

Correction methods for unmeasured confounding in non-interventional studies

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Introduction

Propensity Score Calibration

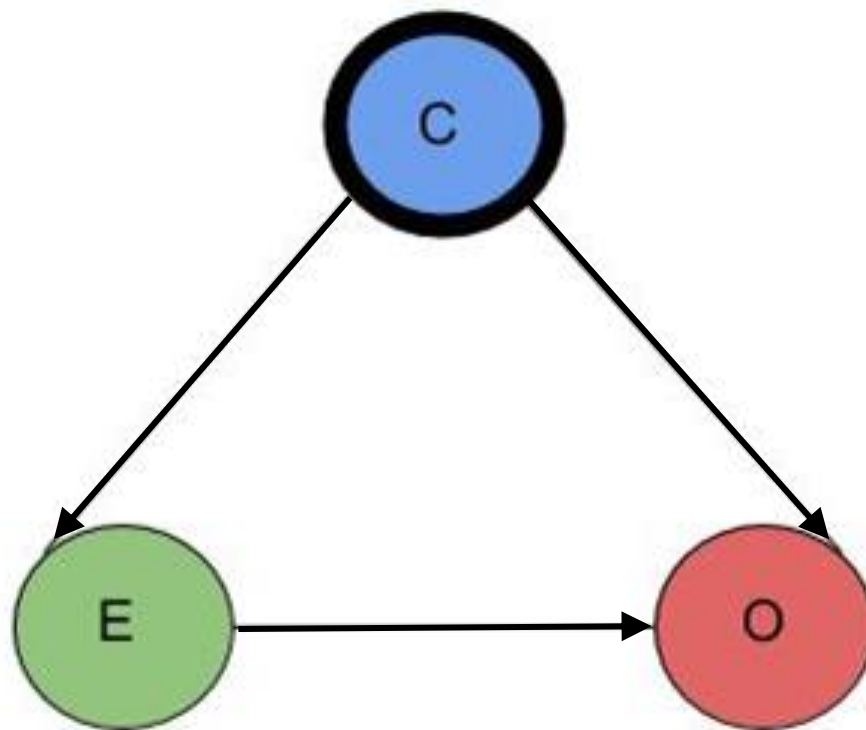
Monte Carlo simulation

Conclusions

Confounding

E = Exposure
O = Outcome
C = Confounder

E = Treatment admin
O = Death
C = Age

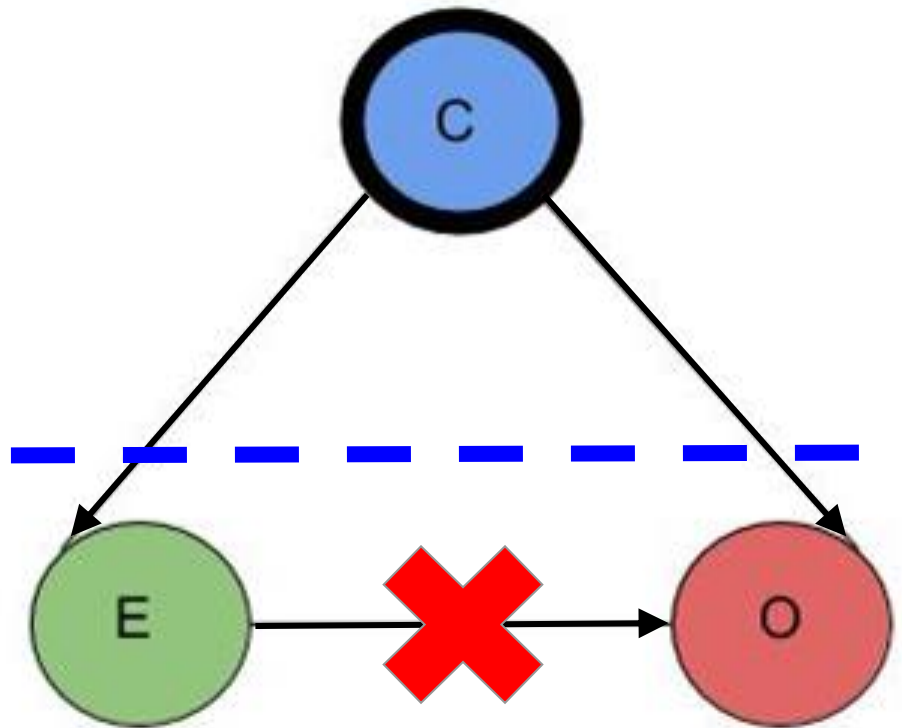


Confounding definitions

“Confounding may be considered a confusion of effects. [...] A variable must be associated with *both the exposure under study and the disease* under study to be a **confounder**. [...] A confounding factor must be associated with disease occurrence *apart* from its association with exposure.”

Confounding

E = Exposure
O = Outcome
C = Confounder



When the confounder is **not** taken into **account** in the analysis, the estimation of the treatment effect is affected by **bias**.

RANDOMIZED CLINICAL TRIALS

Randomization of treatment assignment allows to avoid systematic associations between exposure and patient characteristics, hence confounding.

The estimator of treatment effect is considered UNBIASED (even if...)



NON-INTERVENTIONAL STUDIES

It is often the case that patient characteristics or other factors are **determinants** of treatment assignment and also of the outcome of interest (e.g. confounding by indication).

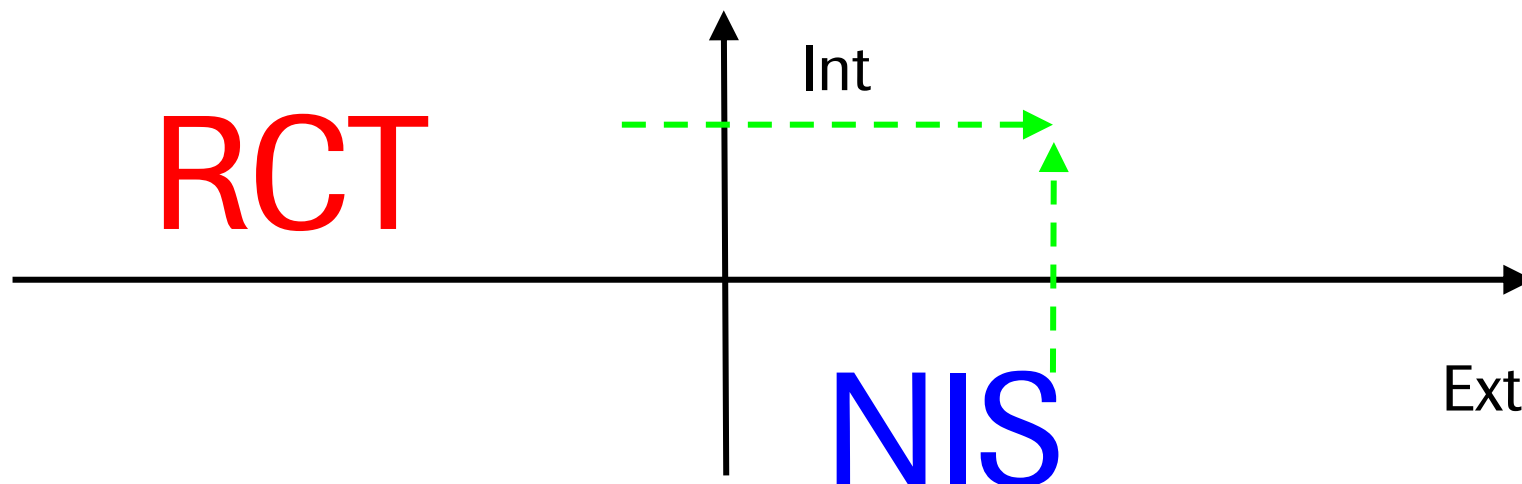
The estimator of treatment effect is considered BIASED.



External vs Internal Validity

Internal : the validity of the inferences drawn as they pertain to the members of the source population of the study. → **Reliable estimates**

External : measure of how much the previous results are **generalizable** to a target population or to a more general set of circumstances.



RCT = Randomized Clinical Trials

NIS= Non-Interventional Studies

Unmeasured confounding

Bias can be avoided accounting for the confounder in the analysis, hence removing the effect of **measured confounding**.

In several contexts this is not possible:

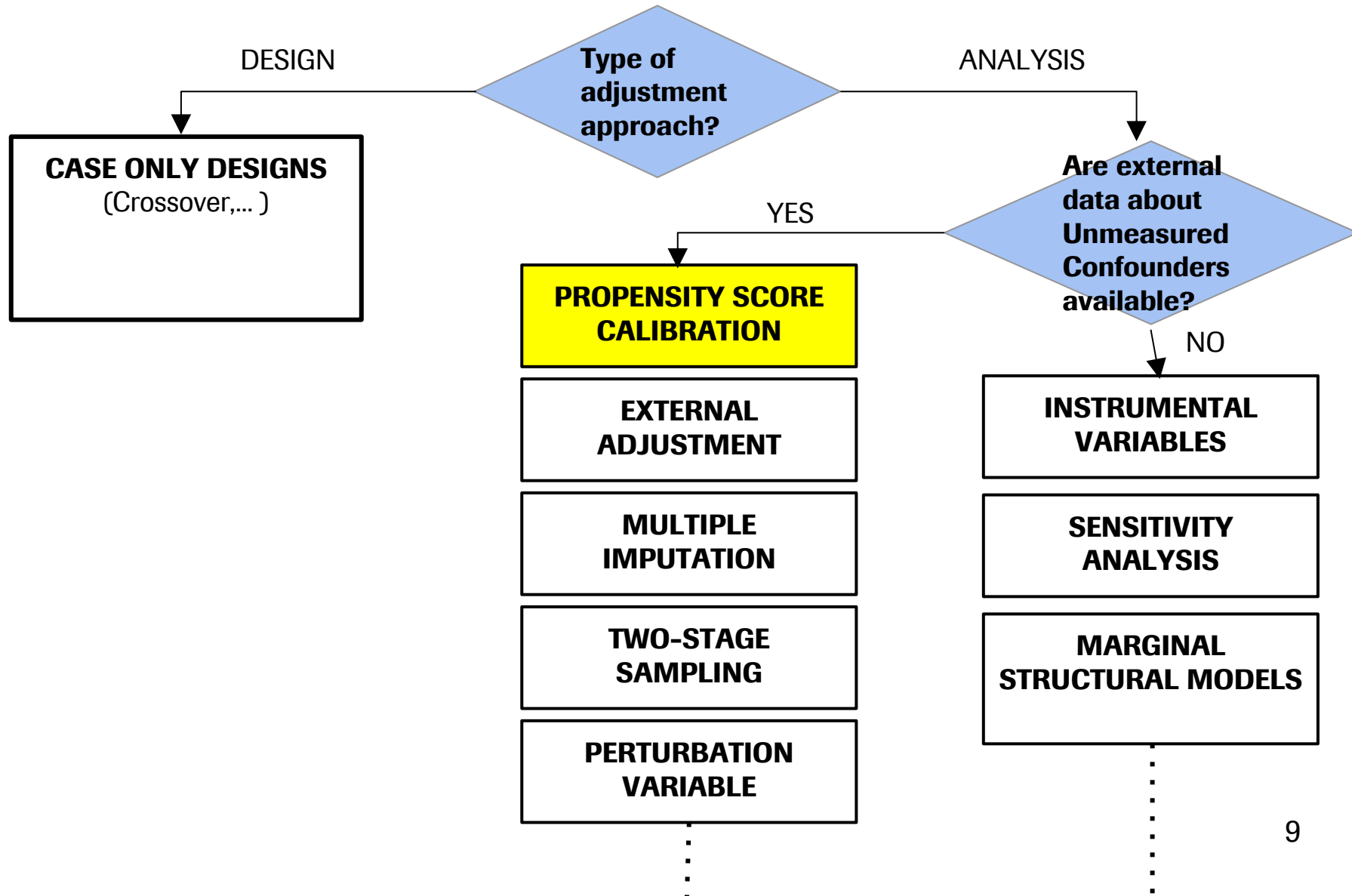
- Measurement errors
- Unfeasible
- Missing
- ...



**Unmeasured
confounding**

Both measured and unmeasured confounding can be present in a study.

Unmeasured confounding: methods



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Propensity Score

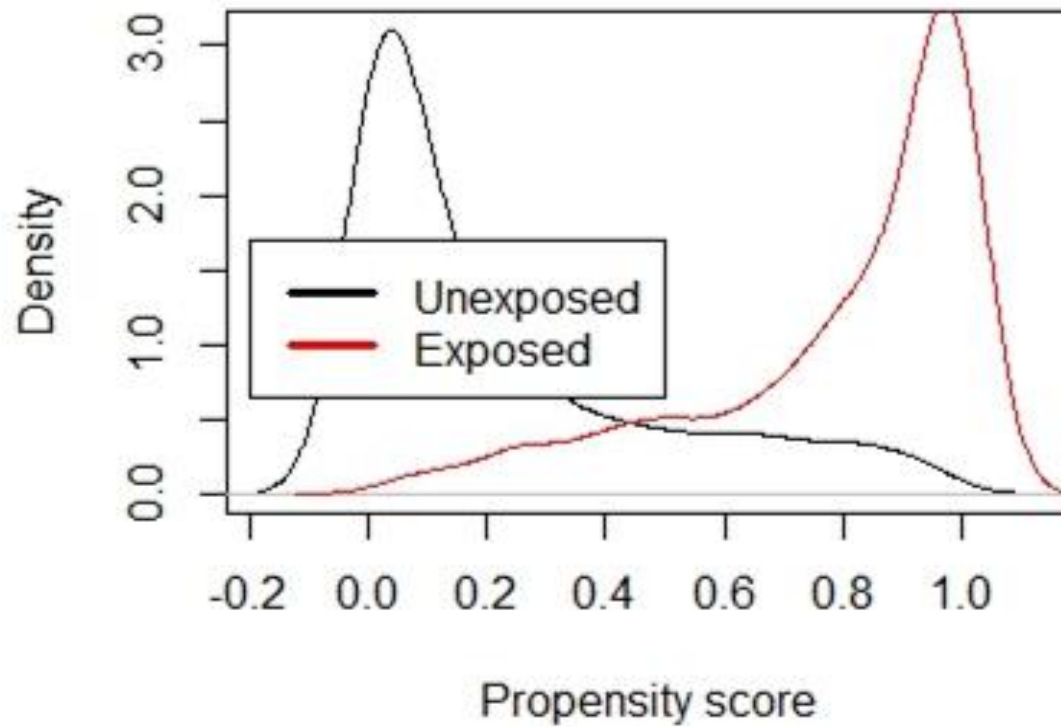
The method aims at **summarizing** covariates through a unique variable (the PS) such that treatment and control groups are **comparable** according to the distribution of covariates within the same PS level and any effect can be attributed to treatment.

$$\text{PS} = e(x) = P(\text{treatment} \mid X=x)$$

where X is the vector of covariates.

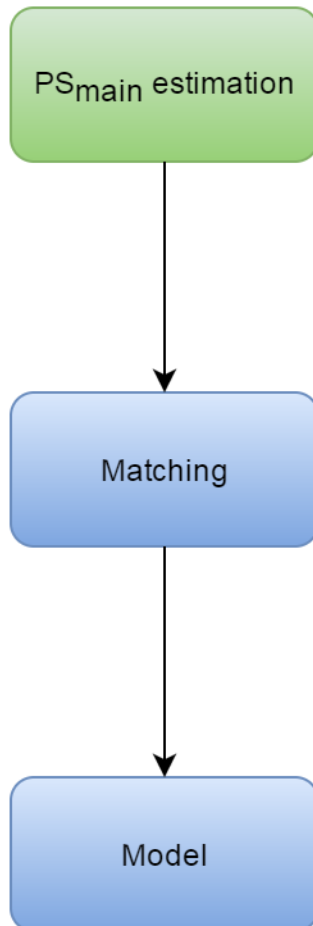
This can be achieved by **logistic regression**.

Propensity Score

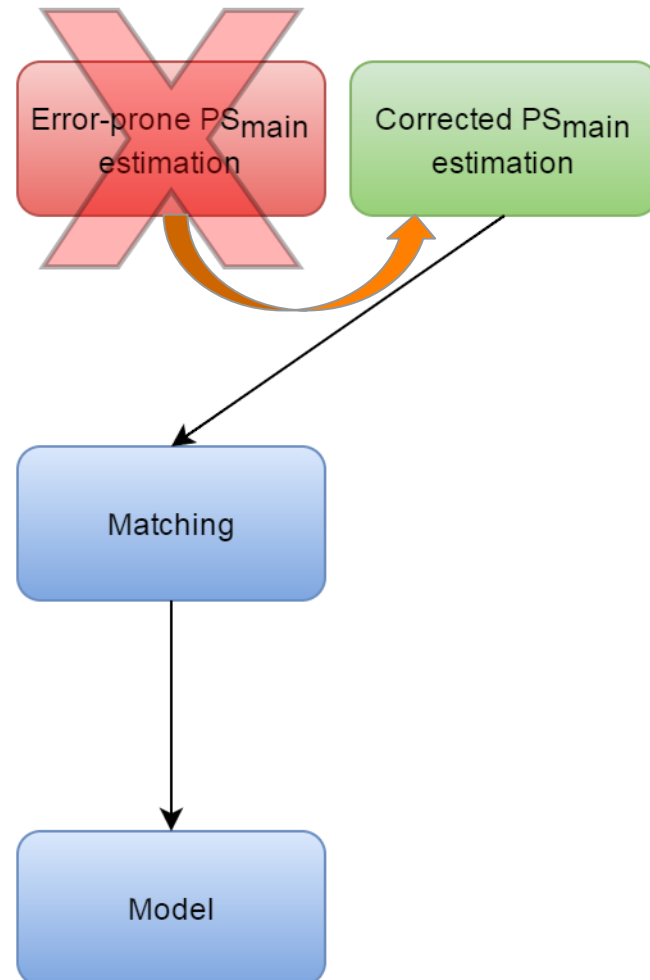


Measured vs Unmeasured Confounding using Propensity Score (PS)

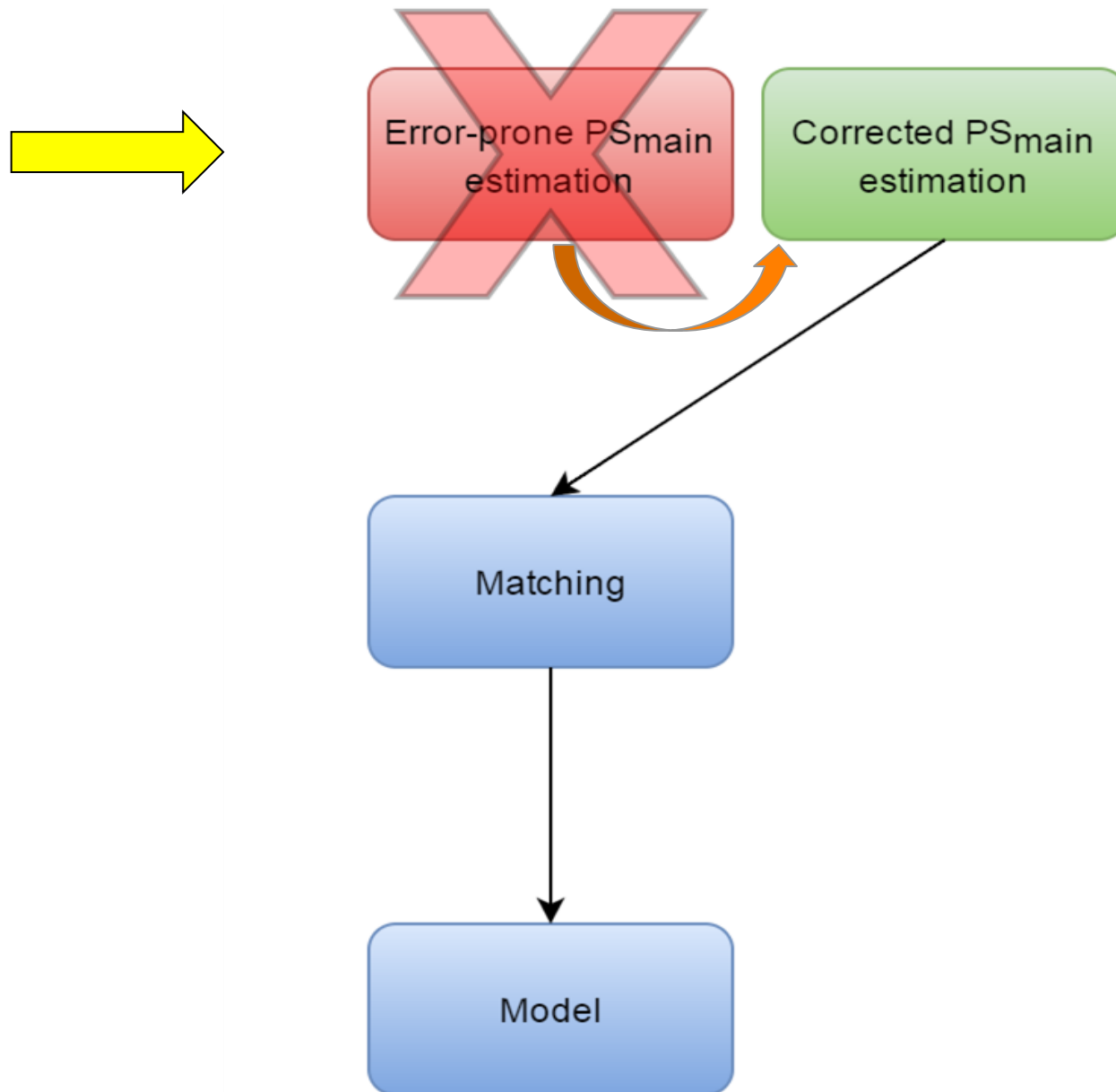
Measured



Unmeasured

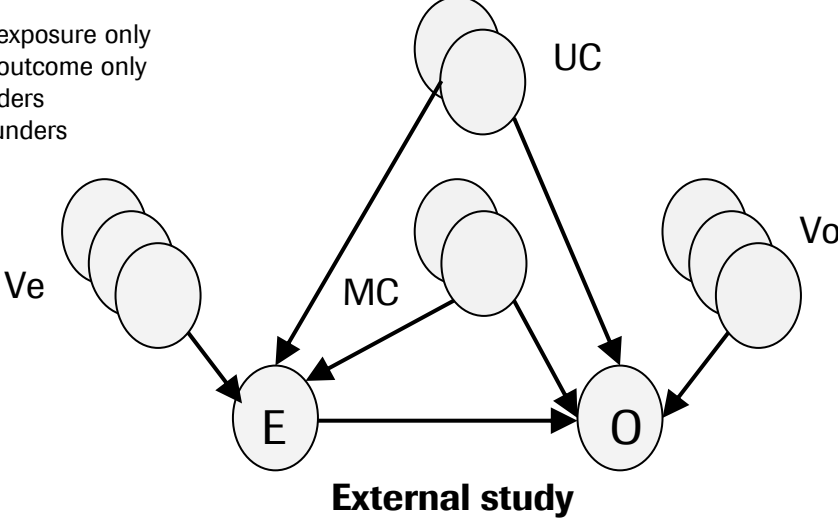
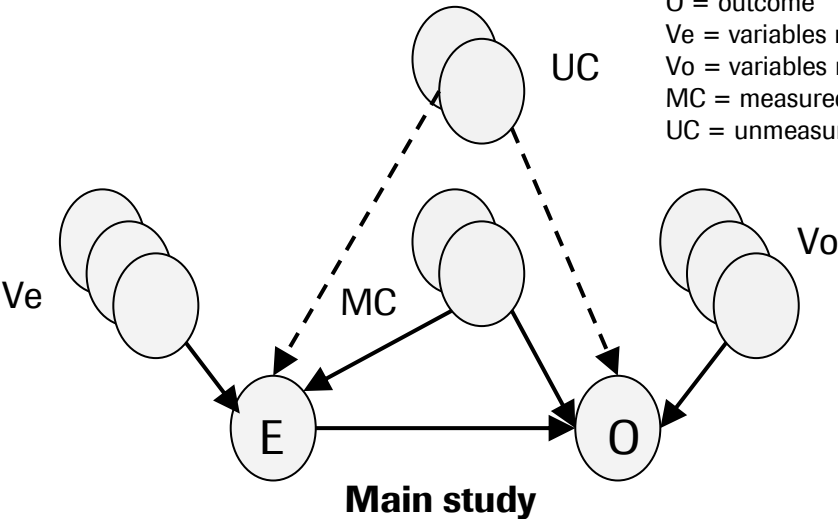


Unmeasured Confounding using PS



Frequentist Propensity Score Calibration

E = exposure
 O = outcome
 Ve = variables related to exposure only
 Vo = variables related to outcome only
 MC = measured confounders
 UC = unmeasured confounders

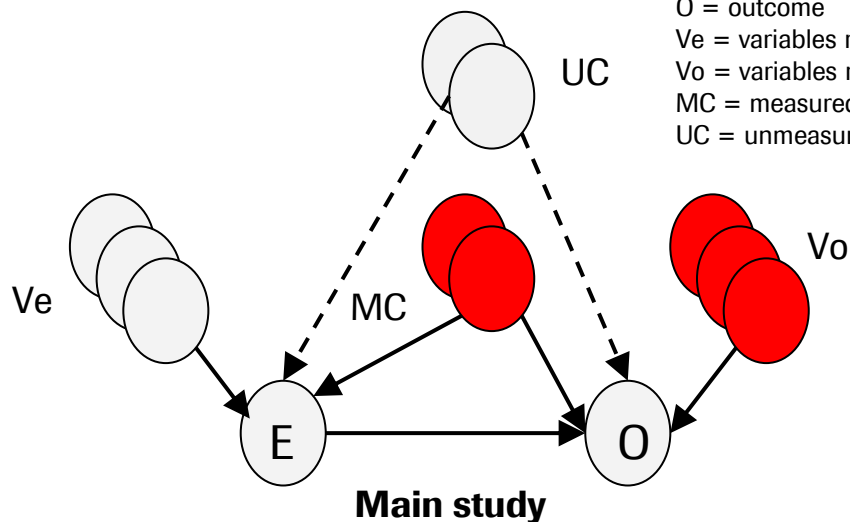


E	O	Ve	Vo	MC	UC
x	x	x	x	x	?
x	x	x	x	x	?
x	x	x	x	x	?

E	O	Ve	Vo	MC	UC
x	(x)	x	x	x	x
x	(x)	x	x	x	x
x	(x)	x	x	x	x

Frequentist Propensity Score Calibration

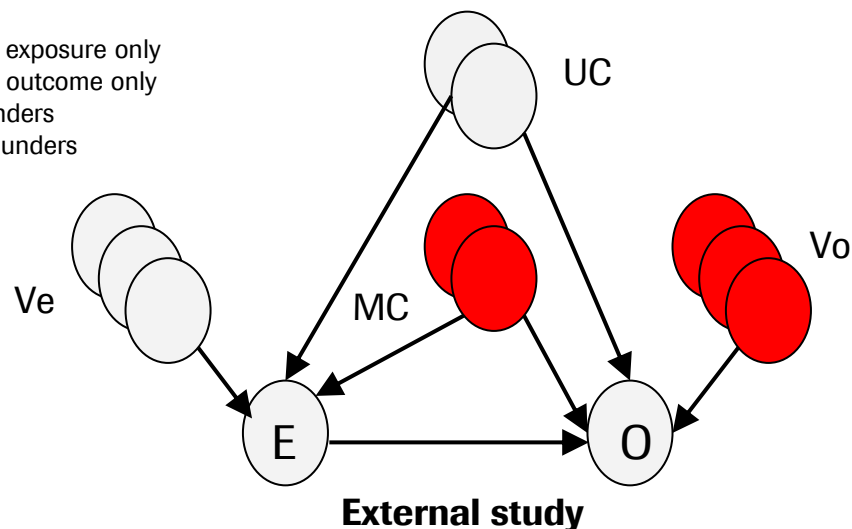
E = exposure
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 UC = unmeasured confounders



Error-prone PS_{main}

Logistic regression

$$E \sim MC + Vo$$



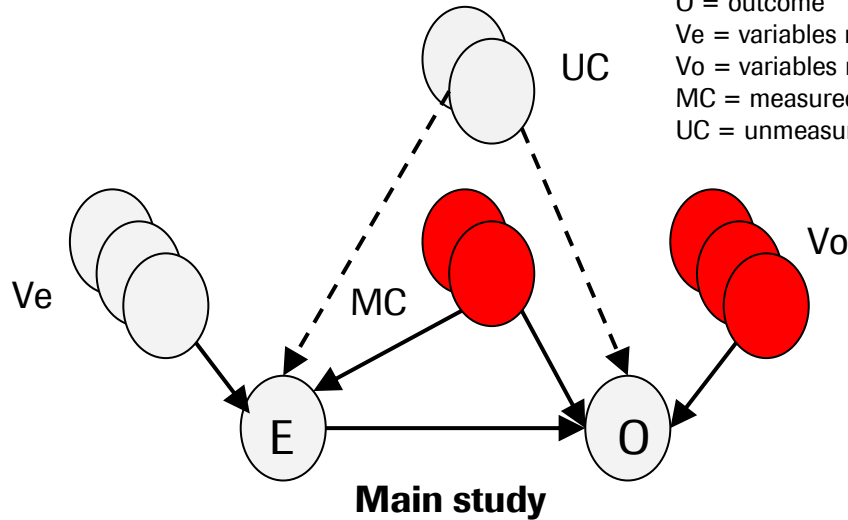
Error-prone PS_{external}

Logistic regression

$$E \sim MC + Vo$$

Frequentist Propensity Score Calibration

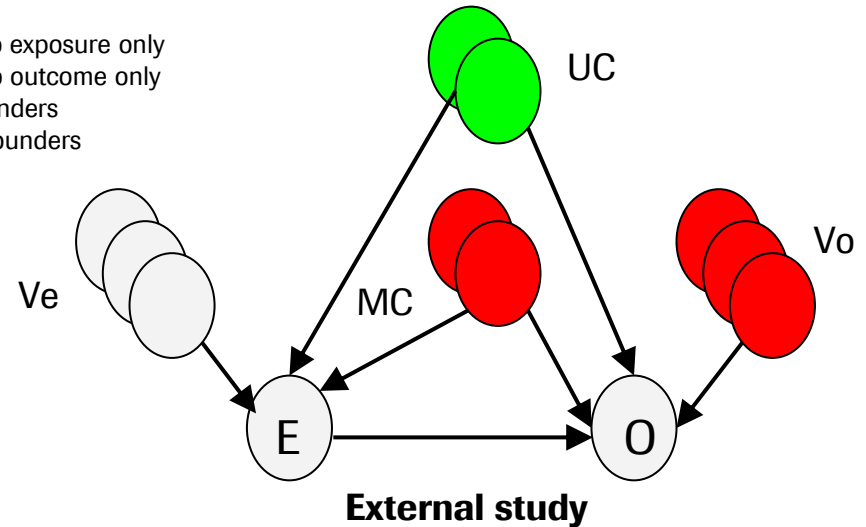
E = exposure
 O = outcome
 Ve = variables related to exposure only
 Vo = variables related to outcome only
 MC = measured confounders
 UC = unmeasured confounders



Error-prone PS_{main}

Logistic regression

$$E \sim MC + Vo$$



Error-prone PS_{external}

Logistic regression

$$E \sim MC + Vo$$

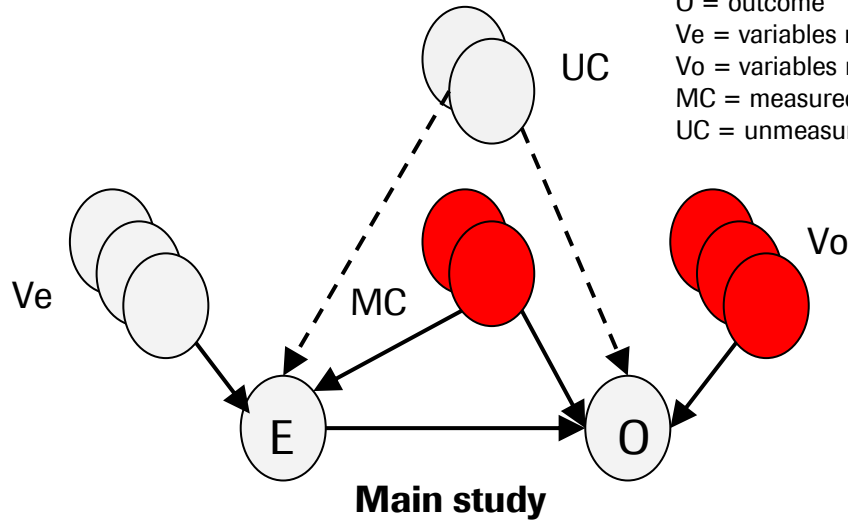
Gold standard PS_{external}

Logistic regression

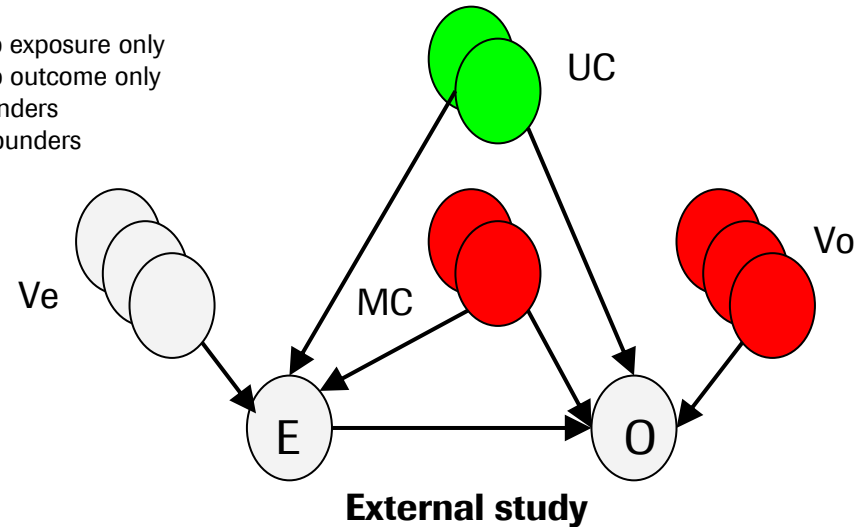
$$E \sim MC + UC + Vo$$

Frequentist Propensity Score Calibration

E = exposure
 O = outcome
 Ve = variables related to exposure only
 Vo = variables related to outcome only
 MC = measured confounders
 UC = unmeasured confounders



Error-prone PS_{main}



Error-prone PS_{external}

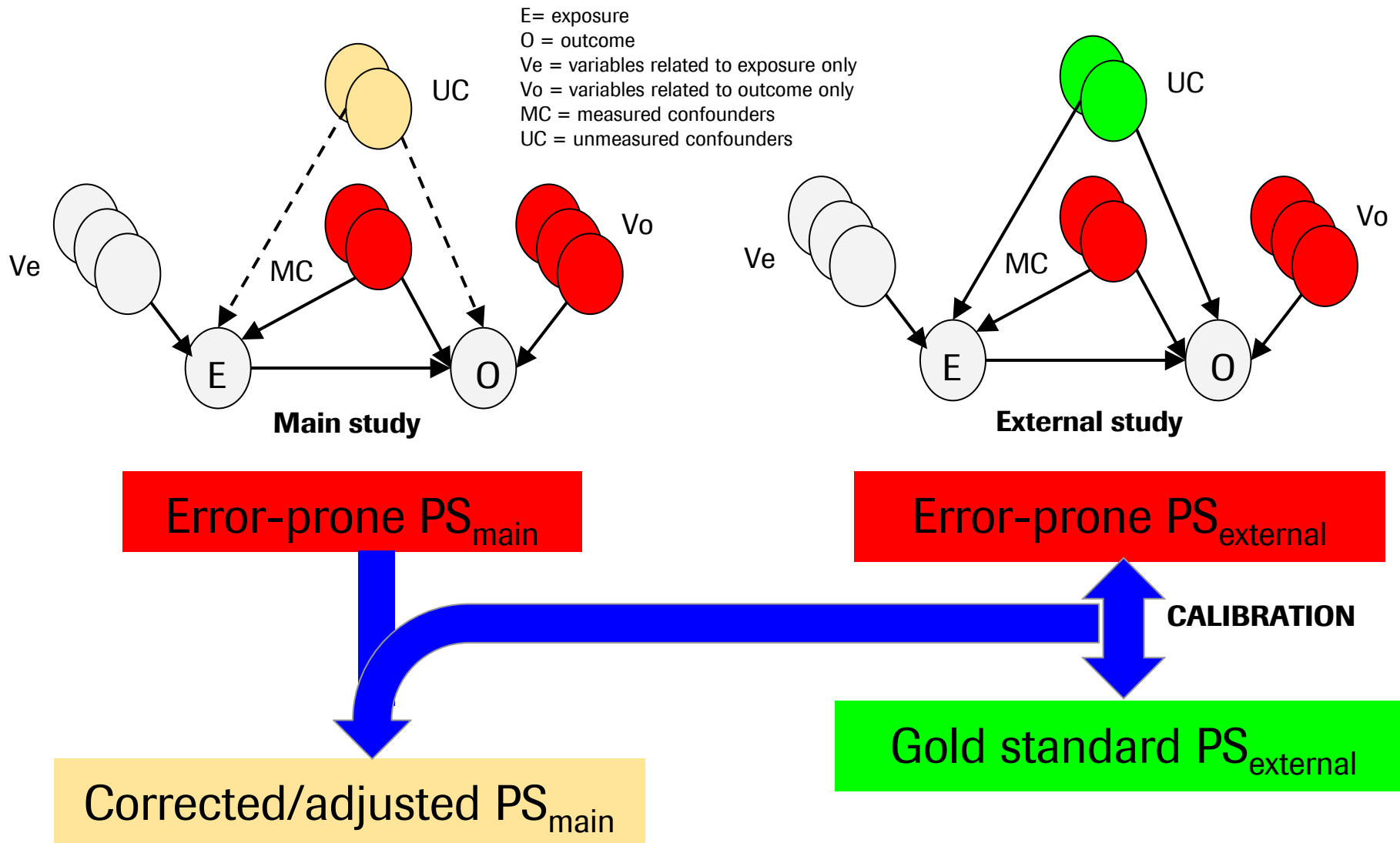
Linear regression

$$PS_{\text{GS}} \sim E + PS_{\text{EP}}$$

CALIBRATION

Gold standard PS_{external}

Frequentist Propensity Score Calibration



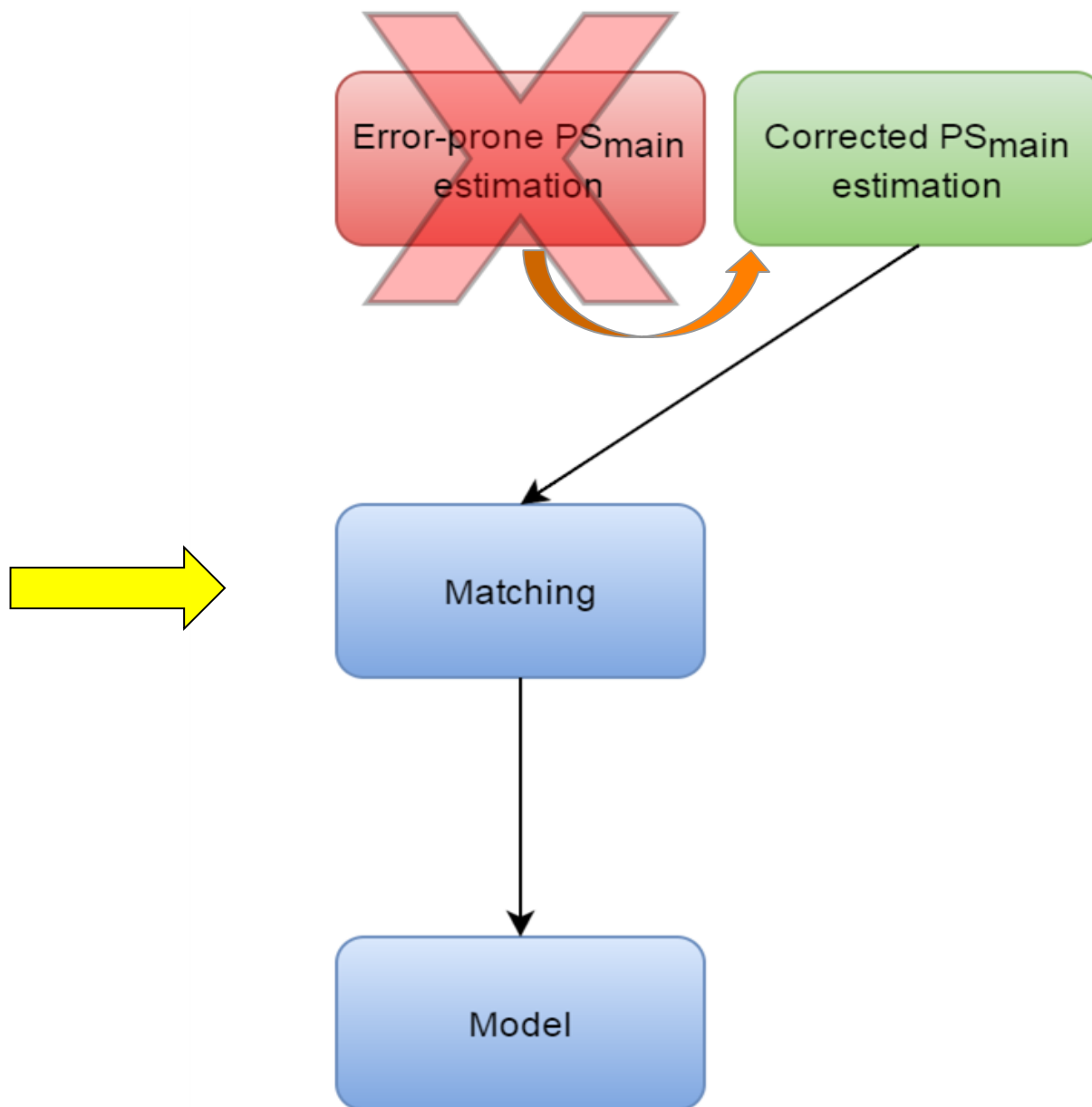
Surrogacy hypothesis

The effectiveness of the Propensity Score Calibration (PSC) strongly relies on the **surrogacy hypothesis**, requiring that

“the error-prone propensity score is independent of the outcome of interest given the gold standard PS and the exposure”.
(i.e. the error-prone PS is a **surrogate** of the gold standard PS).

O independent **PS_{EP} | PS_{GS}, E**

Unmeasured Confounding using PS

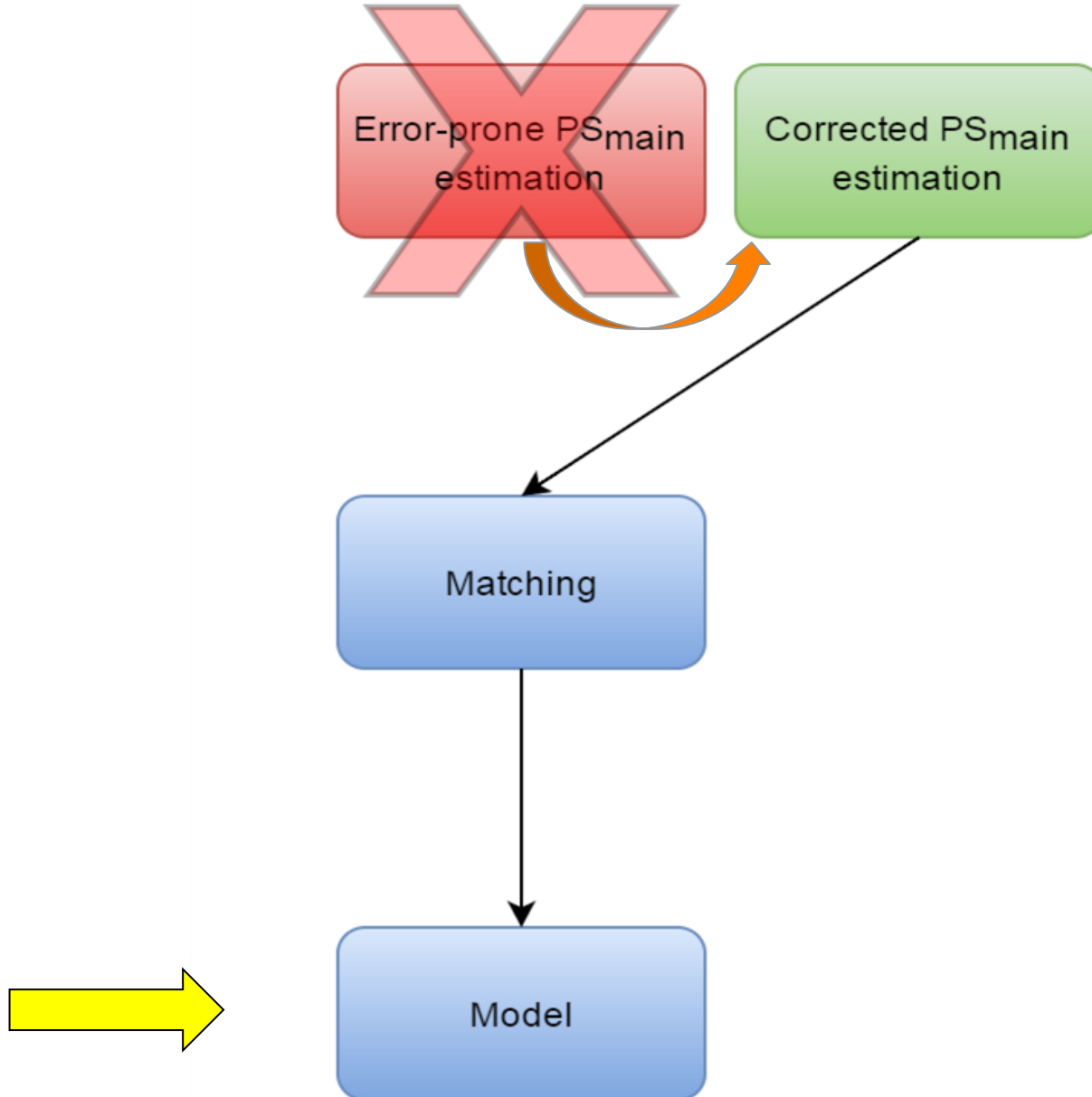


- Use of external data
- Calibration

Matching

1:1 with no replacement within a fixed caliper
by greedy algorithm

Unmeasured Confounding using PS



- Use of external data
- Calibration
- Matching in the main study population

Model

Marginal treatment effect estimated by the appropriate model on the matched population (e.g. Cox model)

$$\mathbf{O} \sim \mathbf{E}$$

O= Outcome E=Exposure

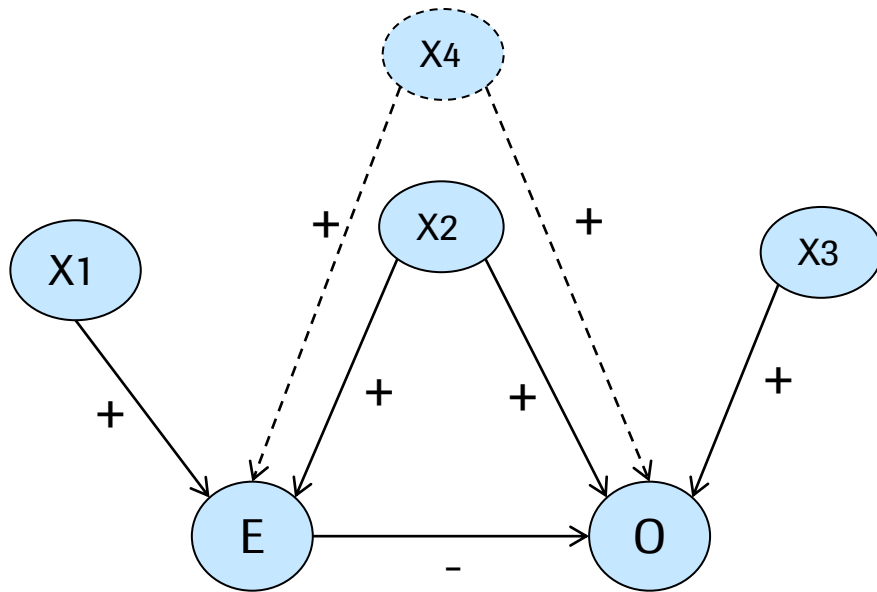
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Propensity Score Calibration

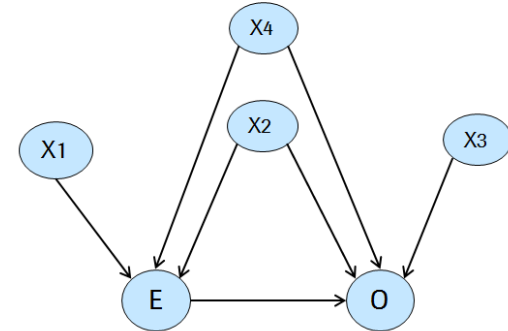
Monte Carlo simulation

Conclusions

Simulation for Unmeasured Confounding



Main study
1000 subjects



External study
250 subjects

$X_1, X_2, X_3, X_4 \sim \text{Bernoulli}(0.5)$

$\text{logit}(p) = 1.5 \cdot X_1 + 1.5 \cdot X_2 + 1.5 \cdot X_4 + \text{logit}(0.05)$

$E \sim \text{Bernoulli}(p)$

$\text{log}(r) = \text{log}(0.76) \cdot E + \text{log}(2) \cdot X_2 + \text{log}(2) \cdot X_4 + \text{log}(2) \cdot X_3 - 3.5$

$O \sim \exp(r)$

No censoring

Simulation for Unmeasured Confounding

Example of PS models and calibration

log(OR)	Error Prone PS	Gold standard PS
Intercept	-1.17 ***	-1.82 ***
X2	1.49 ***	1.47 ***
X3	-0.26	-0.21
X4	NA	1.08 ***

$p \leq 0.001$ ***

$0.01 < p \leq 0.05$ *

$p > 0.1$

$0.001 < p \leq 0.01$ **

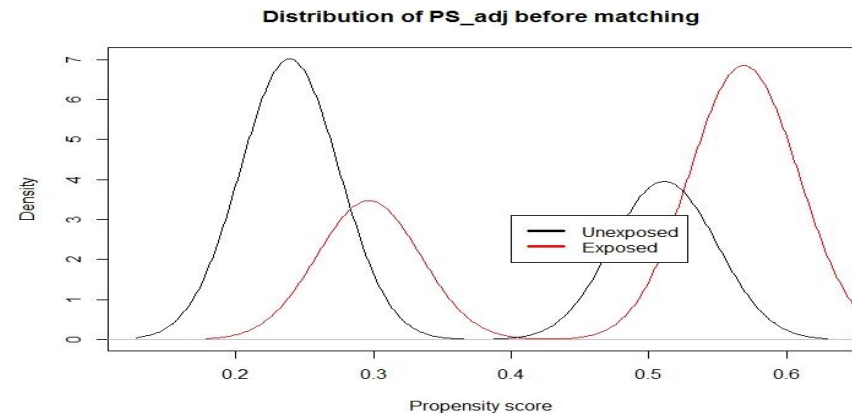
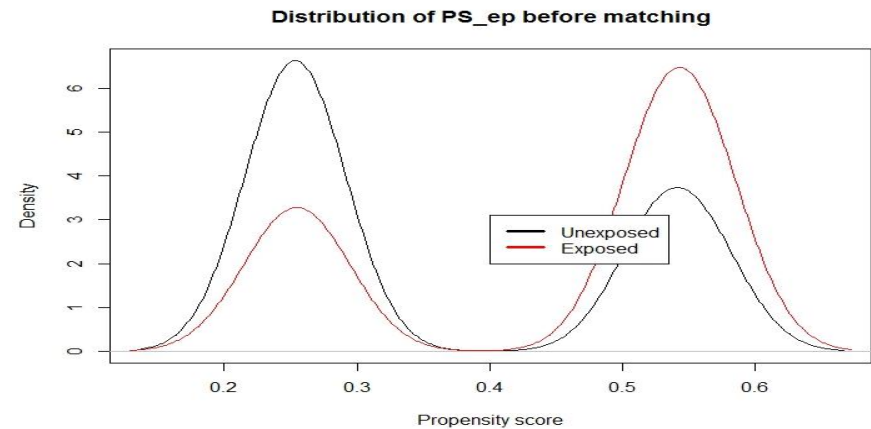
$0.05 < p \leq 0.1$.

	Calibration
Intercept	0.00
E	0.06 ***
PS ep	0.94 ***

Simulation for Unmeasured Confounding

Example of PS models and calibration

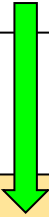
	Calibration
Intercept	0.00
E	0.06 ***
PS ep	0.94 ***



Simulation for Unmeasured Confounding

Results for 1000 repetitions

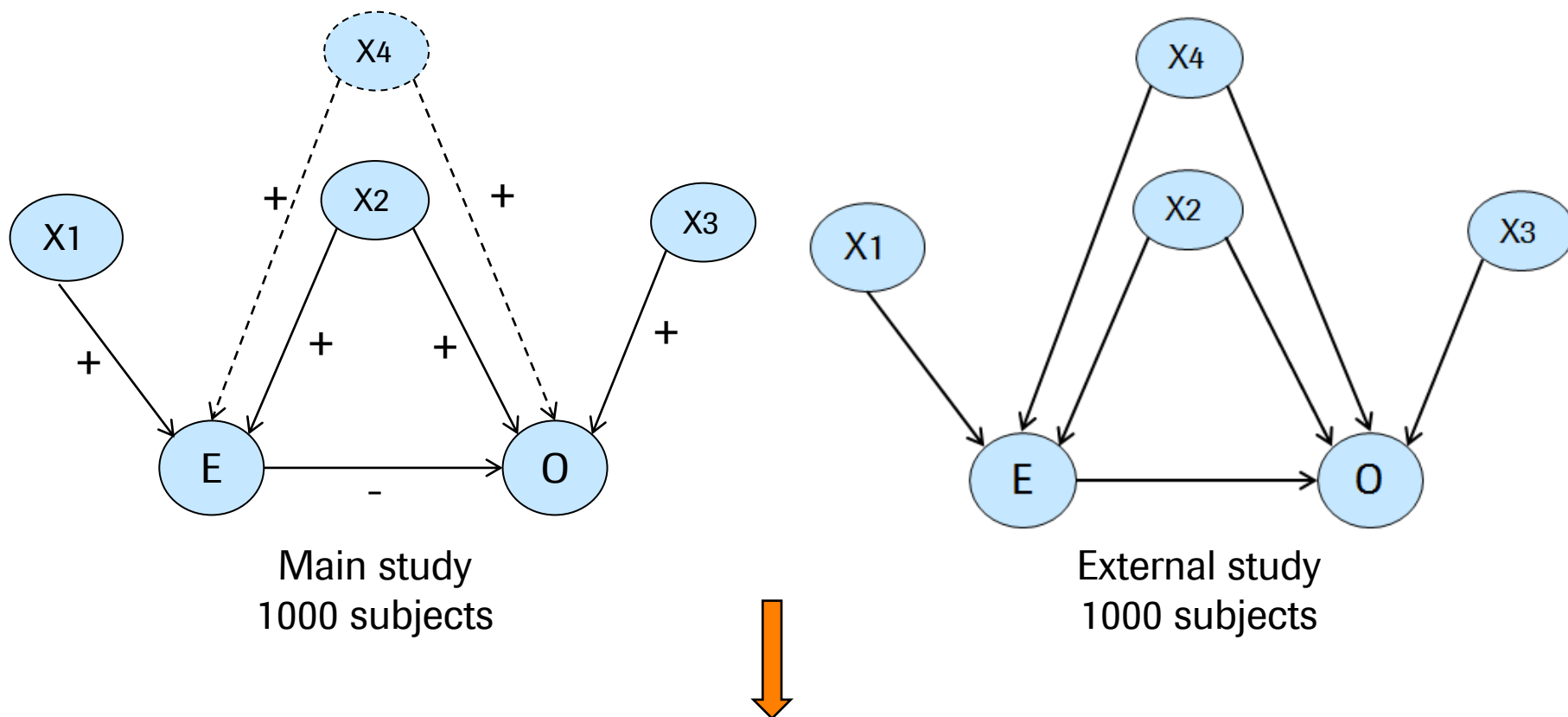
True = 0.80	Point est (HR)	95% confint	Relative Bias	Matched subjects
Crude	1.12	[0.99, 1.29]	40%	NA
Matching (EP)	0.96	[0.84, 1.10]	20%	716
Matching (adj)	0.90	[0.58, 1.18]	18%	657



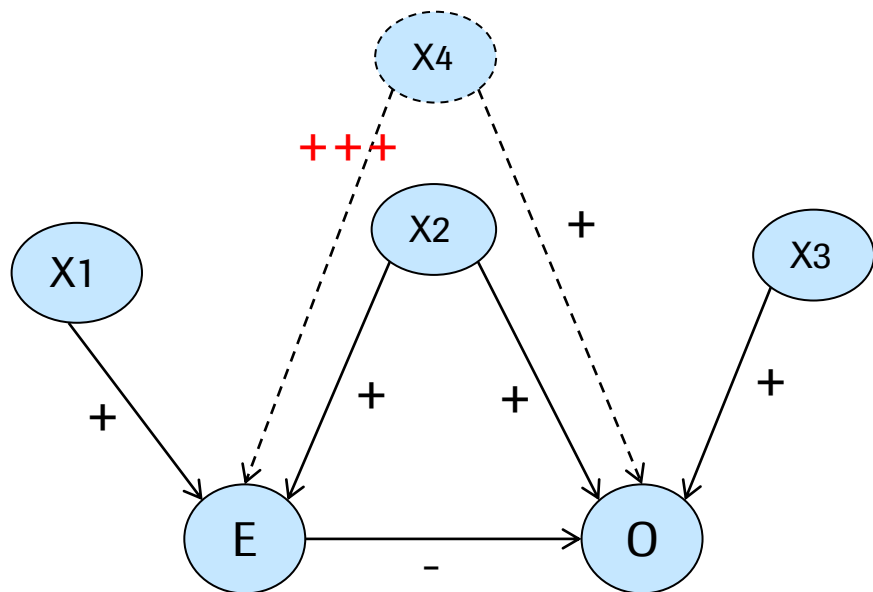
Legend

- Point est: mean Hazard Ratio (HR) across 1000 repetitions
- 95% confint: 95% empirical confidence interval using 2.5% and 97.5% quantiles across 1000 repetitions
- Relative Bias: mean relative absolute bias across 1000 repetitions
- Matched subjects: mean number of matched patients across 1000 repetitions

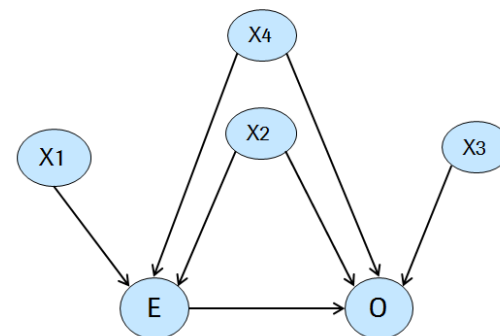
Other scenarios: larger external study



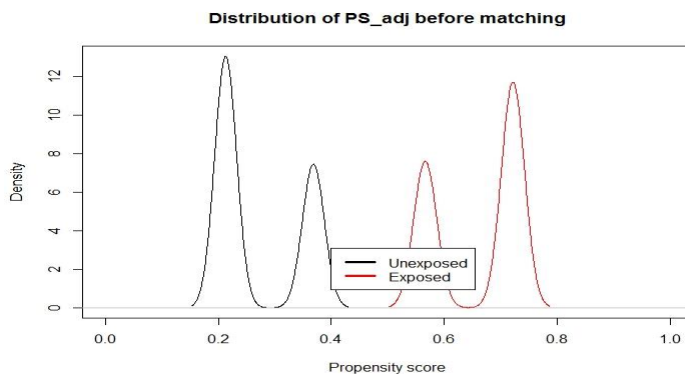
Other scenarios: predominant UC



Main study
1000 subjects

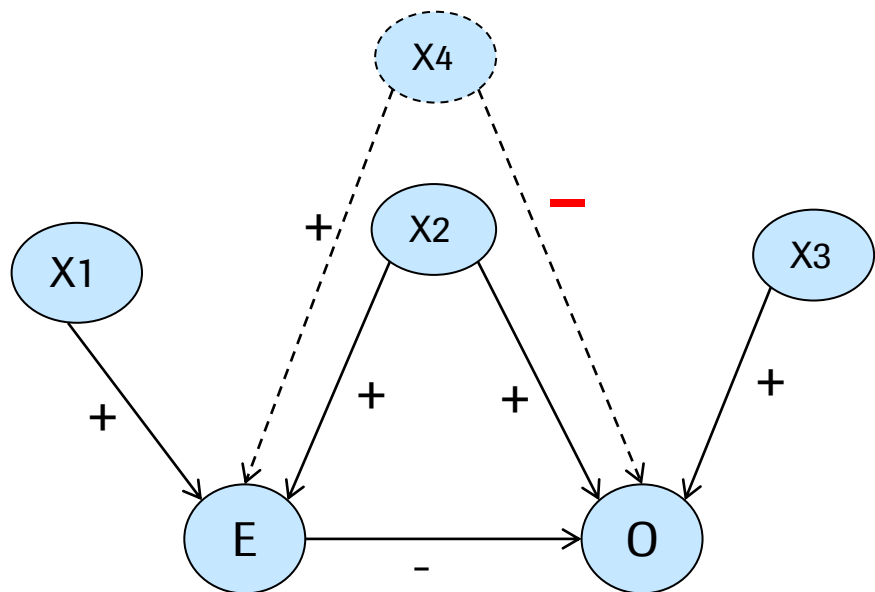


External study
250 subjects



- **No overlapping → matching problem**
- **Balance of confounders is not achieved**

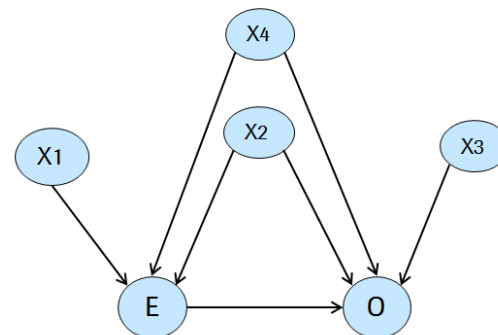
Other scenarios: surrogacy violation



Main study
1000 subjects

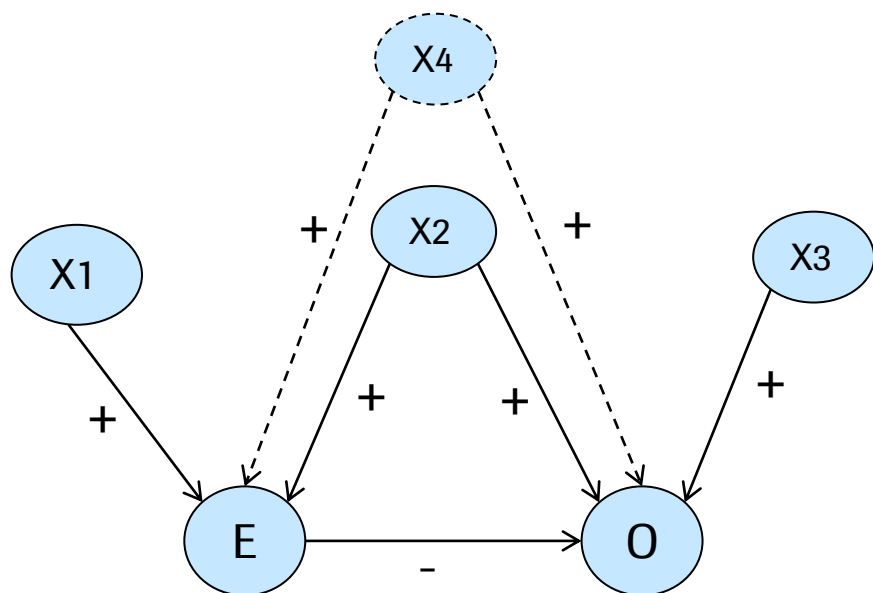


Biased estimate



External study
250 subjects

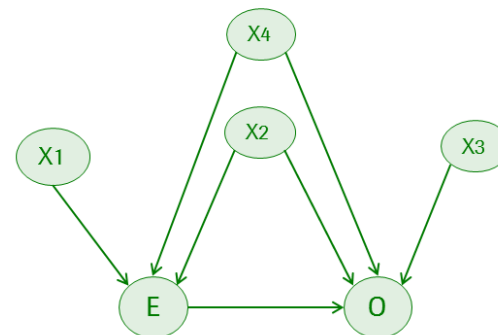
Other scenarios: transportability



Main study
1000 subjects



**PSC correction may
not be significant or
reliable**



External study
250 subjects

Perturbations in:

- **Exposure prevalence**
- **Exposure Causal Parameters**

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- ❖ The **surrogacy test** enables to define a context where PSC **reduces** confounding bias
- ❖ This allows to use PSC to perform sensitivity analyses, once the related practical elements have been properly defined



- The **matching algorithm** and the **caliper** may significantly impact on the performance of the method: different options may be undertaken
- The choice of the **external database** (and its size) plays an important role, even if this point is often related to practical feasibility constraints and it is difficult to check
- Correlations and different strengths of covariates effects on treatment and outcome require further assessment

Doing now what patients need next