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Probabilistic Graphical Models and Causal Inference

*Episode 2:
Structural Causal Models
(from Interventions to Counterfactuals)*

Marco Piastra

This presentation can be downloaded at:
<https://vision.unipv.it/AI/AIRG.html>

- **Causal Inference in Statistics**

A Primer

Judea Pearl, Madelyn Glymour and Nicholas P. Jewell

Wiley, 2016



CAUSAL INFERENCE IN STATISTICS

A Primer

Judea Pearl
Madelyn Glymour
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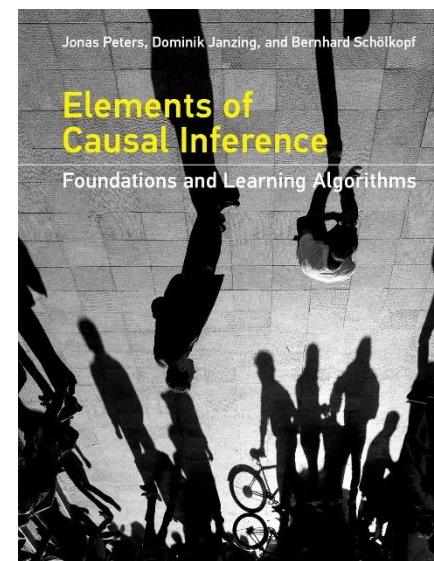
WILEY

- **Elements of Causal Inference**

Foundations and Learning Algorithms

Jonas Peters, Dominik Janzing and Bernhard Schölkopf

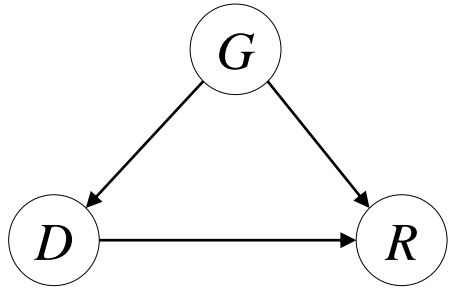
MIT Press, 2017



Causal Graphical Models and Interventions (*recap*)

Causes and Effects: *say it with graphs*

- Probabilistic Graphical Model



G is biological gender (= Male/Female)

D is drug administration (= Yes(1)/No(0))

R is recovery from illness (= Yes(1)/No(0))

$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$

<i>Females</i>	$R = 0$	$R = 1$		Recovery Rate
$D = 0$	25	55	80	69%
$D = 1$	71	192	263	73%
	96	247	343	

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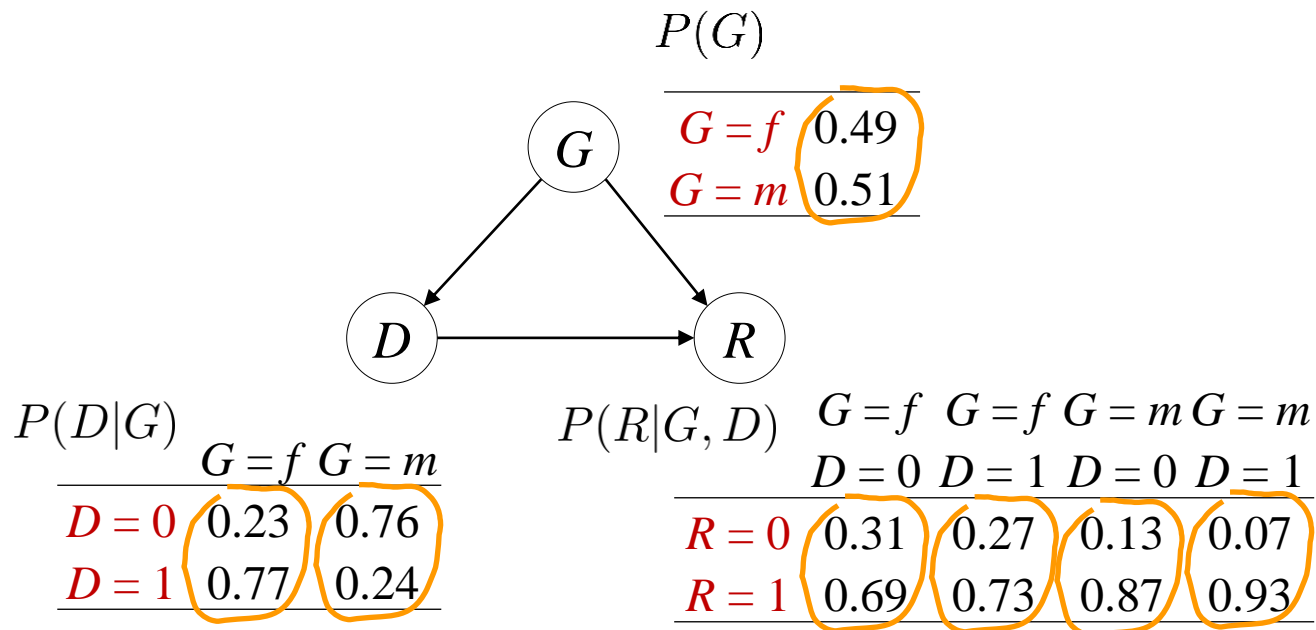
[Data from Pearl, J. et al., "Causal Inference in Statistics: A Primer", Wiley, 2016]

Causes and Effects: *say it with graphs*

■ Probabilistic Graphical Model

Maximum Likelihood Estimation (CPTs) of

$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$



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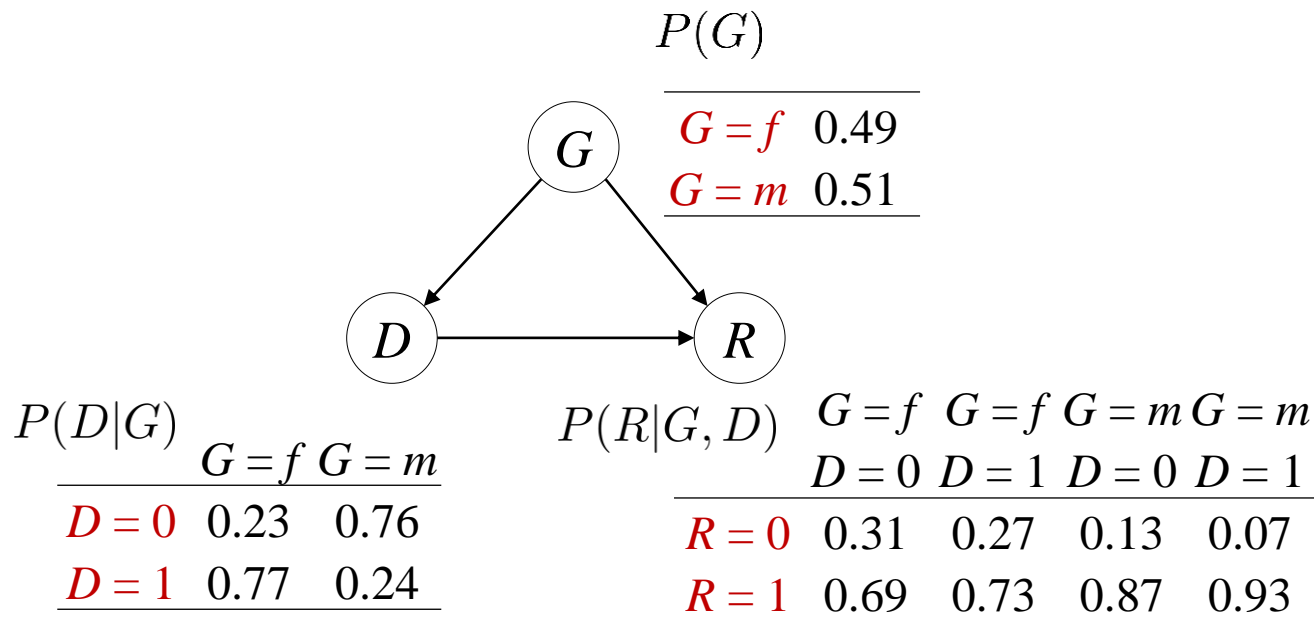
[Data from Pearl, J. et al., "Causal Inference in Statistics: A Primer", Wiley, 2016]

Causes and Effects: *say it with graphs*

■ Probabilistic Graphical Model

Maximum Likelihood Estimation (CPTs) of

$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$



Using Graphical Model as a predictor

Case 1: Gender is observed

$$P(R = 1 | G = 0, D = 0) = 0.69$$

$$P(R = 1 | G = 0, D = 1) = 0.73$$

$$P(R = 1 | G = 1, D = 0) = 0.87$$

$$P(R = 1 | G = 1, D = 1) = 0.93$$

Prescribe drug, regardless

Case 2: Gender is not observed

$$P(R|D) = \frac{\sum_G P(R|G, D)P(D|G)P(G)}{\sum_{G,R} P(R|G, D)P(D|G)P(G)}$$

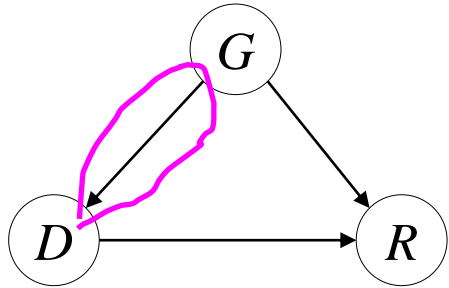
$$P(R = 1 | D = 0) = 0.83$$

$$P(R = 1 | D = 1) = 0.78$$

Do not prescribe drug, regardless
(ridiculous!)

Causes and Effects: *say it with graphs*

■ Probabilistic Graphical Model



G is biological gender (= Male/Female)

D is drug administration (= Yes(1)/No(0))

R is recovery from illness (= Yes(1)/No(0))

How can we solve the problem?

- The problem is due to the discrepancy in drug administration across genders
- An obvious solution would be *to repeat* the experiment with equal administration rates
- *In other words, we would sever **this** link*

<i>Females</i>	$R = 0$	$R = 1$		Recovery Rate
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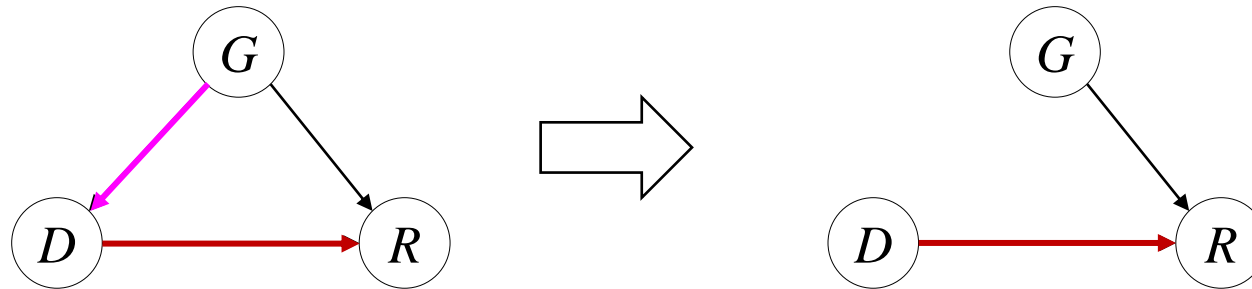
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The Magic of Controlled Experiments

■ When association is causation



In this *Causal Graphical Model*:

1. The **causal effect** we are interested in is that of D over R
2. The **link** between G and D is *problematic*: we know that $P(D|G = 0) \neq P(D|G = 1)$
3. In a *controlled experiment*, D is administered at random, therefore

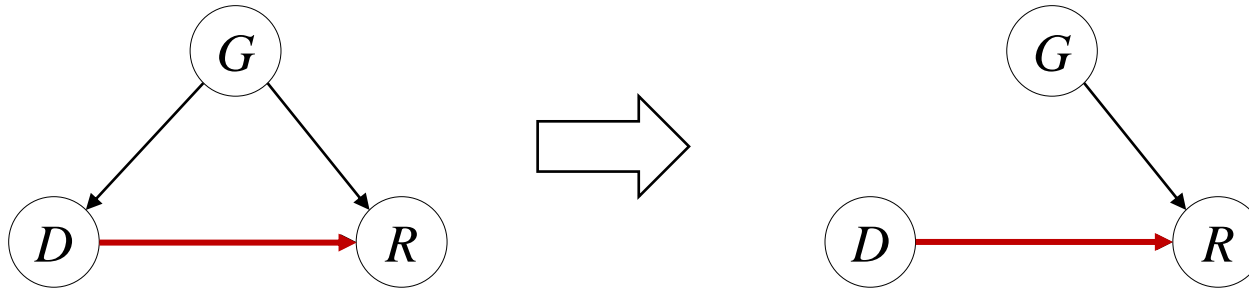
$$\langle D \perp G \rangle \implies P(D|G = 0) = P(D|G = 1) = P(D)$$

4. In other words, the corresponding CGM 'loses' the problematic **link** and the estimate becomes

$$P(R|D) := \sum_G P(G)P(R|G, D)$$

The Magic of Controlled Experiments

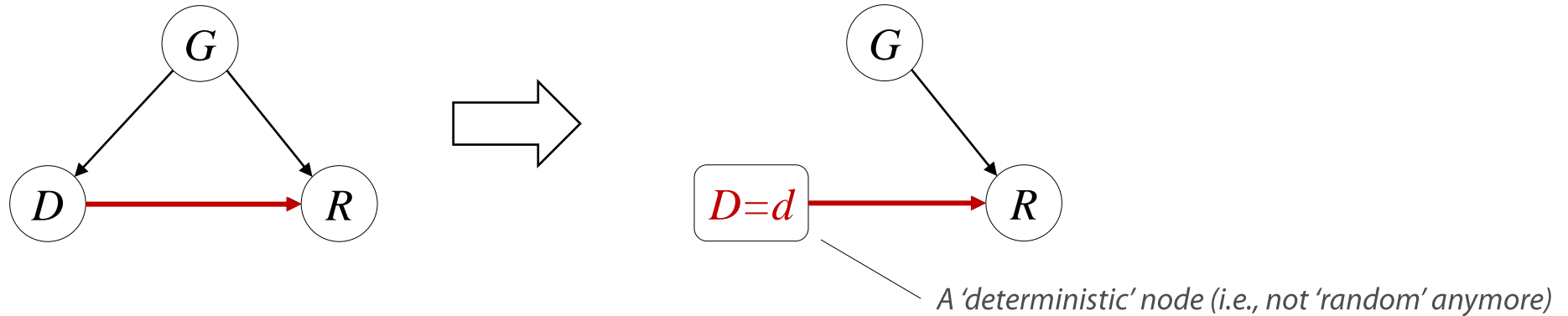
- **When association is causation**



With *controlled experiments* (i.e., the ‘gold standard’ for testing) the principle is more general:

- by *randomizing* the administration of treatment
- we make the *effects* independent of any *confounders* (be them observed or not)

■ From Conditional (pre-intervention) to Intervention Probability



In this *Causal Graphical Model* (for an uncontrolled experiment):

1. Conditional probability:

$$P(R|D = d) = \frac{\sum_G P(G)P(R|G, D = d)P(D = d|G)}{\sum_G P(G)P(D = d|G)}$$

2. Intervention (**do-calculus**, *this is new*)

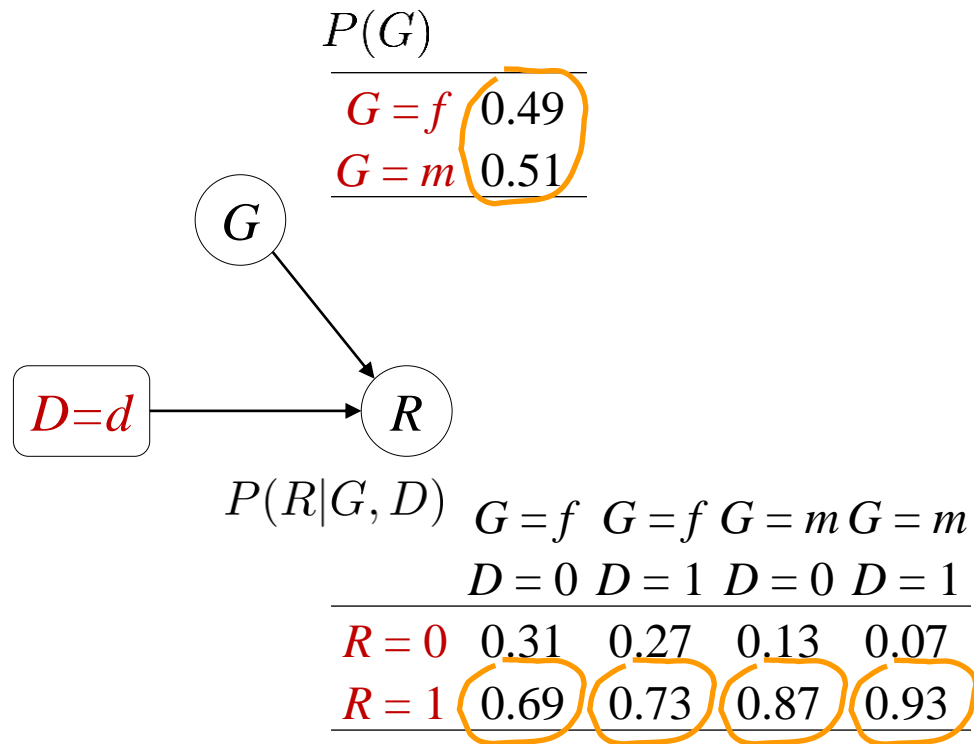
$$P(R|do(D = d)) := \sum_G P(G)P(R|G, D = d)$$

3. This is equivalent to $P(R|D = d)$ in a modified CGM in which we 'enforce intervention'

These two expressions would be identical if
 $P(D = d|G) = 1$
which cannot be true in general

From Conditional (pre-intervention) to Intervention Probability

(same observational probabilities, from MLE)



Using do-calculus

$$P(R = 1 | do(D = 0)) = \sum_G P(G) P(R = 1 | G, D = 0)$$

$$= 0.49 \cdot 0.69 + 0.51 \cdot 0.87 = 0.78$$

$$P(R = 1 | do(D = 1)) = \sum_G P(G) P(R = 1 | G, D = 1)$$

$$= 0.49 \cdot 0.73 + 0.51 \cdot 0.93 = 0.83$$

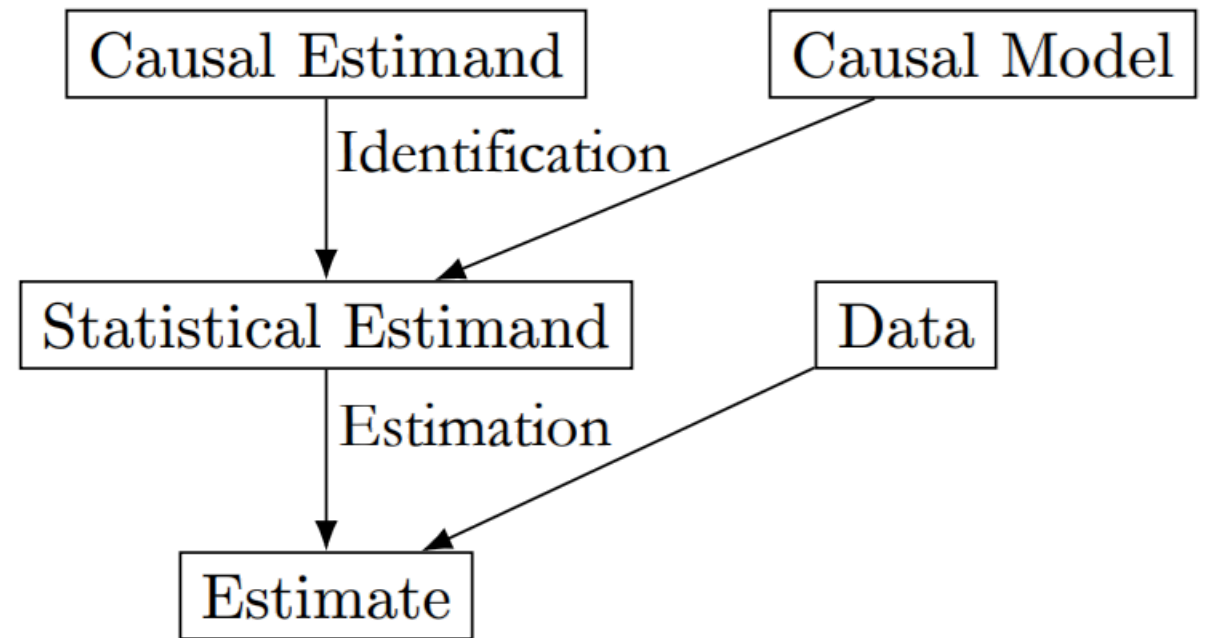
Prescribe drug, regardless

Causation and Conditionals

■ Causal Model and Estimation

Basic principles:

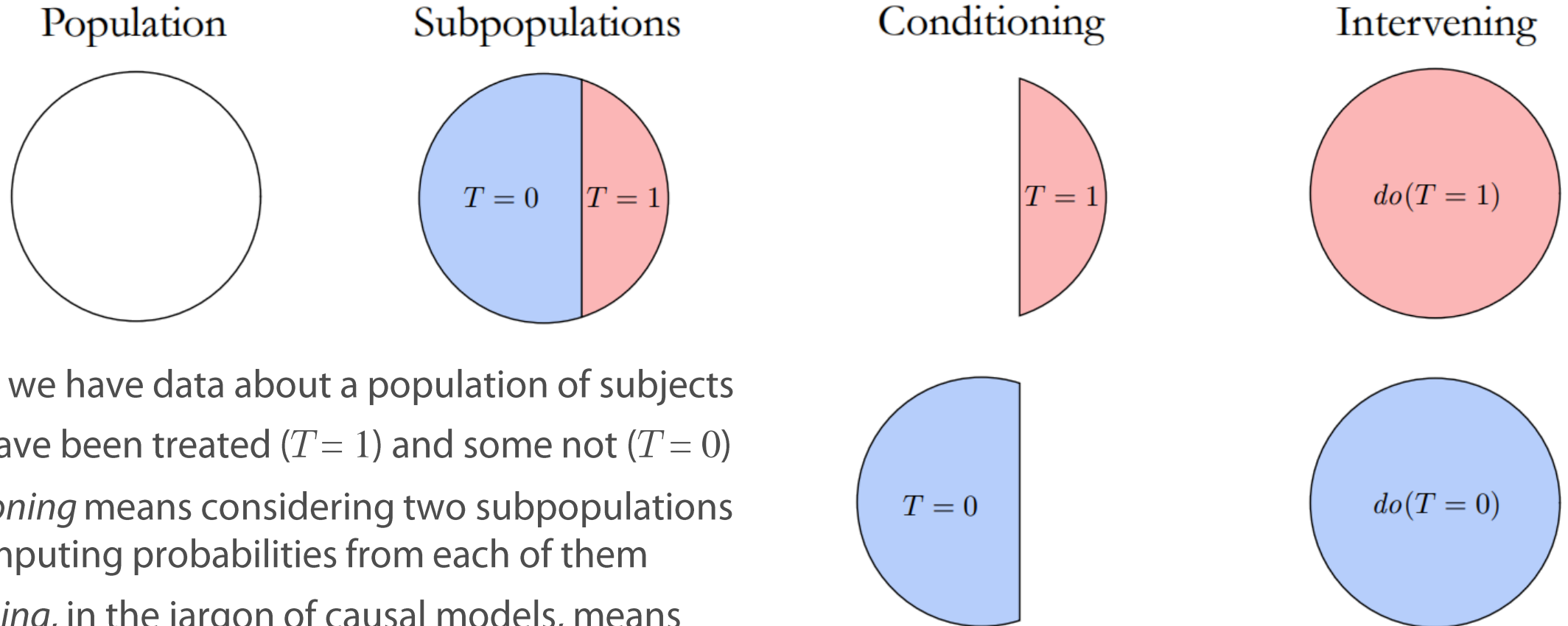
1. Having selected what kind of causal effect we want to estimate
2. We start from a *Causal Graphical Model* (CGM)
3. To translate the estimate into a **statistical estimand**, (*Identification*)
4. We use then *observational* data to compute the **estimate**: a *probability* or an *expected value*



[Image from <https://www.bradyneal.com/causal-inference-course>]

Causation and Conditionals

■ Conditioning and Intervening



Assume we have data about a population of subjects
Some have been treated ($T = 1$) and some not ($T = 0$)

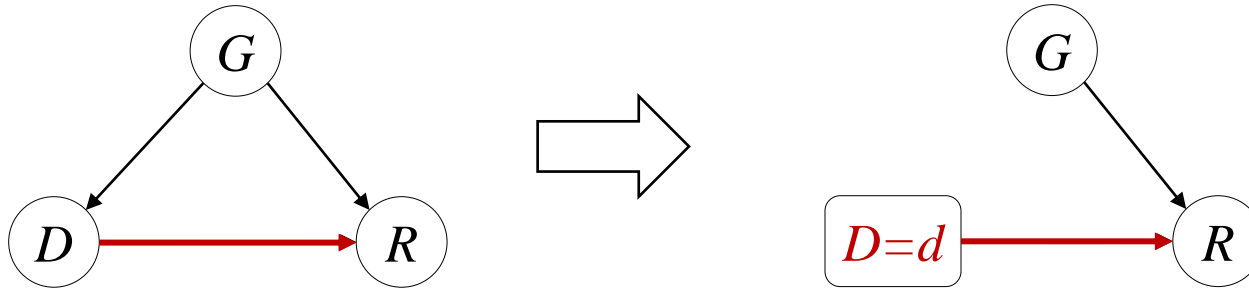
Conditioning means considering two subpopulations
and computing probabilities from each of them

Intervening, in the jargon of causal models, means
assuming that every subject in the population has
been treated or not (*potential outcomes*)

[Image from <https://www.bradyneal.com/causal-inference-course>]

do-Calculus

Compare two expressions



1. Conditioning:

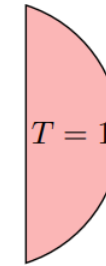
$$P(R|D = d) = \frac{\sum_G P(G)P(R|G, D = d)P(D = d|G)}{\sum_G P(G)P(D = d|G)}$$

2. Intervening:

$$P(R|do(D = d)) := \sum_G P(G)P(R|G, D = d)$$

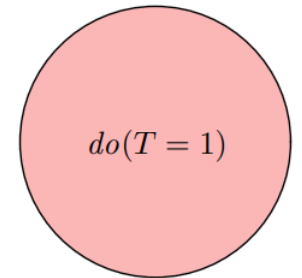
no normalization =
no conditional subspace

Conditioning

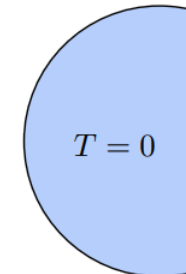


$T = 1$

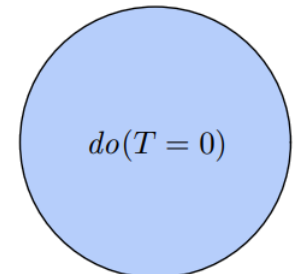
Intervening



$do(T = 1)$



$T = 0$



$do(T = 0)$

Identification

■ Adjustment Set Criterion [Shipster et al. 2010]

In a Causal Graphical Model, the *causal effect* of T over Y is *identifiable* iff it exists an *adjustment set* \mathbf{W} of variables such that:

- no variable in \mathbf{W} is on, or is a descendant of any variables on, a **causal path** (excluding the descendants of T alone)
- the variables in \mathbf{W} block (*in the sense of graphical models*) all the **non-causal paths** between T and Y

This criterion is necessary and sufficient for effect identifiability

Then:

$$P(Y|do(T = t)) = \sum_{\mathbf{W}} P(Y|T = t, \mathbf{W})P(\mathbf{W})$$

In words, the causal effect can be estimated statistically, from data

Identification

- **Adjustment Set Criterion with *observed* and *unobserved* variables**

*More in general, in practical cases,
there can be both observed and unobserved (possibly hidden) variables*

An *adjustment set* can be composed of both:

$$\mathbf{W} = \mathbf{W}_{obs} \cup \mathbf{W}_{hid}$$

Then, if \mathbf{W} satisfies the Adjustment Set Criterion:

$$P(Y|do(T = t), \mathbf{W}_{obs}) = \sum_{\mathbf{W}_{hid}} P(Y|T = t, \mathbf{W}_{hid}, \mathbf{W}_{obs})P(\mathbf{W}_{hid})$$

When there are no *observed* variables in the adjustment set:

$$P(Y|do(T = t)) = \sum_{\mathbf{W}} P(Y|T = t, \mathbf{W})P(\mathbf{W})$$

Likewise, when there are no *unobserved* variables in the adjustment set:

$$P(Y|do(T = t), \mathbf{W}) = P(Y|T = t, \mathbf{W})$$

Estimating Effects

Expected effects of different interventions can be estimated via do-calculus

In general, the *expected effect* on Y of treatment T will be

$$\mathbb{E}[Y|T = t, \mathbf{W}_{obs}] := \sum_{y \in \mathcal{Y}} y P(Y|do(T = t), \mathbf{W}_{obs})$$

where $\mathbf{W} = \mathbf{W}_{obs} \cup \mathbf{W}_{hid}$ is a valid *adjustment set*

Differences in effects can be measured by comparing expected effects.

As a special case, when $T \in \{0, 1\}$

- The *Conditional Average Treatment Effect* (CATE) is defined as:

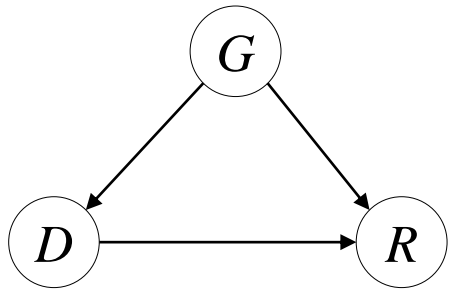
$$\tau(\mathbf{W}_{obs}) := \mathbb{E}[Y|T = 1, \mathbf{W}_{obs}] - \mathbb{E}[Y|T = 0, \mathbf{W}_{obs}]$$

- The *Average Treatment Effect* (ATE) is defined as:

$$\mathbb{E}[\tau(\mathbf{W})] := \mathbb{E}[Y|T = 1] - \mathbb{E}[Y|T = 0]$$

Structural Causal Models

Probabilistic Graphical Model



G is biological gender (= Male/Female)

D is drug administration (= Yes(1)/No(0))

R is recovery from illness (= Yes(1)/No(0))

$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$

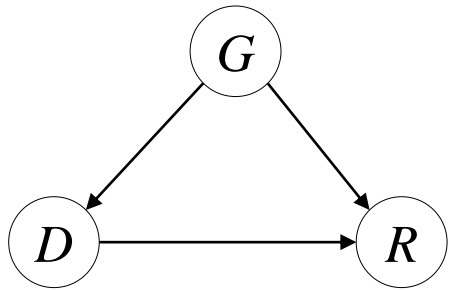
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[Data from Pearl, J. et al., "Causal Inference in Statistics: A Primer", Wiley, 2016]

From Graphical Model to Structural Equations



G is biological gender (= Male/Female)
 D is drug administration (= Yes(1)/No(0))
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$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$

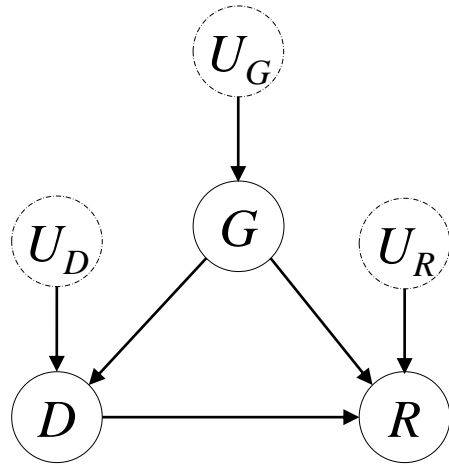
Structural Equations * first approximation

$$D = f_D(G)$$

$$R = f_R(G, D)$$

How can these two things
be reconciled?
Functions are deterministic

From Graphical Model to Structural Equations



G is biological gender (= Male/Female)

D is drug administration (= Yes(1)/No(0))

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$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$

U_G , U_D and U_R are unobservable, random variables

The probability distribution is the observable aspect of the structural equations

Structural Equations ^{* second approximation}

$$G = U_G$$

$$D = f_D(G, U_D)$$

$$R = f_R(G, D, U_R)$$

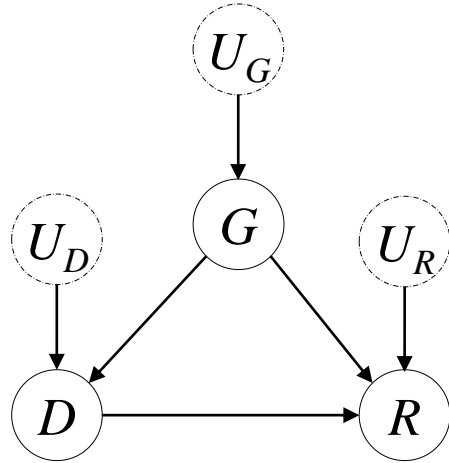
Causal? Functions could be invertible

Example:

$$D = k + \beta_g G + U_D$$

$$G = \frac{1}{\beta_g} (D - k - U_D)$$

From Graphical Model to Structural Causal Model



G is biological gender (= Male/Female)

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$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$

U_G , U_D and U_R are unobservable, random variables

The probability distribution is the observable aspect of the structural causal model

Structural Causal Model (SCM)

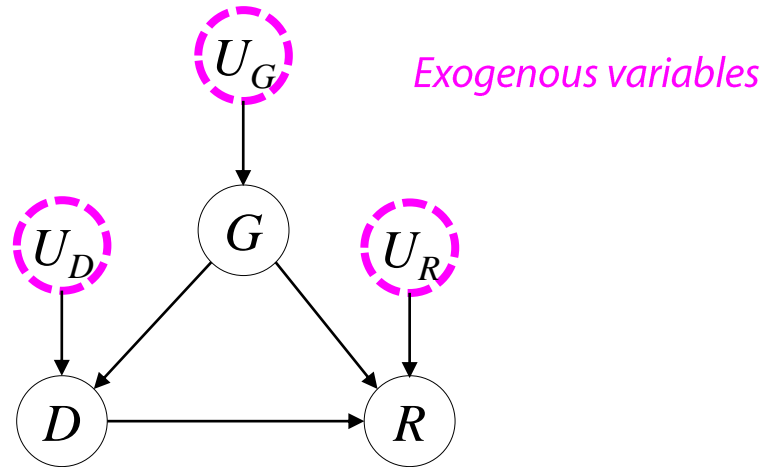
$$G := U_G$$

$$D := f_D(G, U_D)$$

$$R := f_R(G, D, U_R)$$

Force directions, in keeping with causation assumptions

From Graphical Model to Structural Causal Model



Structural Causal Model (SCM)

$$G := U_G$$

$$D := f_D(G, U_D)$$

$$R := f_R(G, D, U_R)$$

G is biological gender (= Male/Female)

D is drug administration (= Yes(1)/No(0))

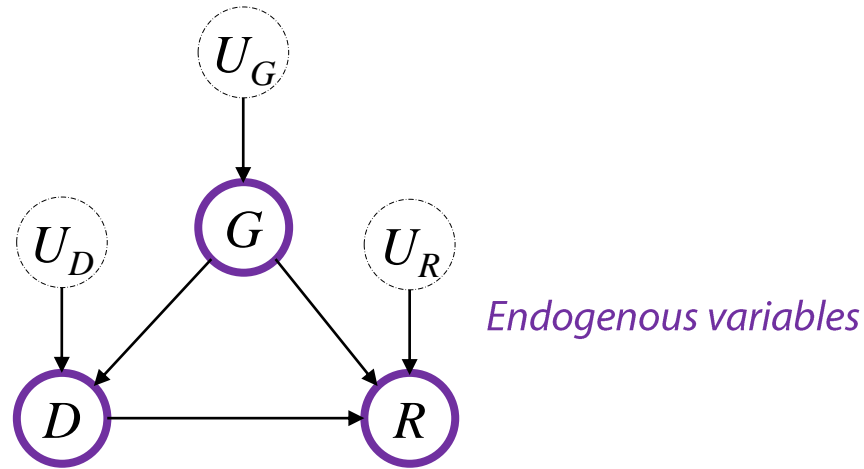
R is recovery from illness (= Yes(1)/No(0))

$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$

U_G , U_D and U_R are unobservable, random variables

The probability distribution is the observable aspect of the structural causal model

From Graphical Model to Structural Causal Model



G is biological gender (= Male/Female)

D is drug administration (= Yes(1)/No(0))

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$$P(G, R, D) = P(G)P(D|G)P(R|G, D)$$

U_G , U_D and U_R are unobservable, random variables

The probability distribution is the observable aspect of the structural causal model

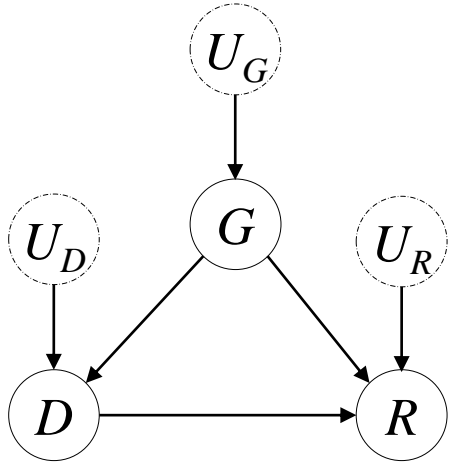
Structural Causal Model (SCM)

$$G := U_G$$

$$D := f_D(G, U_D)$$

$$R := f_R(G, D, U_R)$$

Structural Causal Model



Structural Causal Model (SCM) definition

- 1) A set of *endogenous* variables
- 2) A set of *exogenous* variables
- 3) A set of *structural equations*

An SCM *induces* a graphical model with a probability distribution P over *endogenous* variables

Structural Causal Model (SCM)

$$G := U_G$$

$$D := f_D(G, U_D)$$

$$R := f_R(G, D, U_R)$$

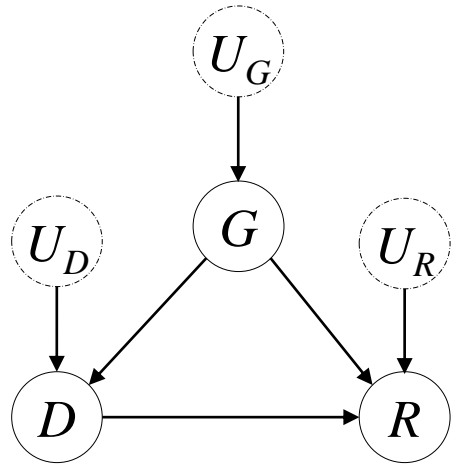
Structural Causal Model (formal definition)

Structural Causal Model (SCM), formally

- 1) A set of *endogenous* variables $\mathbf{V} := \{V_1, V_2, \dots, V_n\}$
- 2) A set of *exogenous* variables $\mathbf{U} := \{U_1, U_2, \dots, U_n\}$
- 3) A set of *structural equations* $\mathbf{f} := \{f_1, f_2, \dots, f_n\}$

An SCM \mathcal{M} induces a graphical model \mathcal{G} with a probability distribution $P(\mathbf{V})$

Structural Causal Model



Structural Causal Model (SCM)

$$G := U_G$$

$$D := f_D(G, U_D)$$

$$R := f_R(G, D, U_R)$$

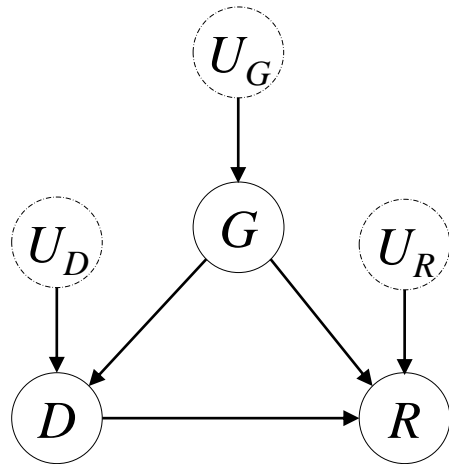
induces

The graphical model induced is uniquely defined

Further questions:

- 1) Which functions?
- 2) How are the random variables U_G , U_D and U_R distributed?
- 3) Are they dependent (or correlated)?
- 4) Is the SCM identifiable from observed data?

Structural Equation Model (linear functions)



a special case, linear

Structural Equation Model (SEM)

$$G := U_G$$

$$D := k_1 + \beta_1 G + U_D$$

$$R := k_2 + \beta_2 G + \beta_3 D + U_R$$

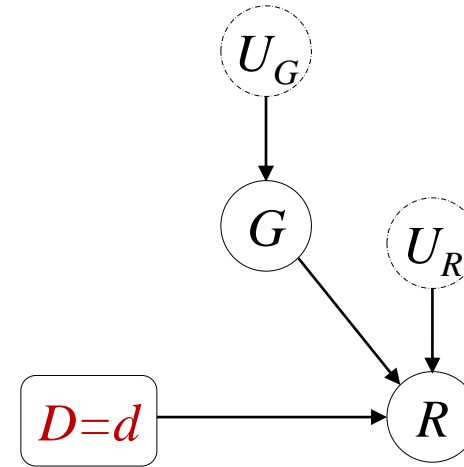
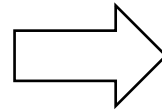
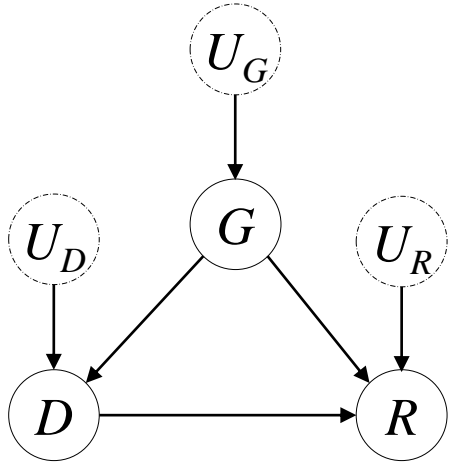
Assumptions:

- 1) *All functions are linear*
- 2) *All random variables U_G , U_D and U_R are normally distributed*
- 3) *All random variables are uncorrelated*

Under further, specific conditions a SEM is identifiable from observed data

more in general, however, this is not true of any SCM

Intervention in a Structural Causal Model

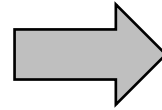


Structural Equation Model (SEM)

$$G := U_G$$

$$D := f_D(G, U_D)$$

$$R := f_R(G, D, U_R)$$



$$G := U_G$$

$$D := d$$

$$R := f_R(G, D, U_R)$$

*An intervention on an SCM creates a new sub-model by changing one or more structural equations
It induces a new graph*

Counterfactuals

Counterfactuals?

The ladder of causal inference [J. Pearl, *Causation*, Cambridge University Press, 2009]

▪ Prediction

Given the probability distribution $P(\mathbf{V})$ and some observations $\mathbf{V}_o = \mathbf{v}_o$
determine the probability $P(\mathbf{V}_u = \mathbf{v}_u \mid \mathbf{V}_o = \mathbf{v}_o)$ for some unobserved variables \mathbf{V}_u

▪ Intervention

Intervene (i.e., force a change in value) on some variables \mathbf{V}_i and determine
the probability of effects $P(\mathbf{V}_e = \mathbf{v}_e \mid do(\mathbf{V}_i = \mathbf{v}_i))$

▪ Counterfactual

Having observed $\mathbf{V}_o = \mathbf{v}_o$ and its effects $\mathbf{V}_e = \mathbf{v}_e$, what could be the probability
of different effects $\mathbf{v}'_e \neq \mathbf{v}_e$ if some conditions $\mathbf{V}_c \subseteq \mathbf{V}_o$ were different?

Counterfactual Inference

■ Counterfactual

Having observed $V_o = v_o$ and its effects $V_e = v_e$, what could be the probability of different effects $v'_e \neq v_e$ if some conditions $V_c \subseteq V_o$ were different?

A few relevant aspects:

- *Prediction and Intervention occur in the same world, whereas counterfactuals require alternative worlds*
- *Conceptually, counterfactuals relate to potential outcomes (“what could it be the outcome, were the condition different?”)*
- *Counterfactual inference can be performed at either individual or population level (more to follow)*

Counterfactual Inference

■ General Method

A few relevant aspects:

- *Prediction and Intervention occur in the same world, whereas counterfactuals require alternative worlds*
- *Conceptually, counterfactuals relate to potential outcomes (“what could it be the outcome, were the condition different?”)*
- *Counterfactual inference can be performed at either individual or population level (more to follow)*

Counterfactual Inference, deterministic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to determine the values u of the unobservable variables U
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for V_c with $V_c = v_c$ (counterfactual values) *all remaining observed values*
3. Prediction: use the sub-model to compute v_e (the effects) by using u , v_c and $v_o \setminus v_c$

Counterfactual Inference, deterministic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to determine the values u of the unobservable variables U
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for V_c with $V_c = v_c$ (counterfactual values)
3. Prediction: use the sub-model to compute v_e (the effects) by using u , v_c and $v_o \setminus v_c$

Example, in the linear case:

$$G := U_G$$

$$D := k_1 + \beta_1 G + U_D$$

$$R := k_2 + \beta_2 G + \beta_3 D + U_R$$

Assume all parameters are known (complete identification of \mathcal{M})

Counterfactual Inference, deterministic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to determine the values u of the unobservable variables U
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for V_c with $V_c = v_c$ (counterfactual values)
3. Prediction: use the sub-model to compute v_e (the effects) by using u , v_c and $v_o \setminus v_c$

Example, in the linear case:

$$u_G = g_o$$

$$u_D := d_o - k_1 - \beta_1 g_o$$

$$u_R := r_o - k_2 - \beta_2 g_o + \beta_3 d_o$$

Replace with observed values and solve for U

Counterfactual Inference, deterministic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to determine the values u of the unobservable variables U
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for V_c with $V_c = v_c$ (counterfactual values)
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Example, in the linear case:

$$g = u_G$$

$$d = \boxed{d_c} \text{ counterfactual value}$$

$$\text{effect } \boxed{r_e} = k_2 + \beta_2 g_o + \beta_3 d_c + u_R$$

Plug back values u , impose counterfactual value d_c and compute the resulting effect r_e

Counterfactual Inference, probabilistic case

From J. Pearl, Primer, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to determine the values u of the unobservable variables U
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for V_c with $V_c = v_c$ (counterfactual values)
3. Prediction: use the sub-model to compute v_e (the effects) by using u , v_c and $v_o \setminus v_c$

More in general, even keeping the assumption of complete identification of \mathcal{M} , what happens if some functions are not one-to-one for the values of U ?

Counterfactual Inference, probabilistic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to determine the values u of the unobservable variables U
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for V_c with $V_c = v_c$ (counterfactual values)
3. Prediction: use the sub-model to compute v_e (the effects) by using u , v_c and $v_o \setminus v_c$

Example, non-invertible case:

$$G := U_G$$

$$D := f_D(G, U_D)$$

$$R := f_R(G, D, U_R)$$

\mathcal{M} is still completely identified

Counterfactual Inference, probabilistic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to determine the values u of the unobservable variables U
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for V_c with $V_c = v_c$ (counterfactual values)
3. Prediction: use the sub-model to compute v_e (the effects) by using u , v_c and $v_o \setminus v_c$

Example, non-invertible case:

$$G = g_o$$

$$D = d_c$$

$$R_e = f_R(g_o, d_c, U_R)$$

there might exist multiple values U_R compatible with the observed effect r_o

\mathcal{M} is still completely identified

Counterfactual Inference, probabilistic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation v_o to update the probability distribution $P(U | \mathbf{V}_o = v_o)$
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for \mathbf{V}_c with $\mathbf{V}_c = v_c$ (counterfactual values)
3. Prediction: use the sub-model and the updated distribution to compute the probability of each possible effect v_e

Note that these two may be in contrast with each other

Counterfactual Inference, probabilistic case

From J. Pearl, *Primer*, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation \mathbf{v}_o to update the probability distribution $P(\mathbf{U}|\mathbf{V}_o = \mathbf{v}_o)$
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for \mathbf{V}_c with $\mathbf{V}_c = \mathbf{v}_c$ (counterfactual values)
3. Prediction: use the sub-model and the updated distribution to compute the probability of each possible effect \mathbf{v}_e

Example, non-invertible case:

$$G = g_o$$

$$D = d_c$$

$$P(R_e = r_e) = \sum_{\{u_R \mid f_R(g_o, d_c, u_R) = r_e\}} P(u_R | \mathbf{V}_o = \mathbf{v}_o)$$

all the values u_R compatible with the effect r_e

Counterfactual Inference, probabilistic case

From J. Pearl, Primer, 2016 (see suggested readings)

1. Abduction (*i.e., going in reverse*): use a specific observation \mathbf{v}_o to update the probability distribution $P(\mathbf{U} | \mathbf{V}_o = \mathbf{v}_o)$
2. Action: create a sub-model of \mathcal{M} by replacing the structural equations for \mathbf{V}_c with $\mathbf{V}_c = \mathbf{v}_c$ (counterfactual values)
3. Prediction: use the sub-model and the updated distribution to compute the probability of each possible effect \mathbf{v}_e

This is called unit or individual-level counterfactual inference since it starts from the observation (possibly complete) of a specific case

It requires the complete identification of \mathcal{M} (including the distribution $P(\mathbf{U}, \mathbf{V})$)

Otherwise, there are too many degrees of freedom and the inference problem is ill-posed

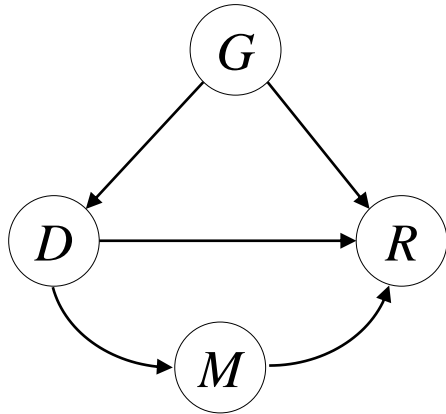
Counterfactual Inference, population level

What kind of counterfactual inference can be performed when the model \mathcal{M} is NOT completely identified?

In other words, when what we have is the distribution $P(\mathbf{V})$ over endogenous variables as derived from actual observations?

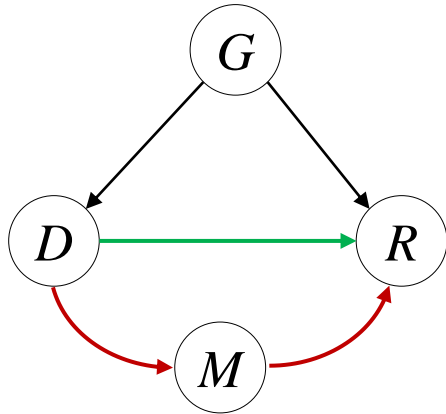
*Perhaps we should change the question somewhat:
what other kind of counterfactual inference could be useful in such case?*

Path-Specific Counterfactual Evidence



Suppose that, as an extension to the previous model, we now assume that drug D has an observable side-effect M which also affects patient's recovery R
It is independent from gender G

Path-Specific Counterfactual Evidence



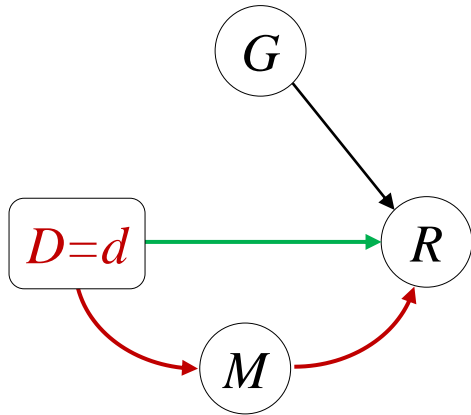
Suppose that, as an extension to the previous model, we now assume that drug D has an observable side-effect M which also affects patient's recovery R

It is independent from gender G

Now the model has two causal paths: one direct and another indirect, mediated by M

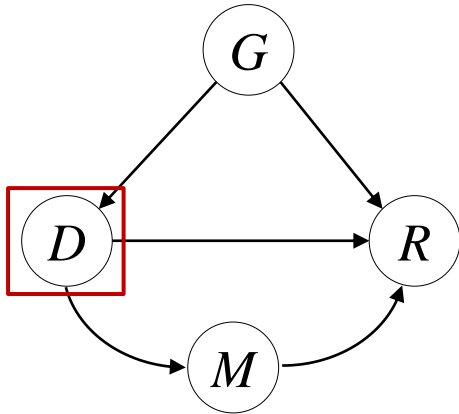
We might want to know what are the causal effects of each path in general, i.e., at the population level

Path-Specific Counterfactual Evidence



Intervention on D alone will not give the answer, as both paths need to be considered at once

Path-Specific Counterfactual Evidence



General idea (intuitive): splitting node D in two and letting different paths 'see' different values

SCM model \mathcal{M}

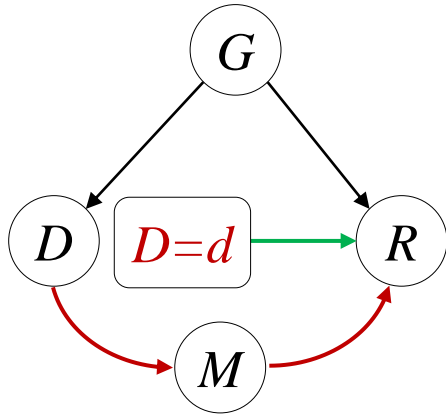
$$G := U_G$$

$$D := f_D(G, U_D)$$

$$M := f_M(D, U_M)$$

$$R := f_R(G, D, M, U_R)$$

Path-Specific Counterfactual Evidence



General idea (intuitive): splitting node D in two and letting different paths 'see' different values

Modified SCM model \mathcal{M}'

$$G := U_G$$

$$D := f_D(G, U_D)$$

When this is feasible, differences in path-specific effects can be evaluated from the distribution $P(\mathbf{V})$ alone

$$M := f_M(D, U_M)$$

$$R := f_R(G, \boxed{D = d}, M, U_R)$$

Counterfactual Inference

(see GeNIe 'berkeley_path_specific' attachment)

Identifiability of Path-Specific Effects

■ Path-Specific Criterion (simplified)

In a SCM model \mathcal{M} , *path-specific effects* of T over Y with mediator M are *identifiable* iff it exists an *adjustment set* W of variables such that:

- no variable in W is on, or is a descendant of any variables on, a **causal path** (excluding the descendants of T alone)
- the variables in W block (*in the sense of graphical models*) all the **non-causal paths** between T and Y
- the variables in W block (*in the sense of graphical models*) all the **non-causal paths** between T and M
- the variables in W block (*in the sense of graphical models*) all the **non-causal paths** between M and Y