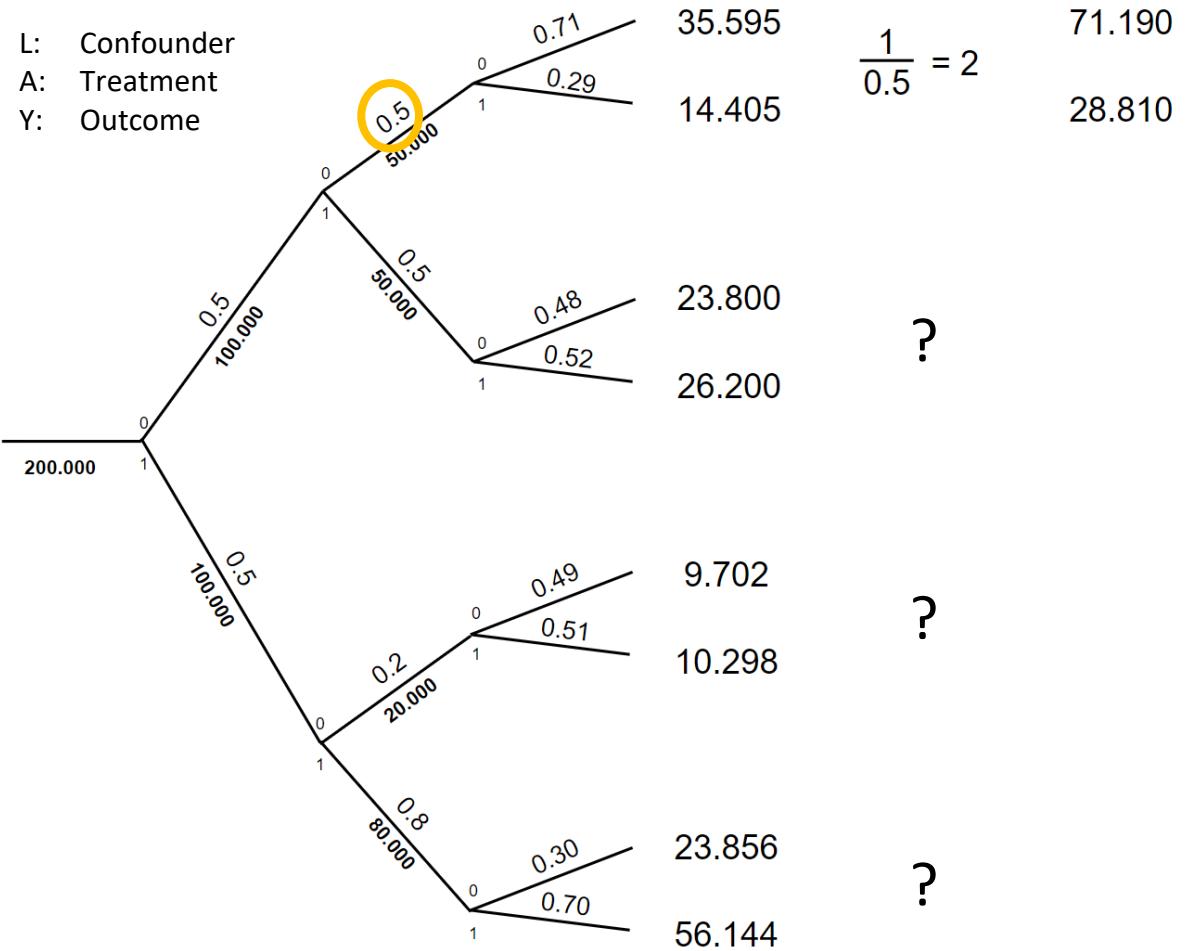
Weights are the inverse of the probability of having received your treatment history given confounders

$$\text{Here: } w_t = \frac{1}{\Pr[A_1|L_1]}$$

Adjusting for baseline confounding with weighting (IPTW)

L_1	A_1	Y	N	w_t	N_w
-------	-------	-----	-----	-------	-------

L: Confounder
 A: Treatment
 Y: Outcome



Inverse Probability of Treatment

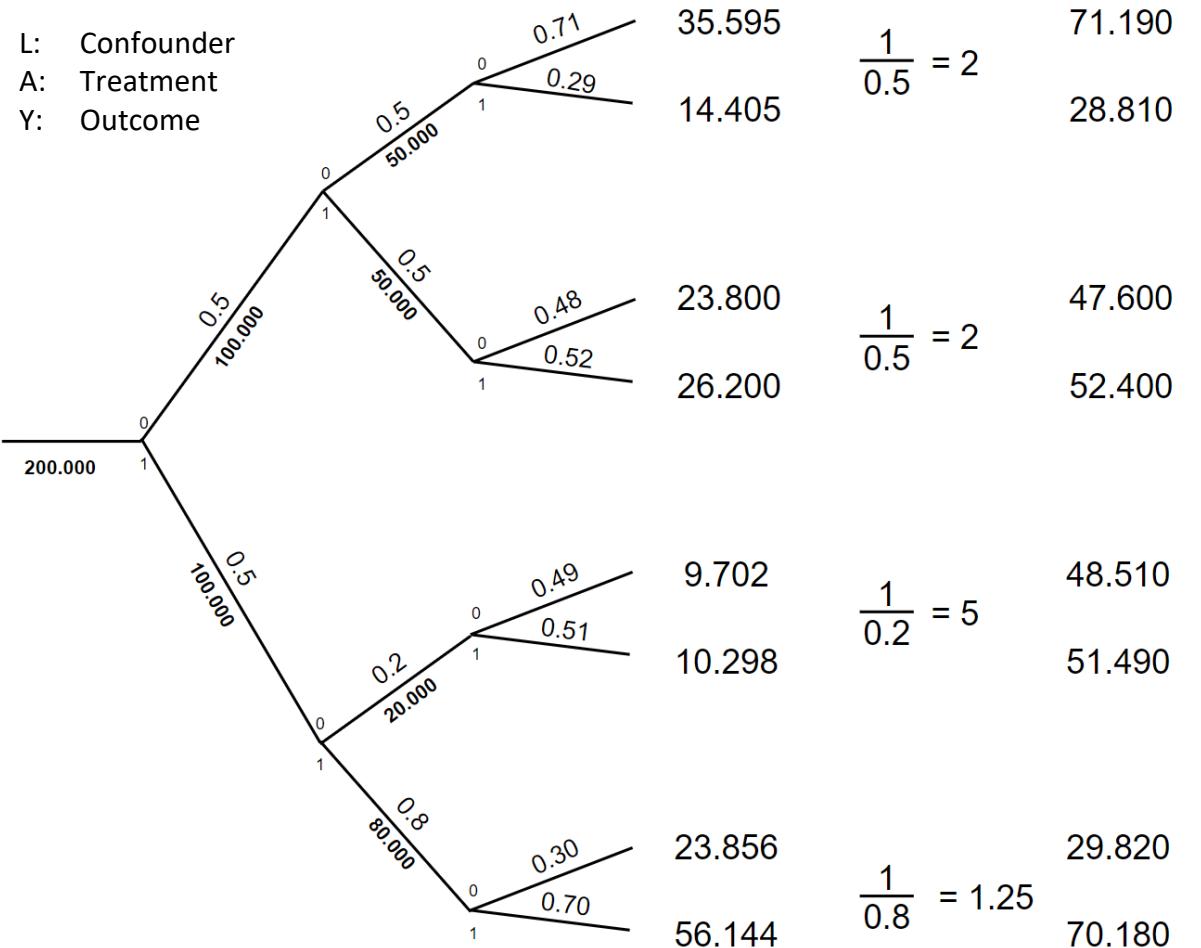
Weights are the inverse of the probability of having received your treatment history given confounders

$$\text{Here: } w_t = \frac{1}{\Pr[A_1|L_1]}$$

Adjusting for baseline confounding with weighting (IPTW)

L_1	A_1	Y	N	w_t	N_w
-------	-------	-----	-----	-------	-------

L: Confounder
 A: Treatment
 Y: Outcome



Inverse Probability of Treatment

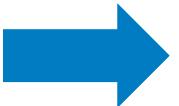
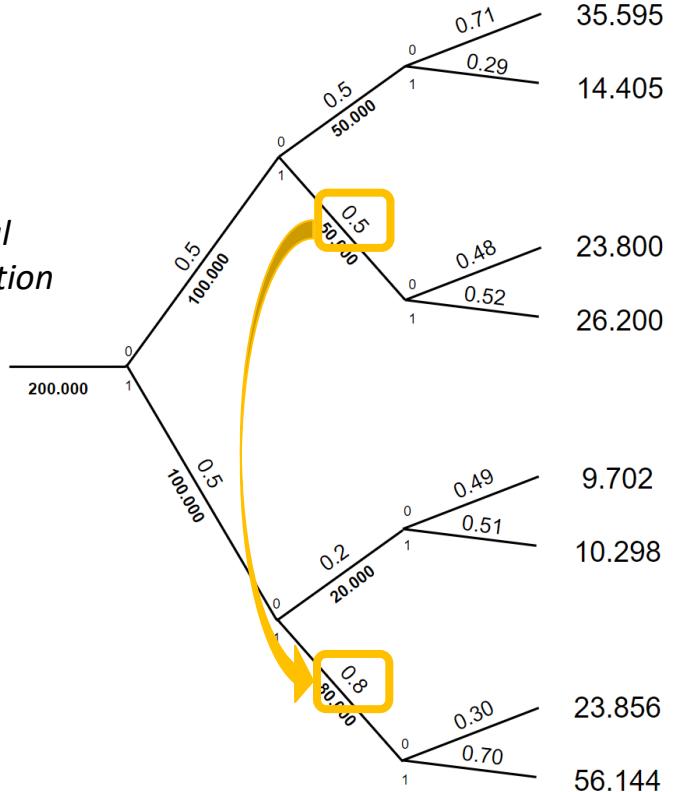
Weights are the inverse of the probability of having received your treatment history given confounders

$$\text{Here: } w_t = \frac{1}{\Pr[A_1|L_1]}$$

Turning our observational study into a randomized trial

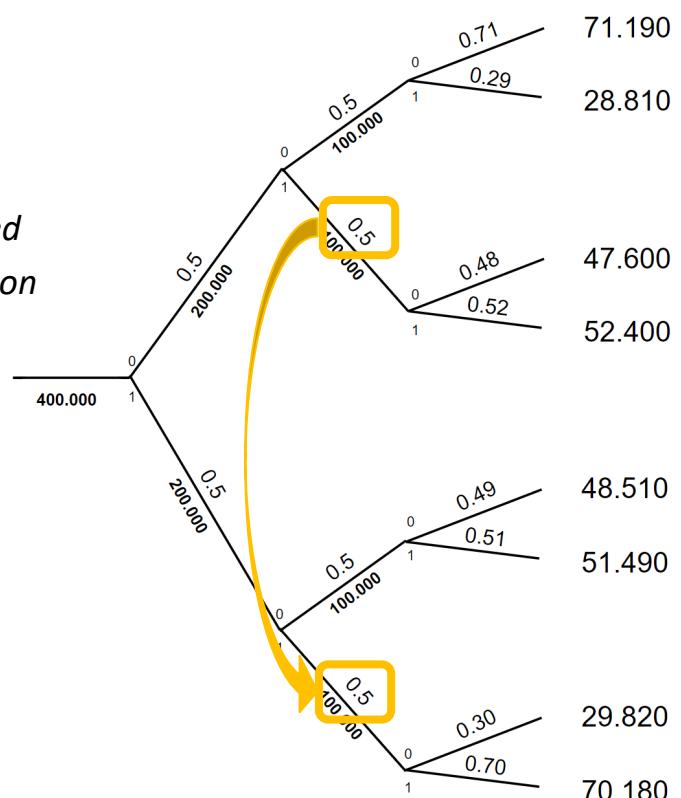
L_1	A_1	Y	N
-------	-------	-----	-----

Original population



L_1	A_1	Y	N_w
-------	-------	-----	-------

Weighted population

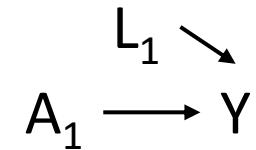
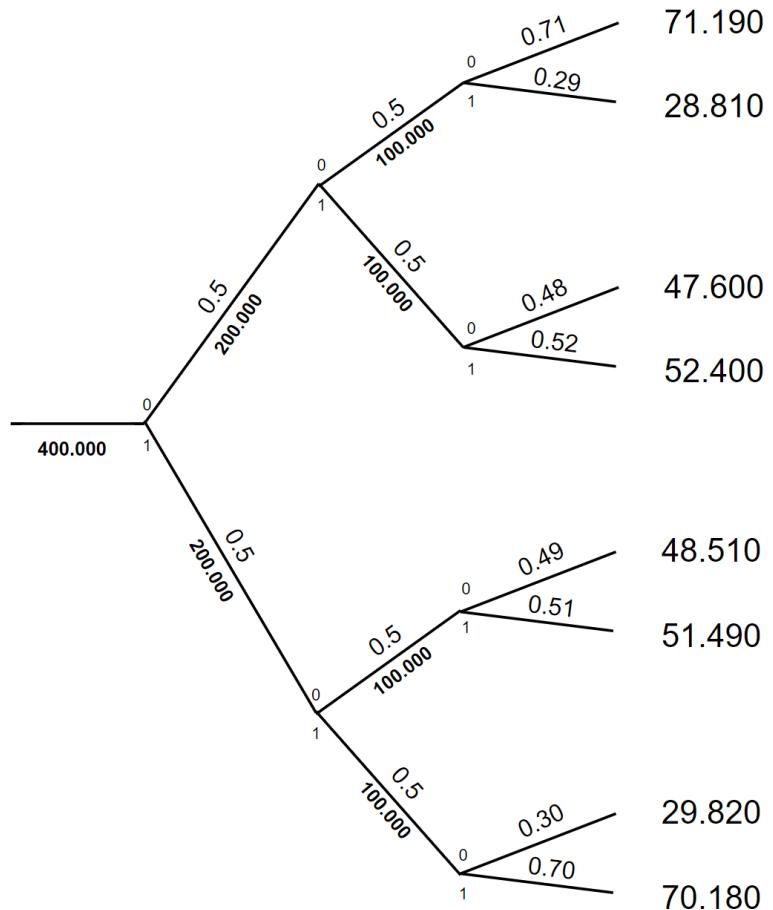


In the original population, treatment is predicted by confounder L_1

In the weighted population, treatment is no longer predicted by confounder L_1

Treatment effect estimation in the weighted pseudopopulation

L_1	A_1	Y	N_w
-------	-------	-----	-------



In weighted pseudopopulation

- Confounder no longer determines whether someone receives treatment or not
- Association is causation in the weighted pseudopopulation

Effect estimation

Risk among untreated

$$(28.810+51.490)/(100.000+100.000) = 0.40 \quad \checkmark$$

Risk among treated

$$(52.400+70.180)/(100.000+100.000) = 0.61 \quad \checkmark$$

$$\text{Causal risk difference: } 0.61 - 0.40 = 0.21 (= 21\%) \quad \checkmark$$

$$\text{Causal risk ratio: } 0.61 / 0.40 = 1.52 \quad \checkmark$$

Some comments on weighting

!

- Note that we only assumed 1 binary confounder – So we could calculate the weights nonparametrically (i.e., without models)
- In practice, there may be many confounders, which may be categorical and continuous → need to **fit models** to estimate the weights (e.g. logistic regression model)
- Note that if there are unmeasured confounders (e.g. if we had not measured L_1), we cannot use them to estimate our inverse probability of treatment weights, and our resulting treatment effects will be biased (then we have not turned our observational study into a randomized trial)

Some comments on outcome model



- In practice, we also fit a model for the outcome (e.g. a *weighted* Cox regression) since survival times are not observed for everyone (there is censoring)
- To obtain correct confidence intervals we need to take into account the weighting, e.g. with robust standard error or bootstrapping

Time-varying confounding

Recap baseline vs. time-varying confounding

Treatment strategies

Point



T_0

- Groups need to be similar at time zero
- Only baseline confounding

Sustained



T_0

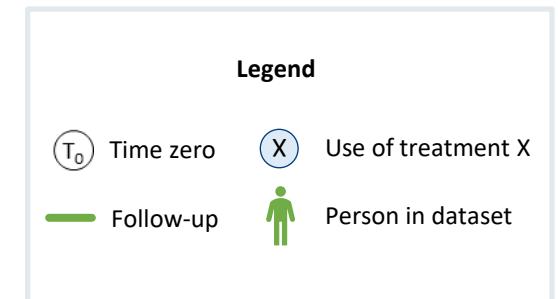
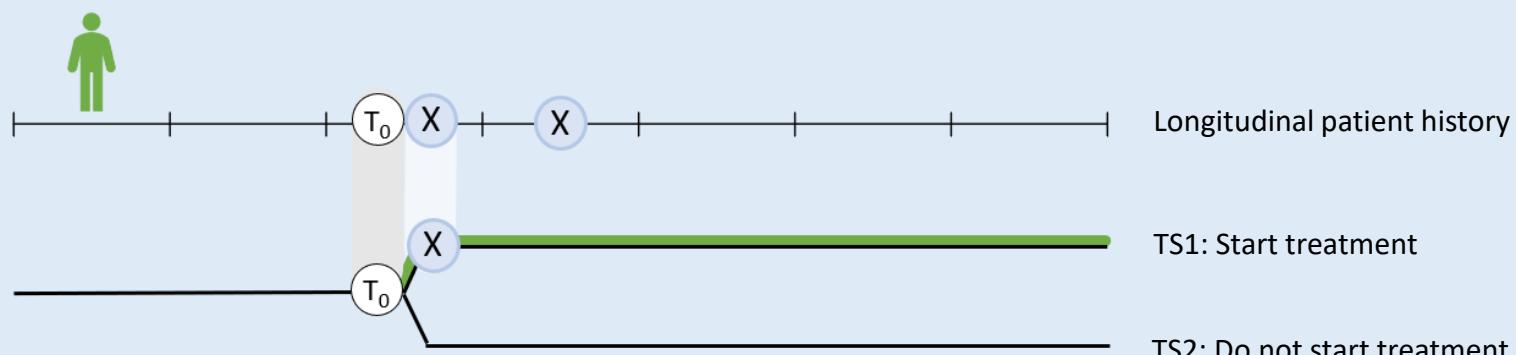
- Groups need to be similar at time zero & during follow-up
- Baseline & **time-varying confounding**

Why the effects of sustained strategies are more interesting

If we compare the point strategies “start treatment” vs. “do not start treatment”, what problems arise?

- Many people in “start treatment” group may stop treatment during follow-up
- Conversely, many people in “do not start treatment” group may start it during follow-up
- We may then find a hazard ratio of 1.0 even for a treatment known to have benefits

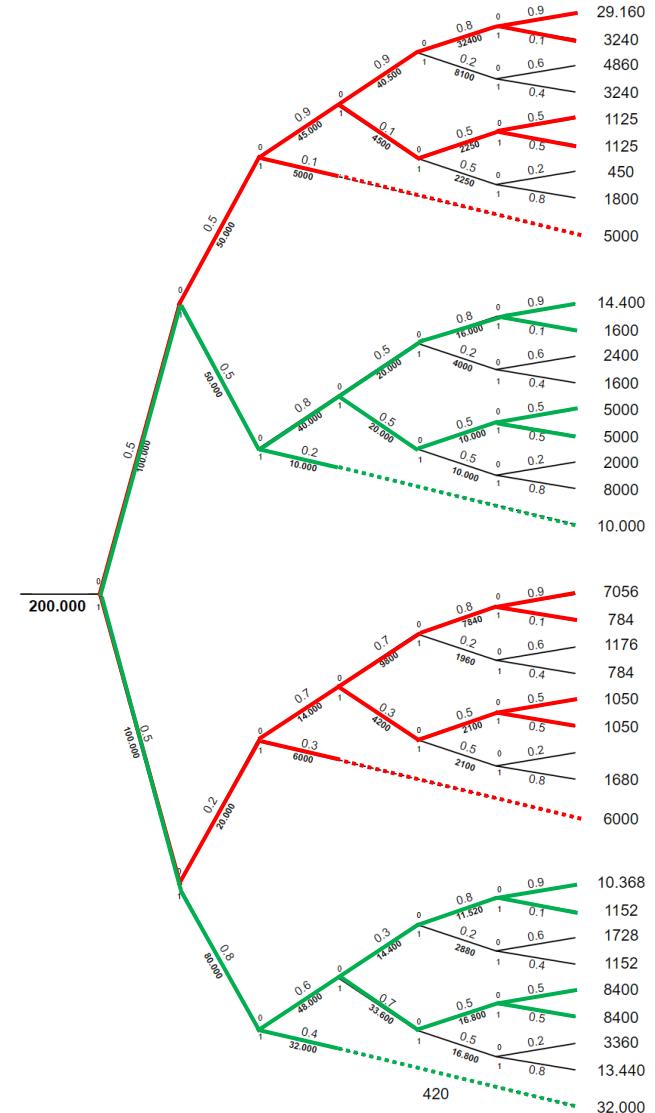
A. Single time zero



Sustained strategies: tree graph with 2+ timepoints

Go to classpoint.app

L_1	A_1	Y_1	L_2	A_2	Y_2	N
-------	-------	-------	-------	-------	-------	-----



Let's say we are interested in the sustained strategies:

- “always treat”
- “never treat”

☰ Multiple Choice

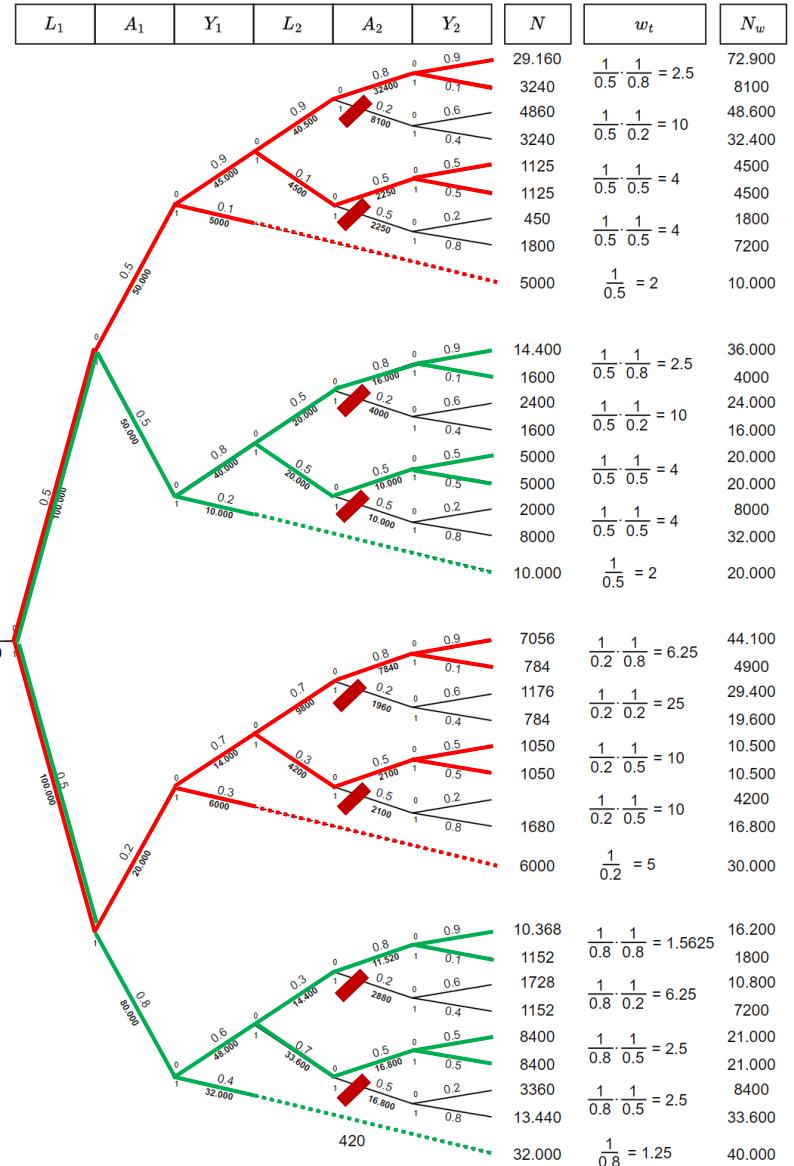
Which strategy is highlighted in the tree?

A: Always treat

B: Never treat

C: Neither

Censoring: focus only on branches of interest

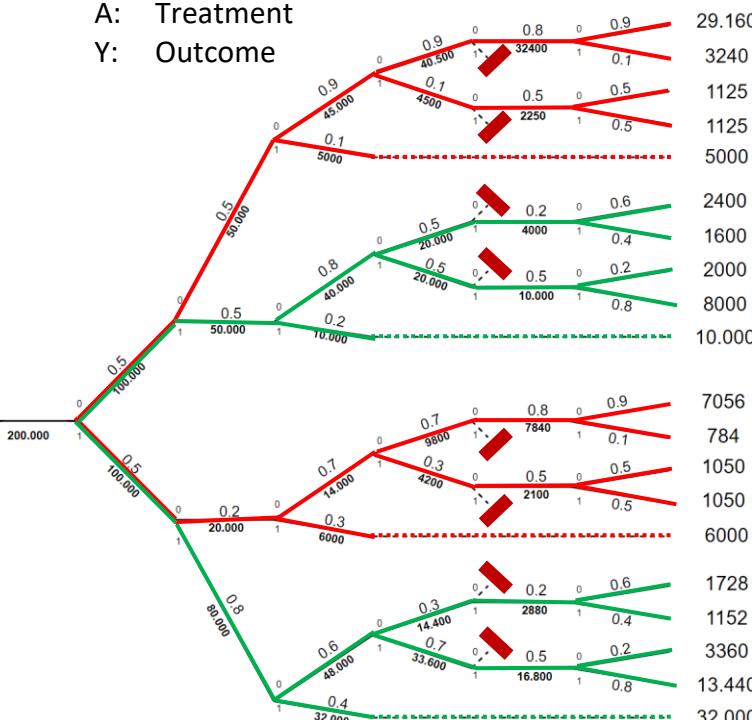


L_1	A_1	Y_1	L_2	A_2	Y_2	N
-------	-------	-------	-------	-------	-------	-----

L: Confounder

A: Treatment

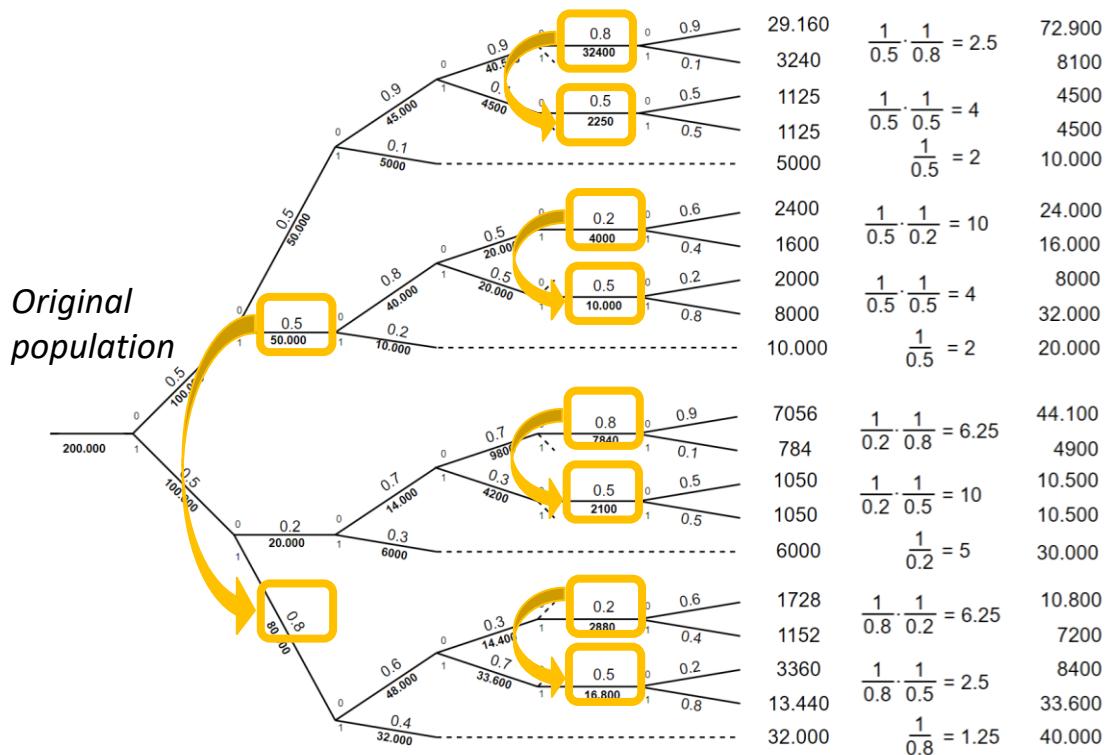
Y: Outcome



Censor patients who deviate from the strategies of interest

Turning our observational study into a sequentially randomized trial

L_1	A_1	Y_1	L_2	A_2	Y_2	N	w_t	N_w
-------	-------	-------	-------	-------	-------	-----	-------	-------

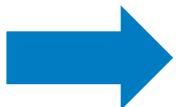
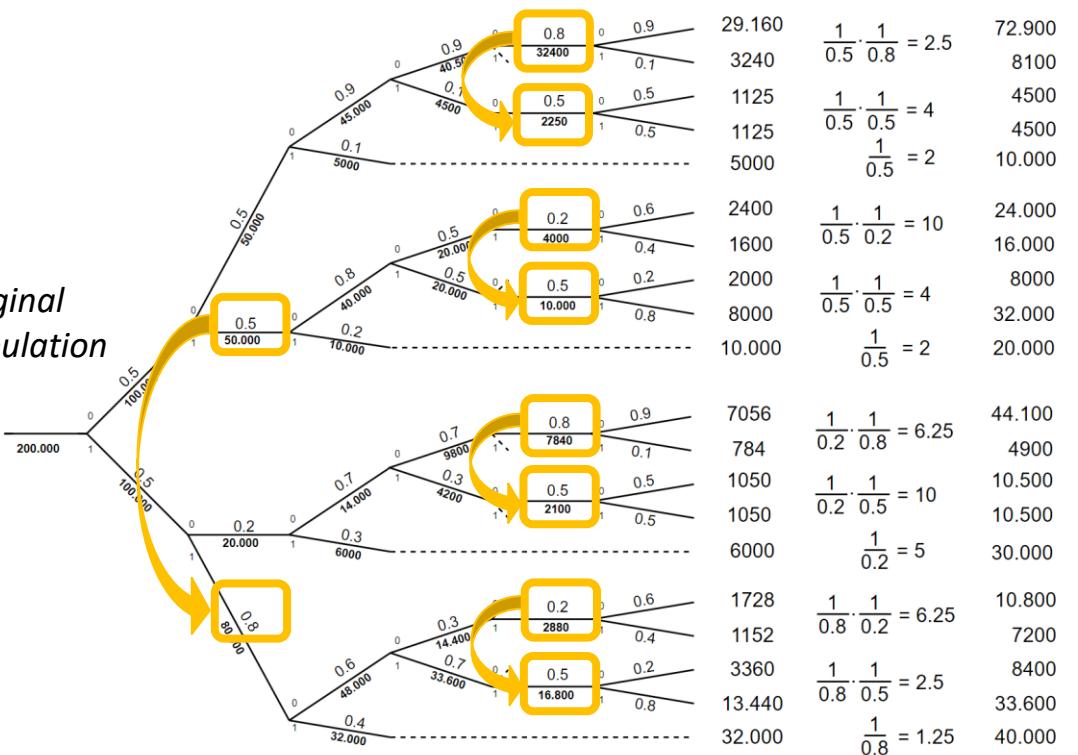


In the original population, treatment A_k is predicted by confounder L_k

Turning our observational study into a sequentially randomized trial

L_1	A_1	Y_1	L_2	A_2	Y_2	N	w_t	N_w
-------	-------	-------	-------	-------	-------	-----	-------	-------

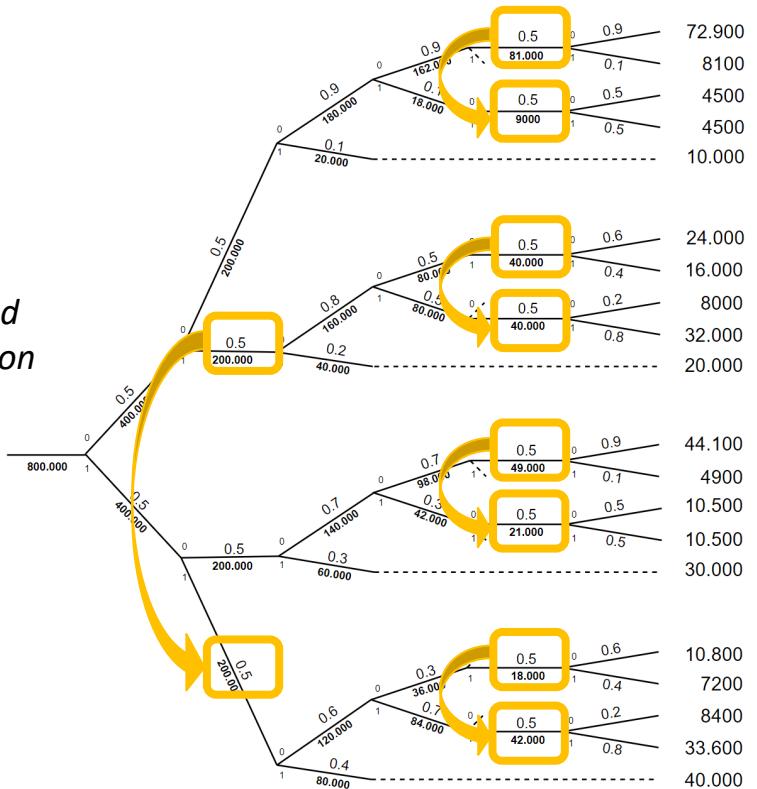
Original population



In the original population, treatment A_k is predicted by confounder L_k

L_1	A_1	Y_1	L_2	A_2	Y_2	N_w
-------	-------	-------	-------	-------	-------	-------

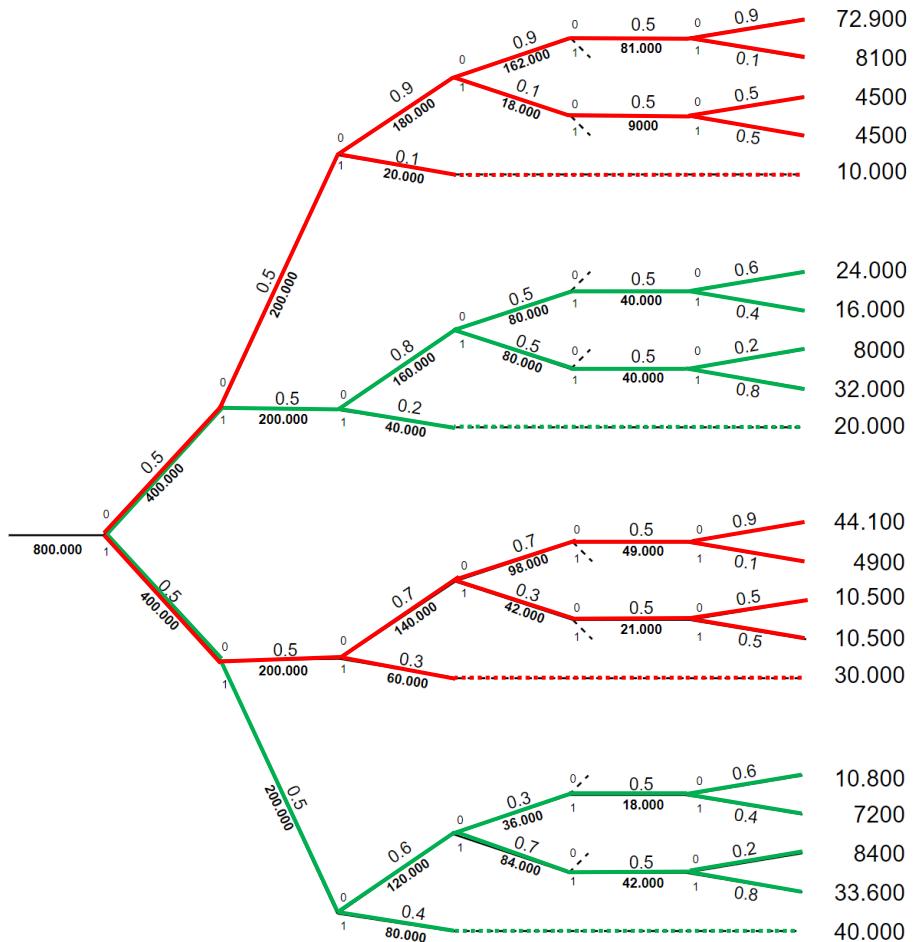
Weighted population



In the weighted population, treatment A_k is no longer predicted by confounder L_k

Treatment effect estimation in the weighted pseudopopulation

L_1	A_1	Y_1	L_2	A_2	Y_2	N_w
-------	-------	-------	-------	-------	-------	-------



Effect estimation sustained strategies

Risk among never treated

$$(8100+4500+10.000+4900+10.500+30.000)/(200.000) = 0.34$$

Risk among always treated

$$(16.000+32.000+20.000+7200+33.600+40.000)/(200.000) = 0.74$$

Causal risk difference: $0.74 - 0.34 = 0.40$ (= 40%)

Causal risk ratio: $0.74 / 0.34 = 2.19$

Effect estimation point strategies

Risk among untreated

$$(28.810+51.490)/(100.000+100.000) = 0.40$$

Risk among treated

$$(52.400+70.180)/(100.000+100.000) = 0.61$$

Causal risk difference: $0.61 - 0.40 = 0.21$ (= 21%)

Causal risk ratio: $0.61 / 0.40 = 1.52$

Conclusions

1. Important distinction between point vs. sustained strategies
2. Always need to adjust for baseline confounding
3. If interested in sustained strategies, also need to adjust for time-varying confounding
4. We showed how weighting can be used to turn the observational data into a randomized or sequentially randomized trial
5. Results are biased if there are unmeasured confounders

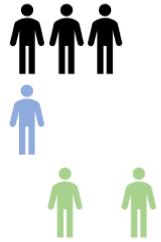
Questions

e.l.fu@lumc.nl



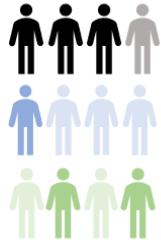
Censoring & weighting on a group-level

Artificial censoring



Censored during follow-up if not following strategy of interest

Weighting



Uncensored replicates (dark color) are upweighted to account for censored replicates (light color) with similar characteristics

Standardization (time-fixed exposure)

Standardization – Identification proof

$$\begin{aligned} E[Y^a] &= \sum_l E[Y^a | L = l] * \Pr[L = l] && \text{By the law of total expectation} \\ &= \sum_l E[Y^a | A = a, L = l] * \Pr[L = l] && \text{By exchangeability assumption} \\ &= \sum_l E[Y | A = a, L = l] * \Pr[L = l] && \text{By consistency assumption} \end{aligned}$$

We have now expressed a **counterfactual quantity** as something that does not contain counterfactuals and that we can **estimate from our data**

Standardization on the tree graph

Standardization as simulation

Equivalence of standardization and IPW

Standardization for ATT



G-formula (aka standardization in the time-varying setting)

G-formula – Identification proof

G-formula as simulation

Equivalence of G-formula and IPW

Probability review (taken from Robins' lecture)

