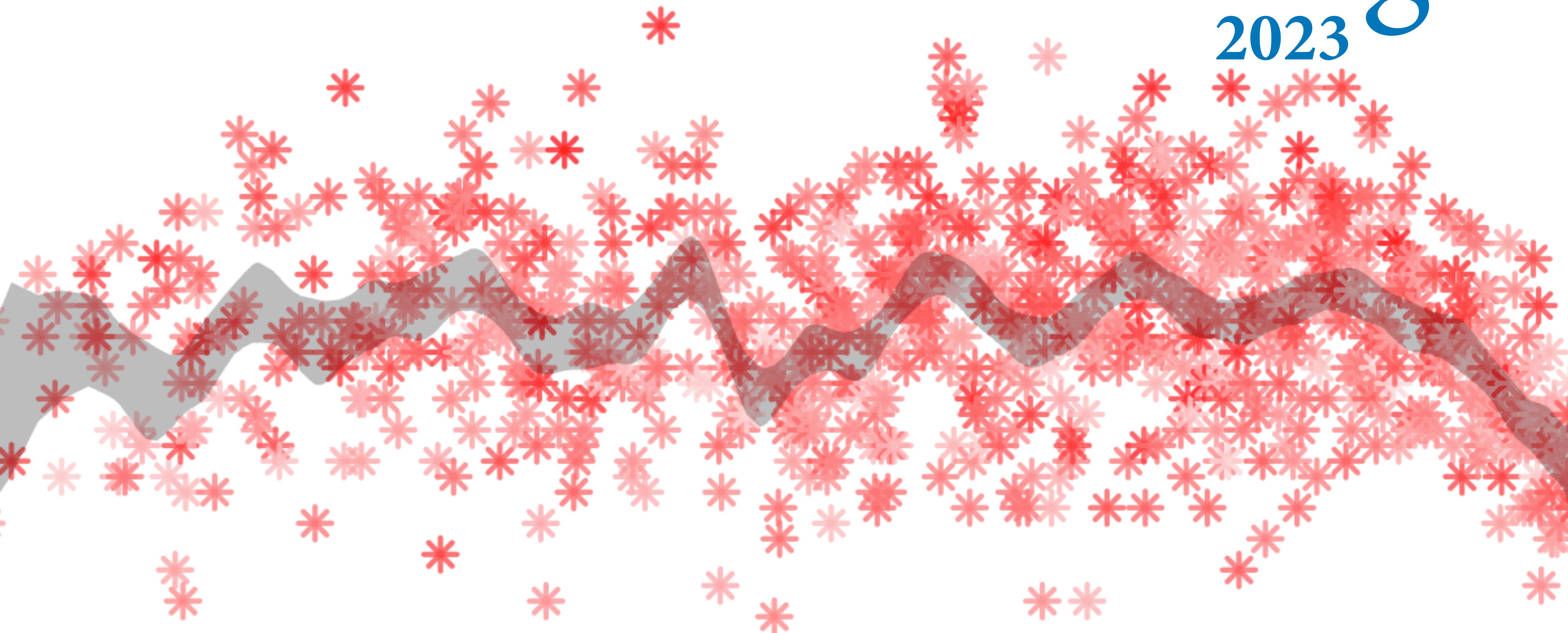


# Statistical Rethinking

2023



6. Good & Bad Controls

# Avoid Being Clever At All Costs

Being clever: unreliable, opaque

Given a causal model, can use logic  
to derive implications

Others can use same logic to verify  
& challenge your work



*Better than clever*



*The Fork*

$$X \leftarrow Z \rightarrow Y$$



*The Collider*

$$X \rightarrow Z \leftarrow Y$$

*The Pipe*

$$X \rightarrow Z \rightarrow Y$$



*The Descendant*

$$X \rightarrow Z \rightarrow Y$$





*The Fork*



$X$  and  $Y$  associated  
unless stratify by  $Z$



*The Pipe*



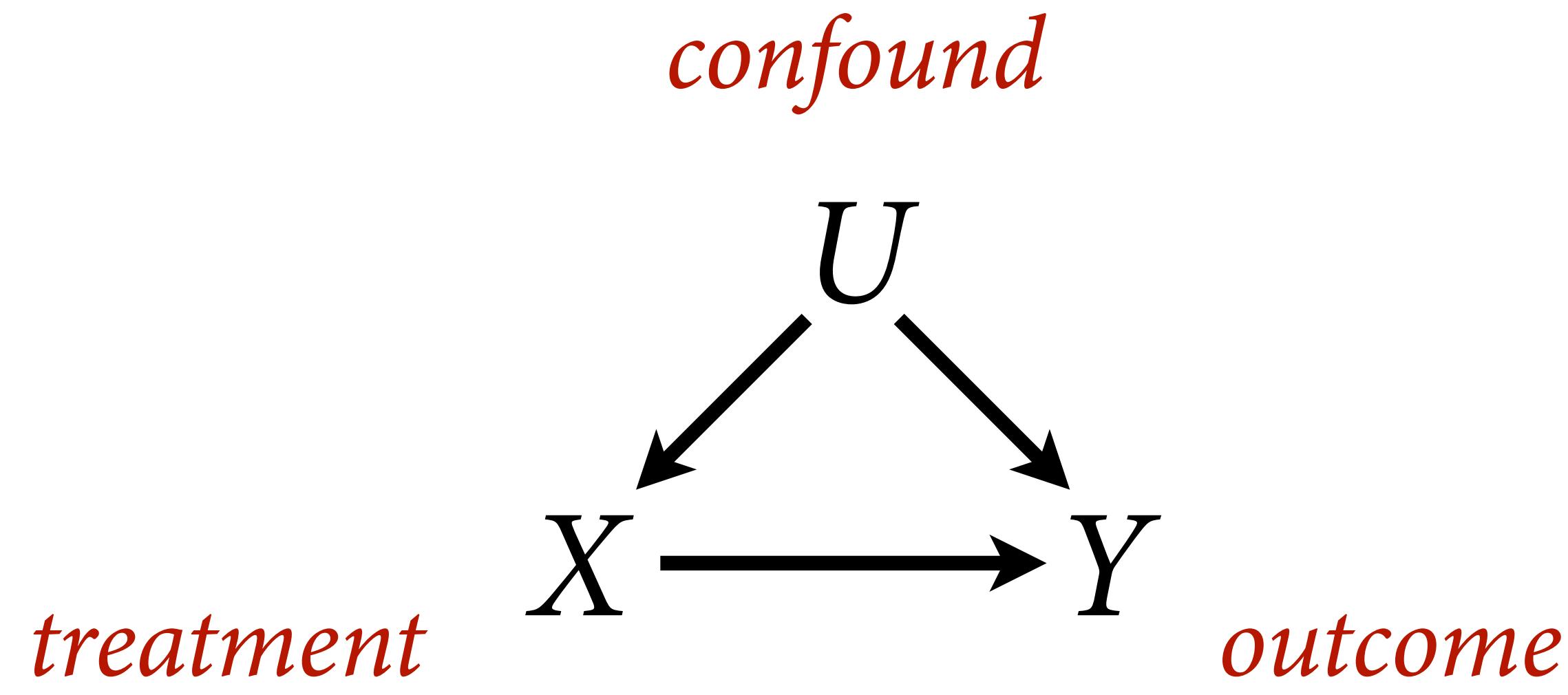
$X$  and  $Y$  associated  
unless stratify by  $Z$



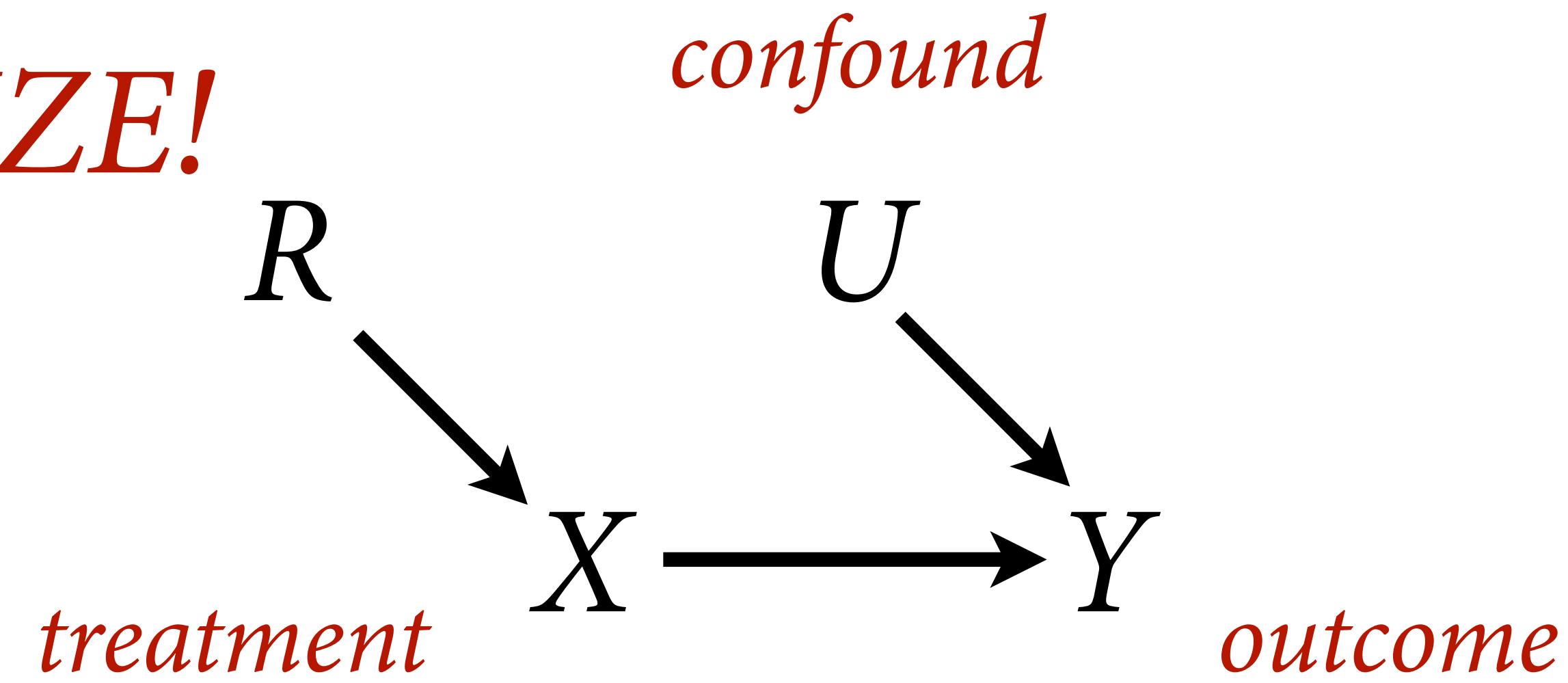
*The Collider*

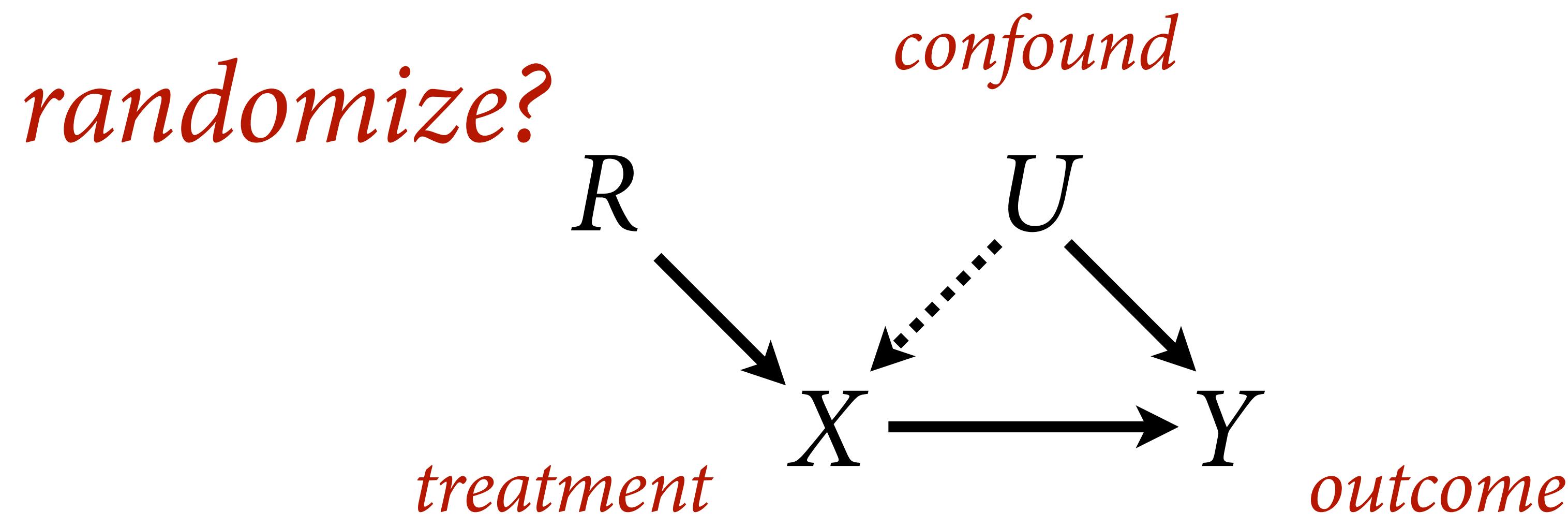


$X$  and  $Y$  not associated  
unless stratify by  $Z$



*RANDOMIZE!*





# Causal Thinking

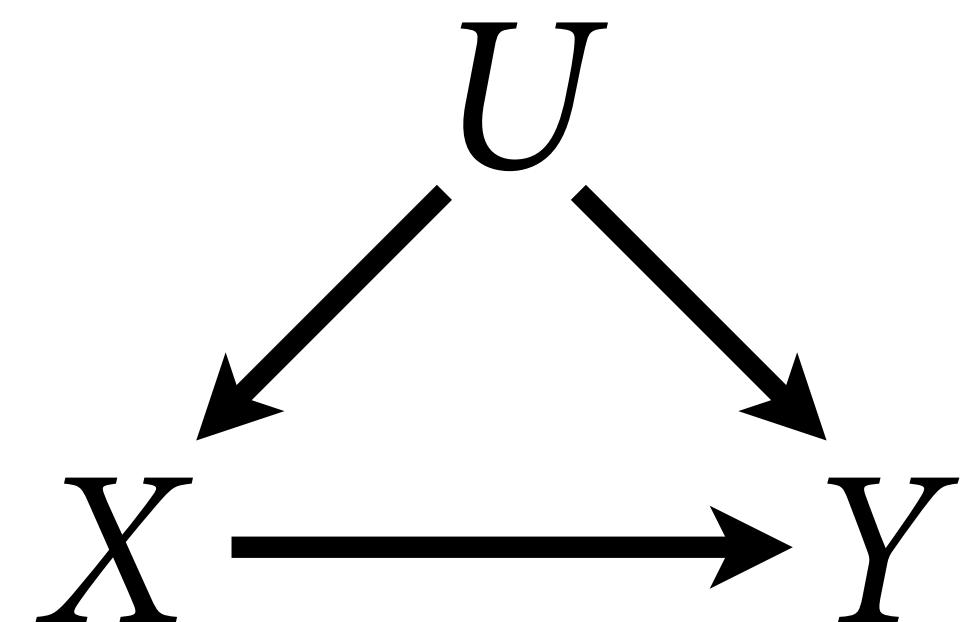
In an experiment, we cut causes of the treatment

We *randomize* (we try at least)

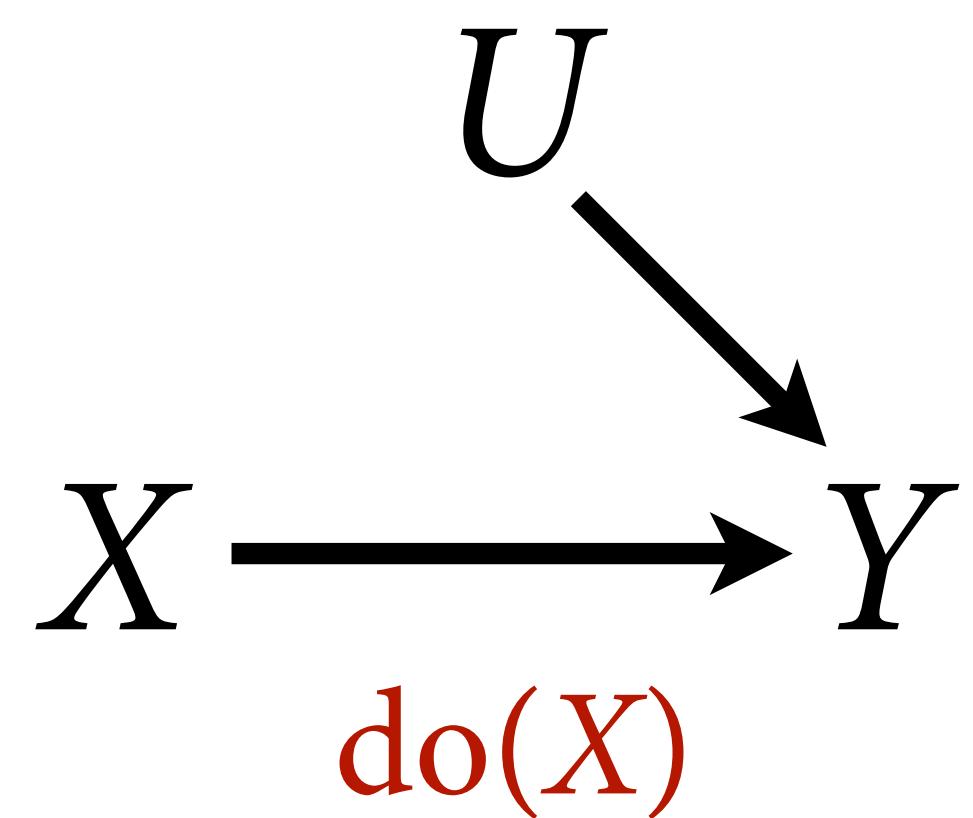
So how does causal inference without randomization ever work?

Is there a statistical procedure that mimics randomization?

*Without randomization*



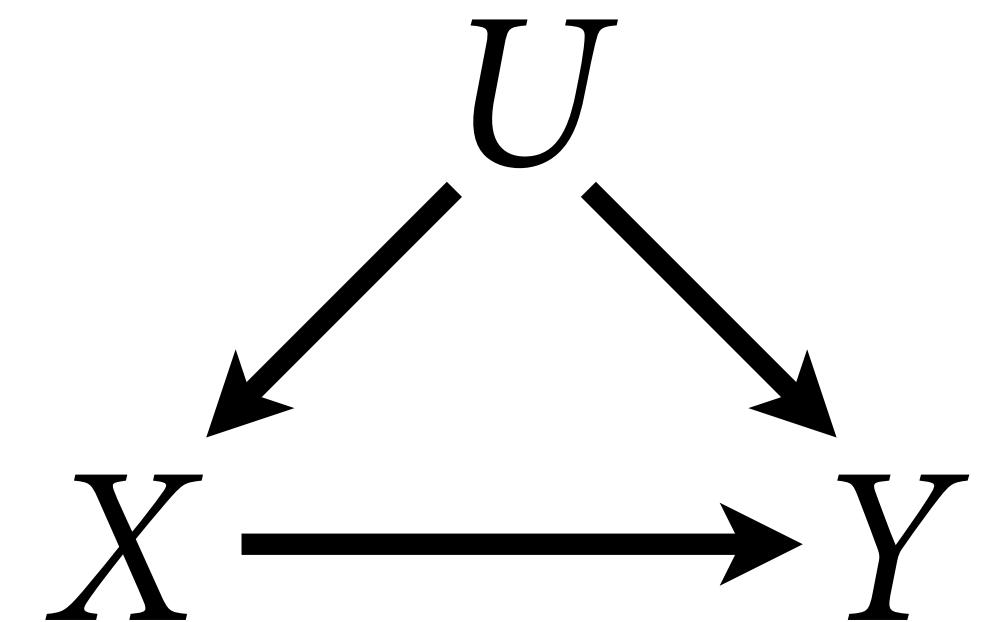
*With randomization*



# Causal Thinking

*Without randomization*

Is there a statistical procedure  
that mimics randomization?

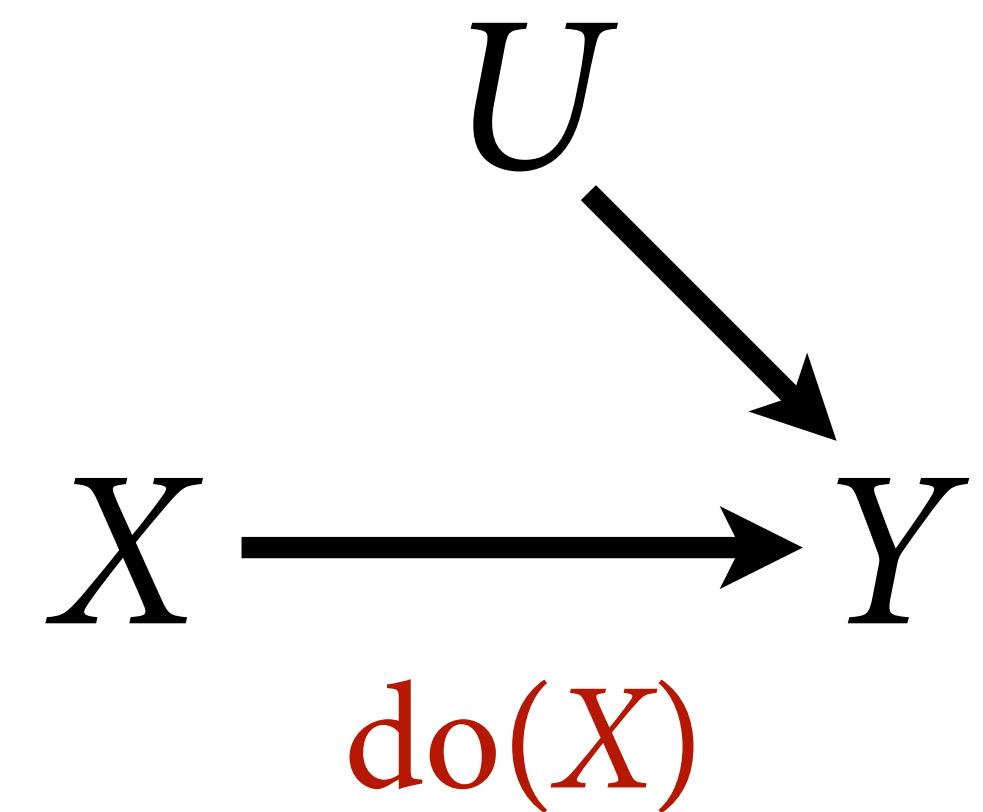


$$P(Y \mid \text{do}(X)) = P(Y \mid ?)$$

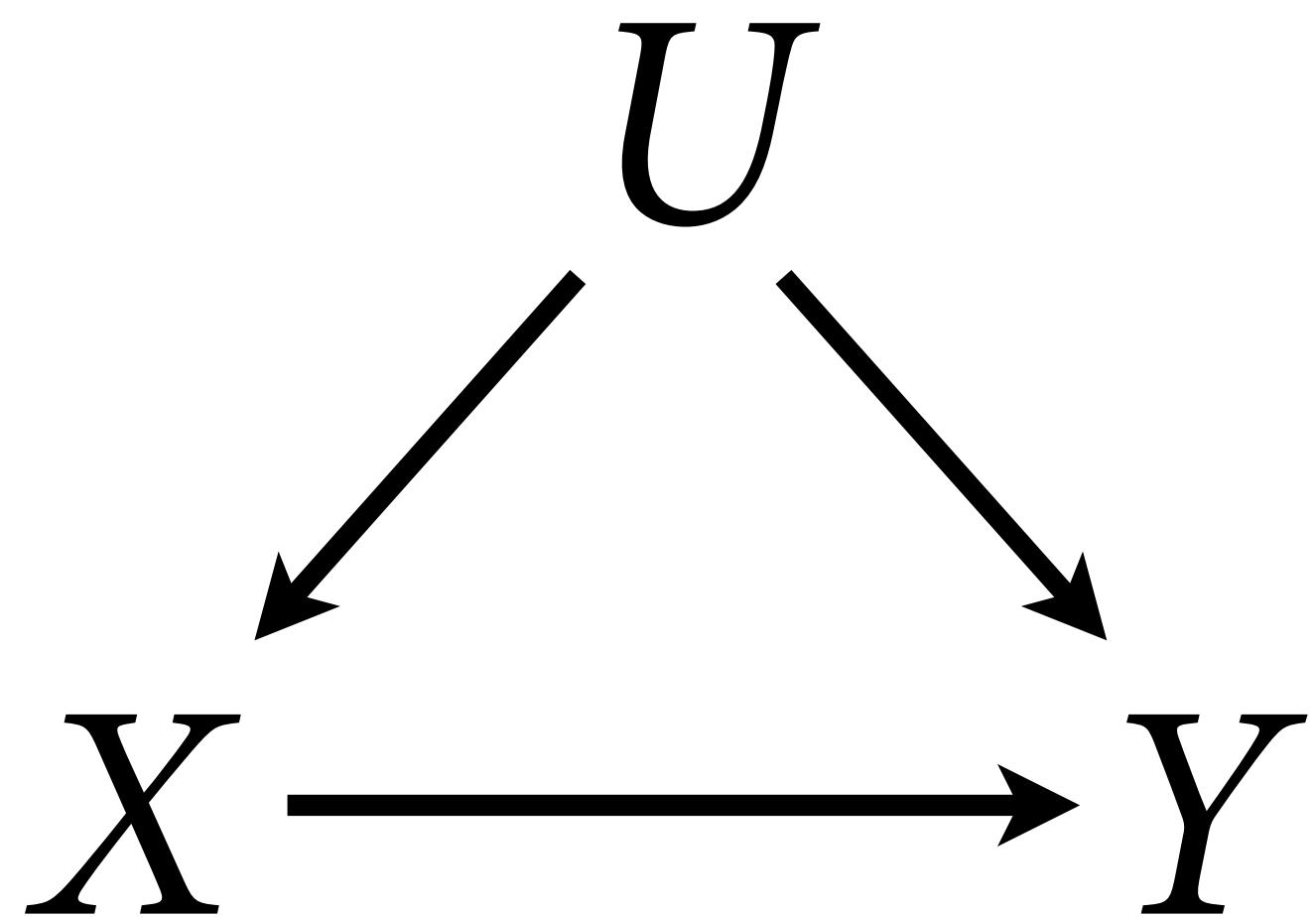
$\text{do}(X)$  means intervene on  $X$

Can analyze causal model to  
find answer (if it exists)

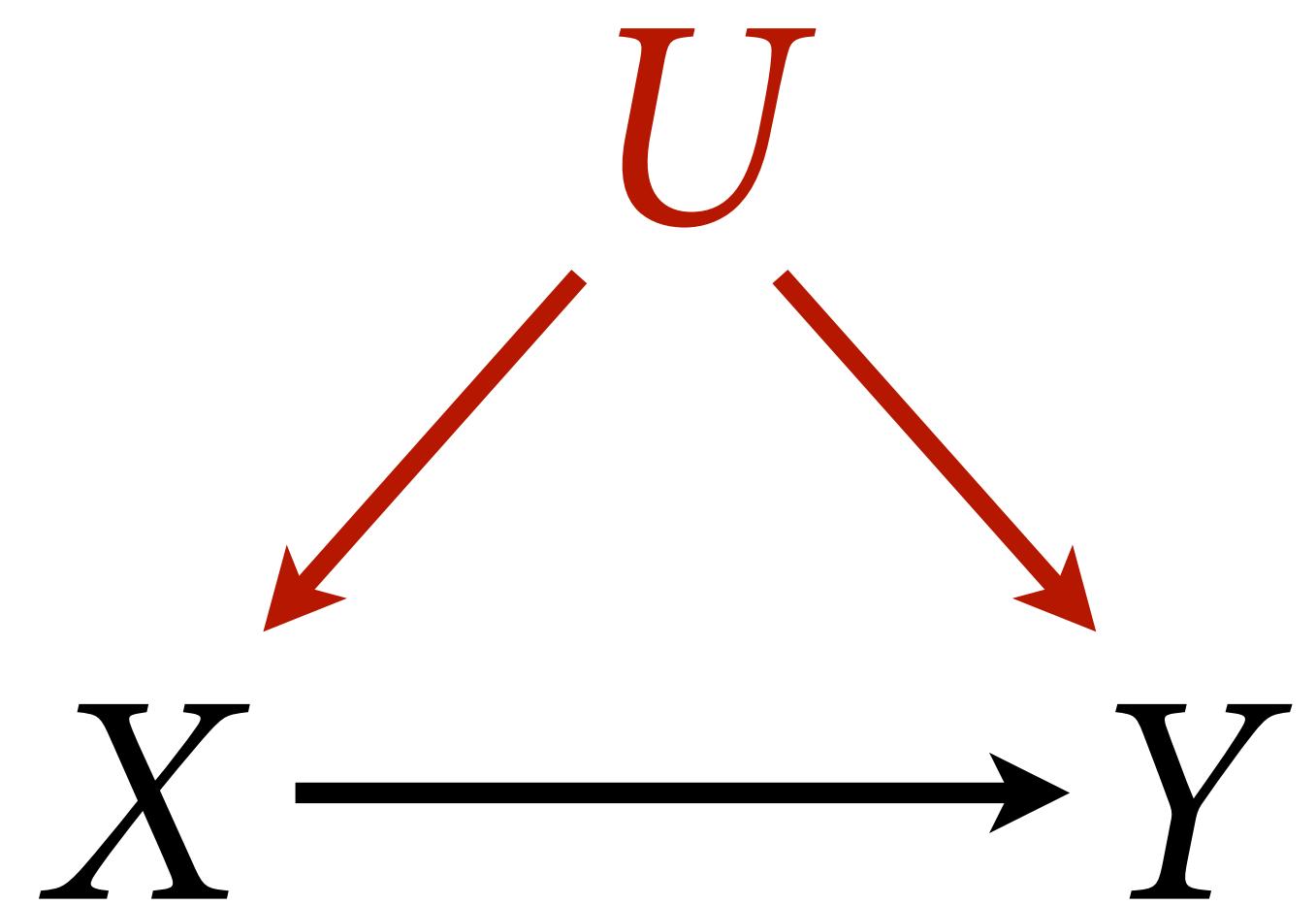
*With randomization*



# Example: Simple Confound



# Example: Simple Confound



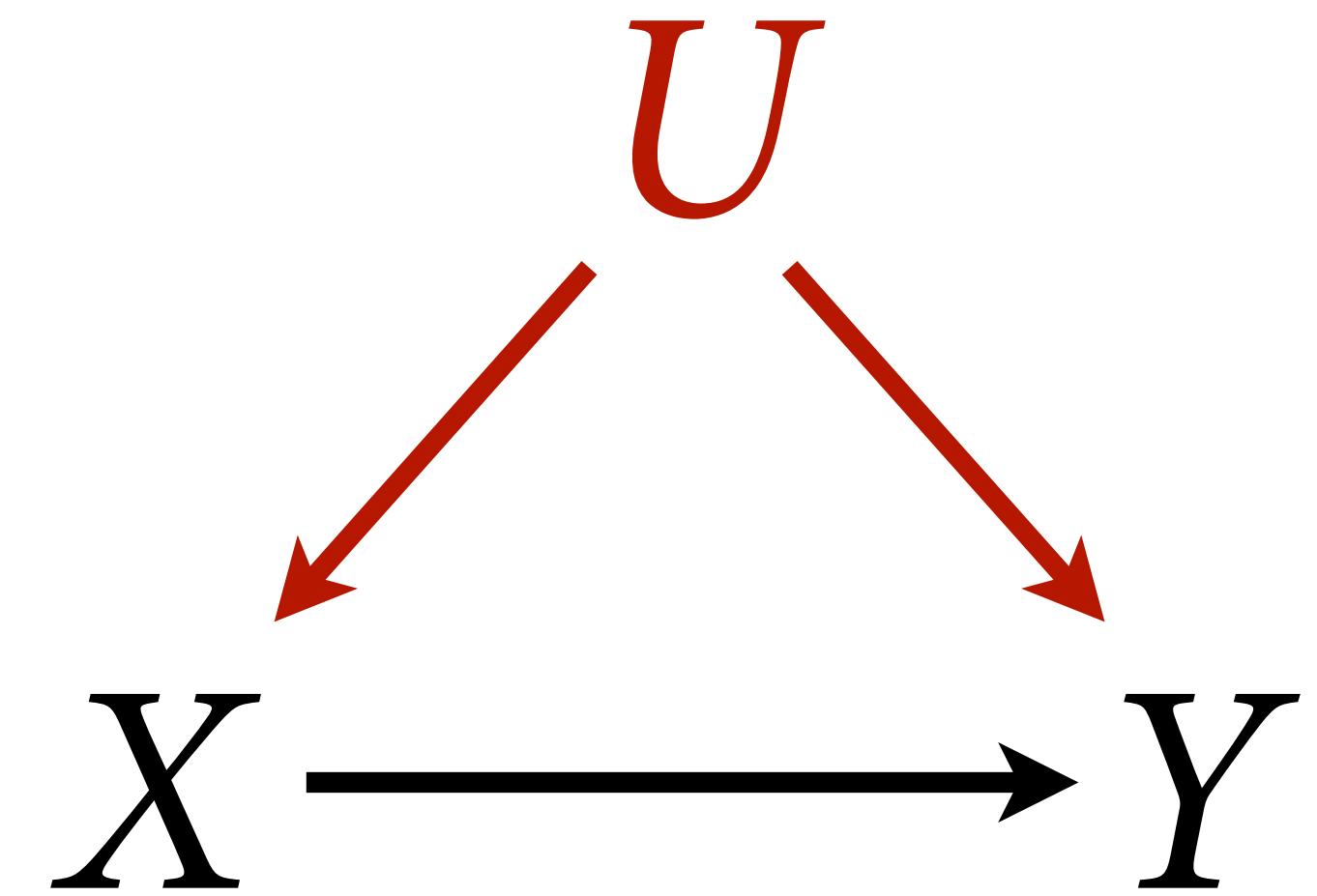
*Non-causal path*

$X <- U -> Y$

*Close the fork!*

*Condition on U*

# Example: Simple Confound



*Non-causal path*

$X <- U -> Y$

*Close the fork!  
Condition on U*

$$P(Y | \text{do}(X)) = \sum_U P(Y | X, U)P(U) = \mathbb{E}_U P(Y | X, U)$$

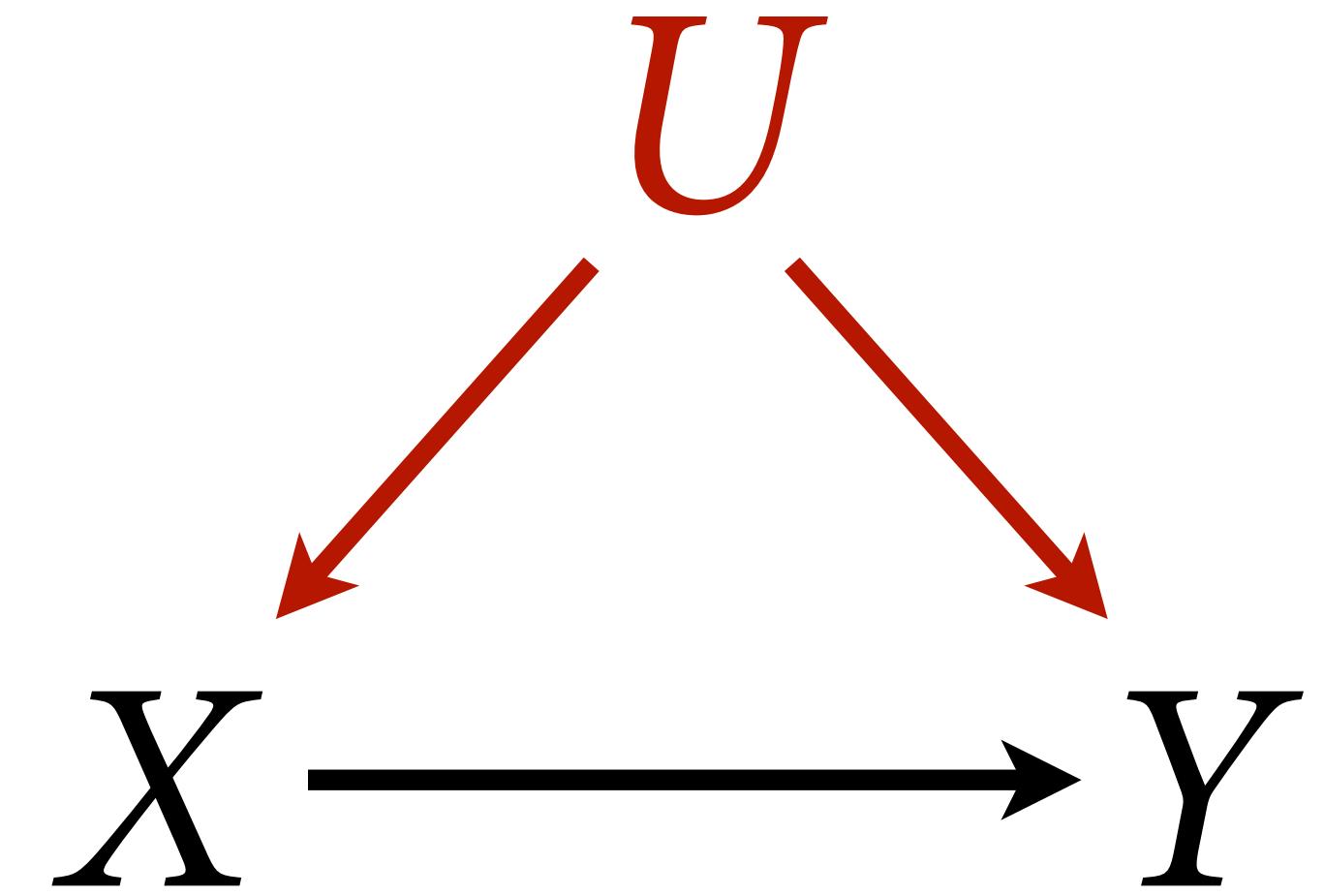
*“The distribution of Y, stratified by X and U,  
averaged over the distribution of U.”*

$$P(Y | \text{do}(X)) = \sum_U P(Y | X, U)P(U) = \text{E}_U P(Y | X, U)$$

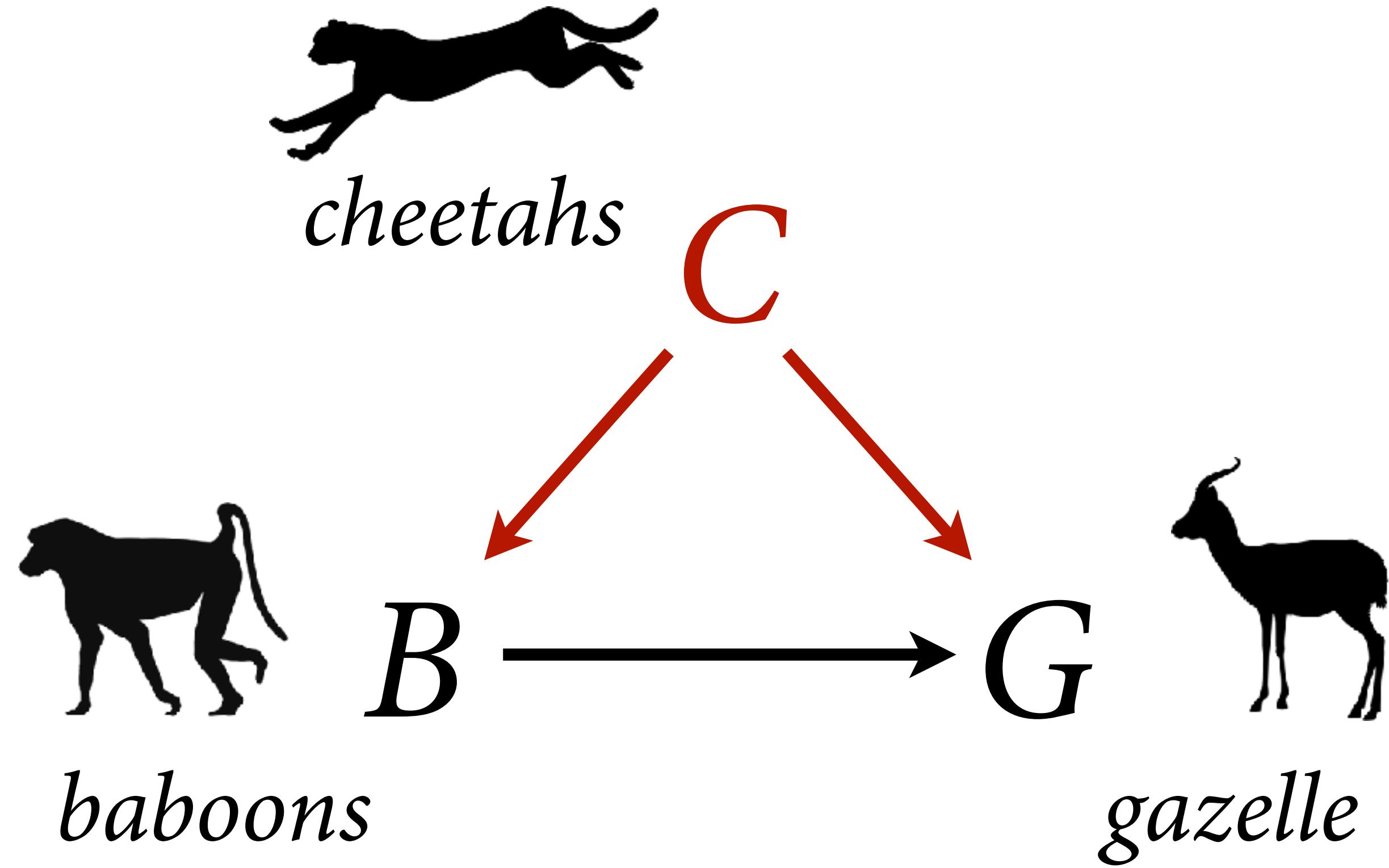
*“The distribution of  $Y$ , stratified by  $X$  and  $U$ , averaged over the distribution of  $U$ .“*

The causal effect of  $X$  on  $Y$  is **not** (in general) the **coefficient** relating  $X$  to  $Y$

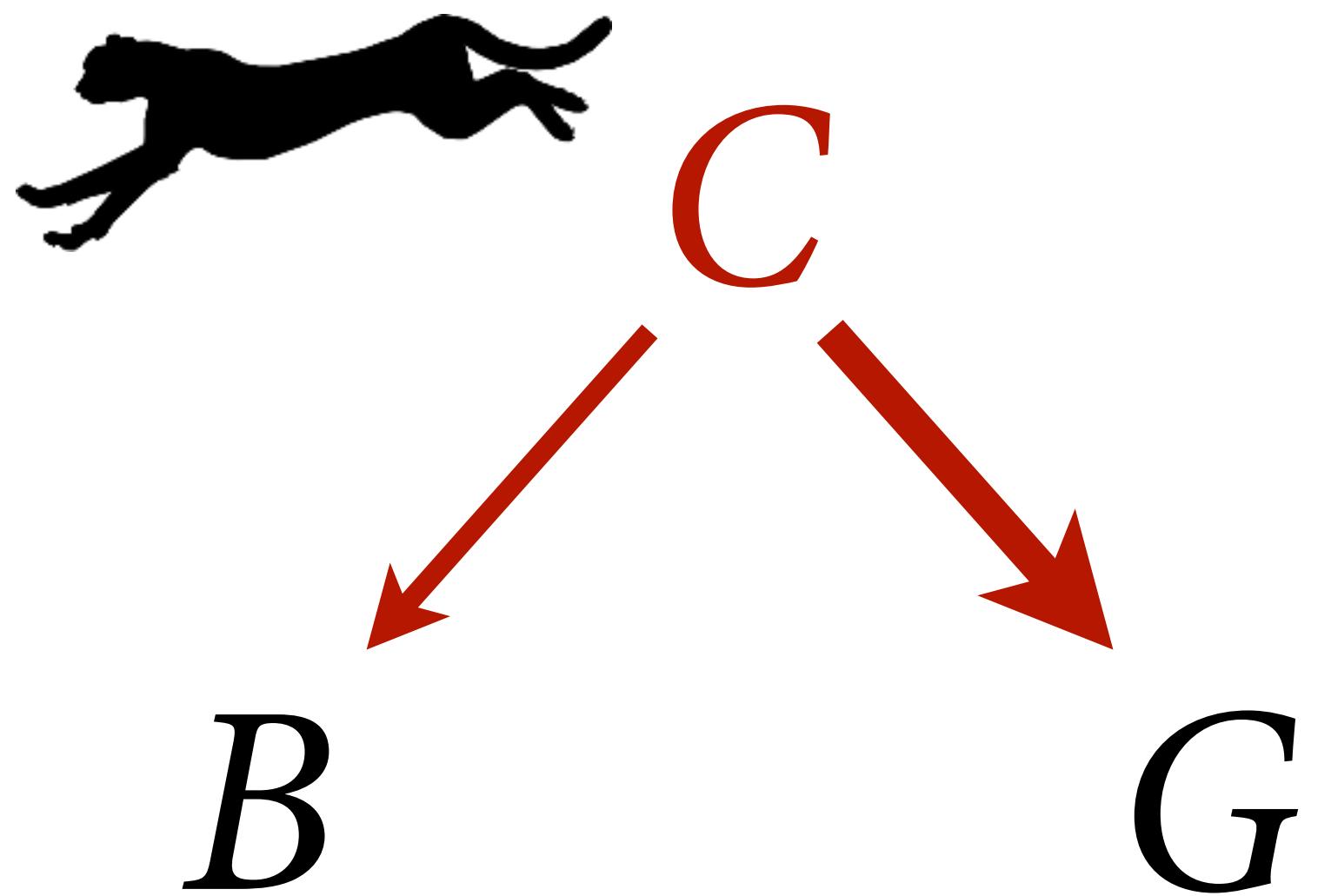
It is the distribution of  $Y$  when we change  $X$ , **averaged** over the distributions of the control variables (here  $U$ )



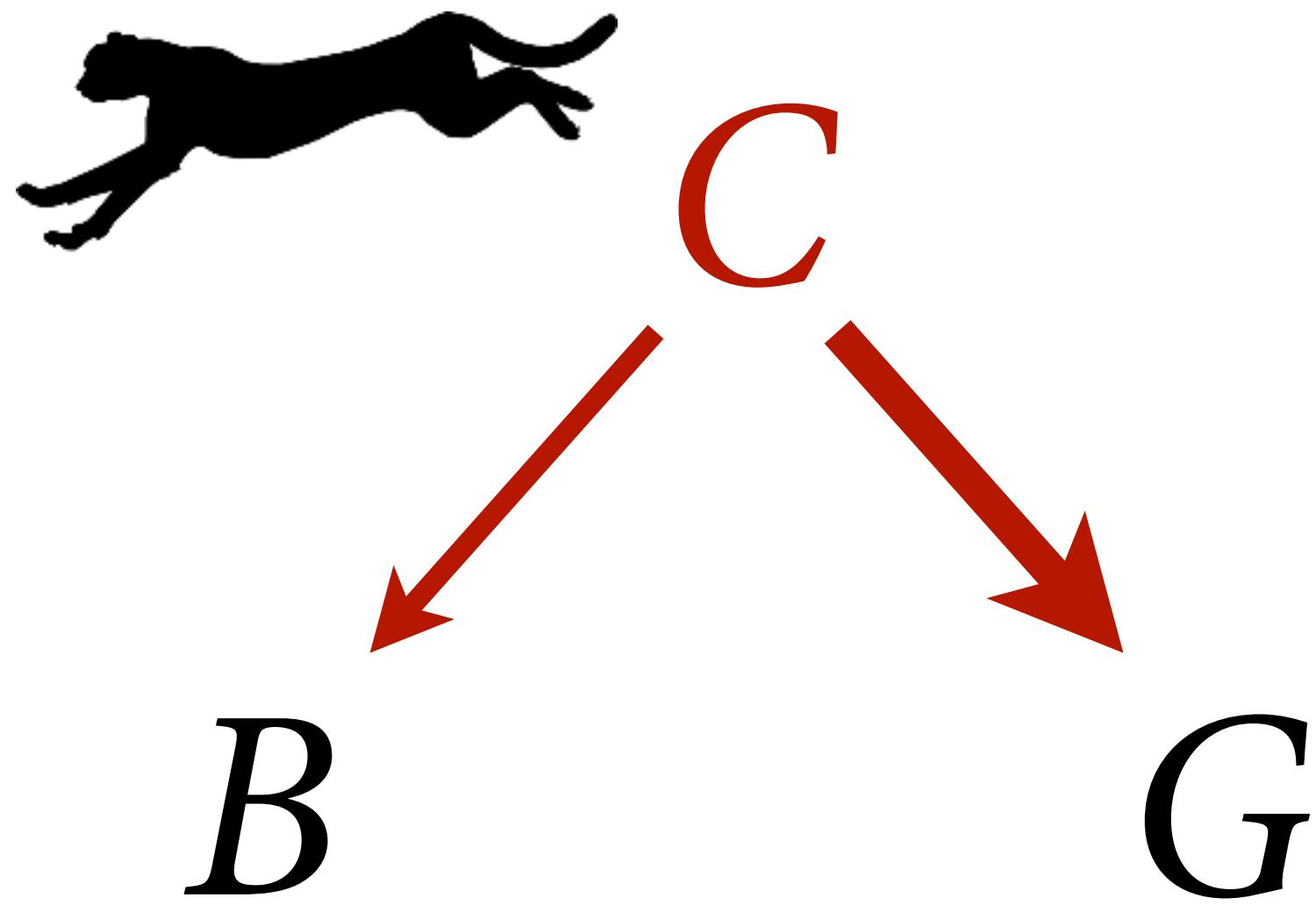
# Marginal Effects Example



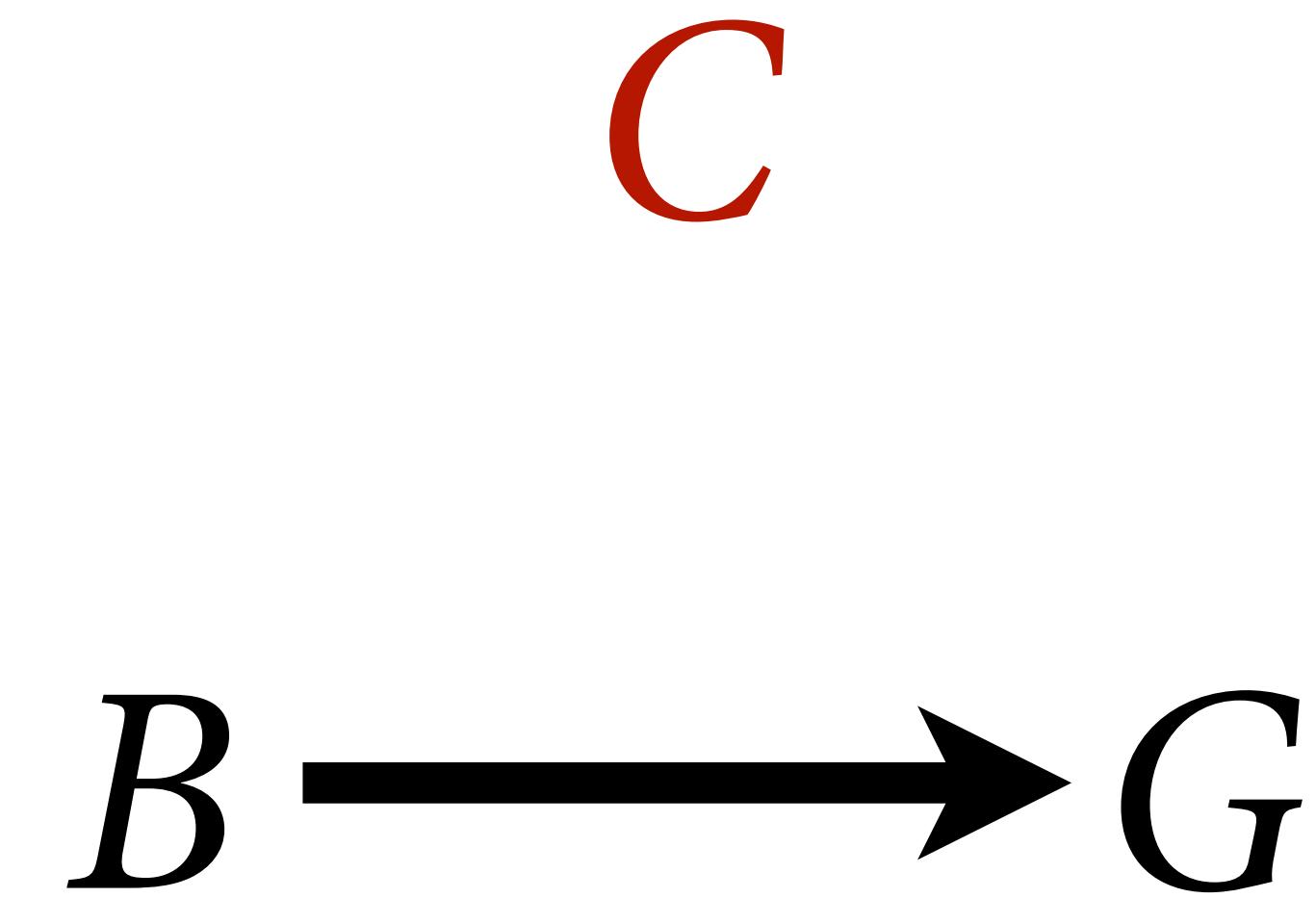
*cheetahs present*



*cheetahs present*



*cheetahs absent*



*Causal effect of baboons depends upon distribution of cheetahs*

# do-calculus

For DAGs, rules for finding  
 $P(Y|do(X))$  known as **do-calculus**

do-calculus says what is possible  
to say **before** picking functions

Justifies **graphical analysis**

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THE BOOK OF WHY

## DO-CALCULUS AT WORK

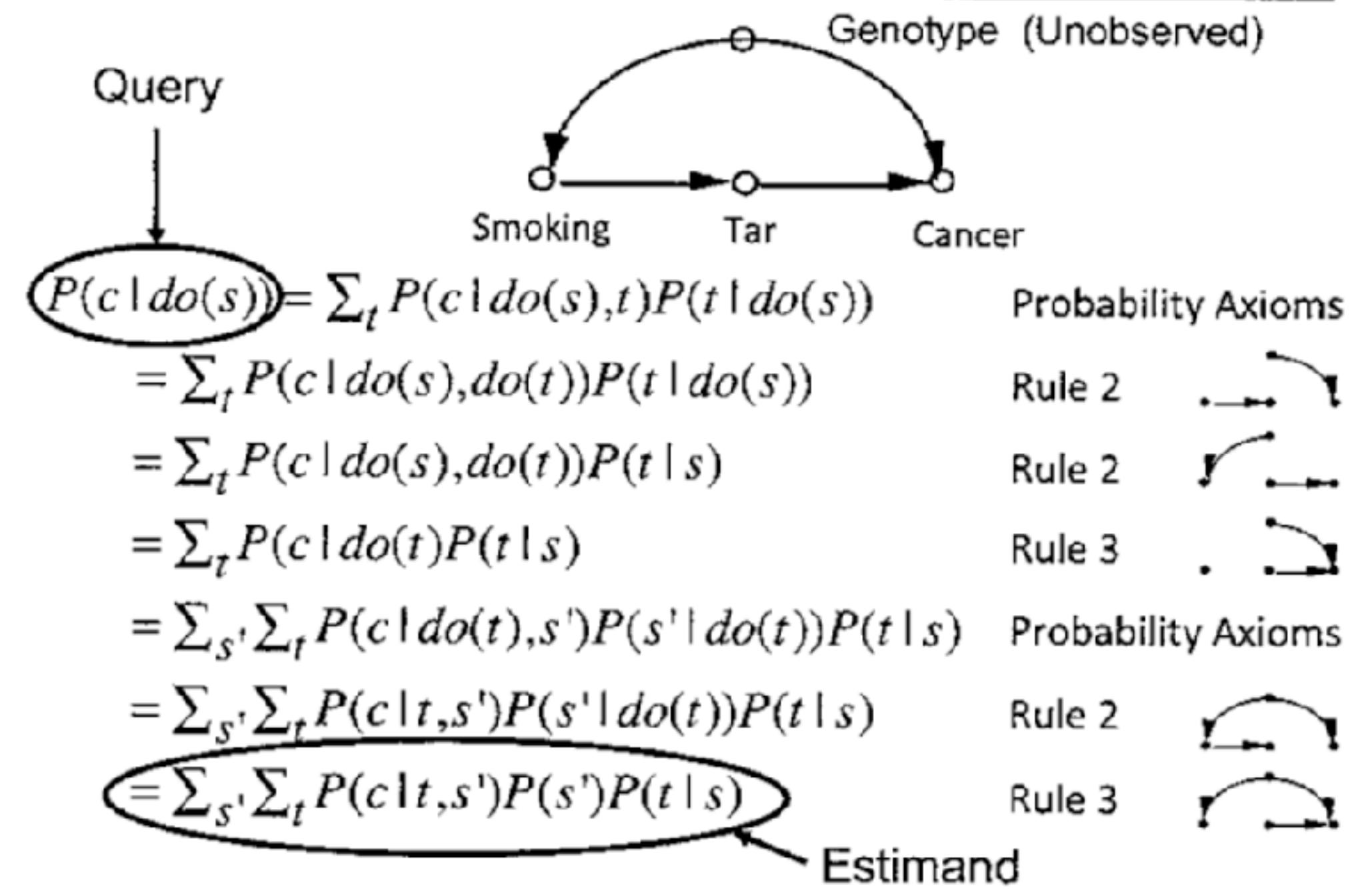


FIGURE 7.4. Derivation of the front-door adjustment formula from the rules of *do*-calculus.

*Do calculus, not too much, mostly graphs*

# do-calculus

do-calculus is **worst case**:  
additional assumptions often  
allow stronger inference

do-calculus is **best case**:  
if inference possible by do-  
calculus, does not depend on  
special assumptions

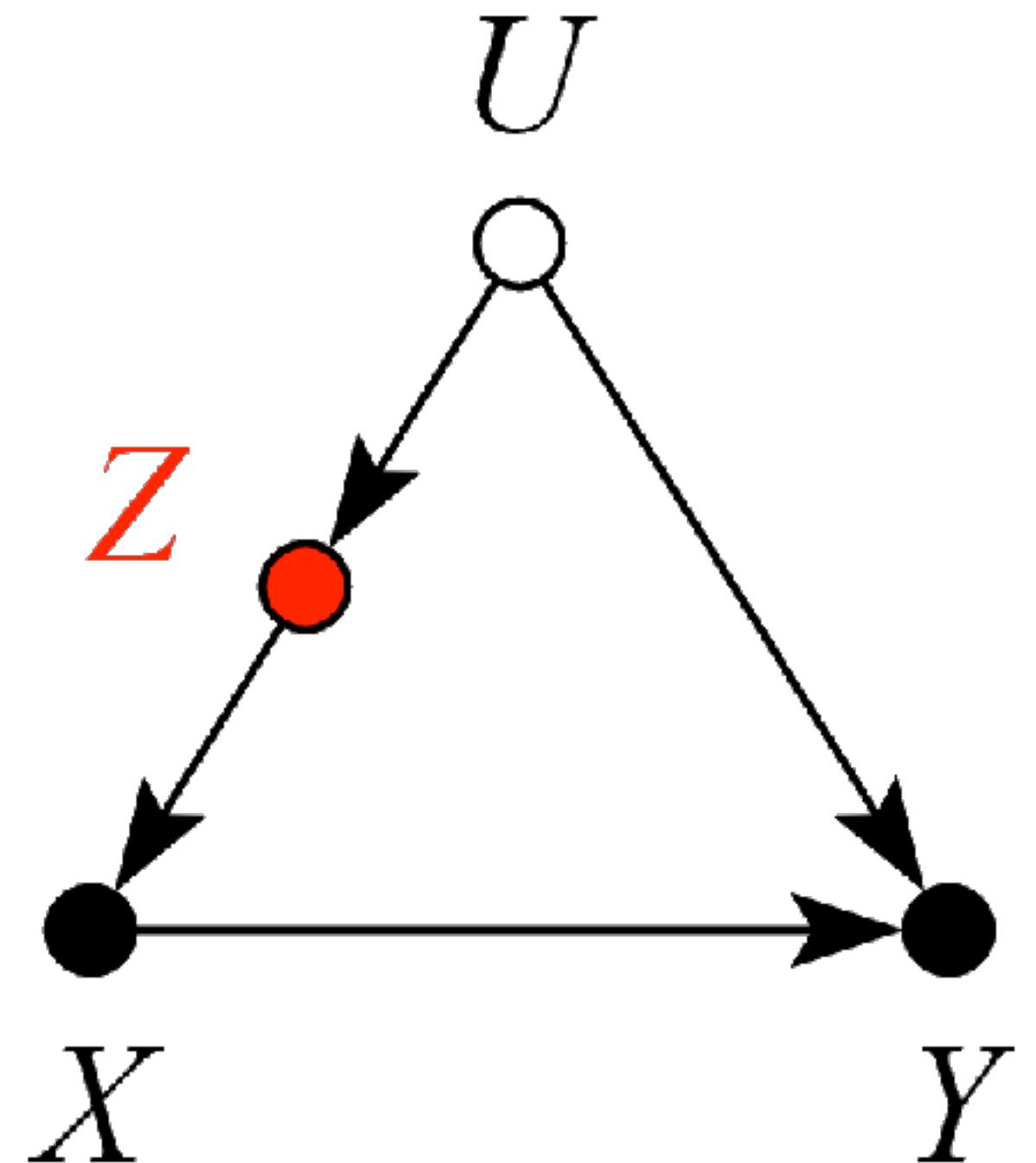


*Judea Pearl, father of do-calculus (1966)*

# Backdoor Criterion

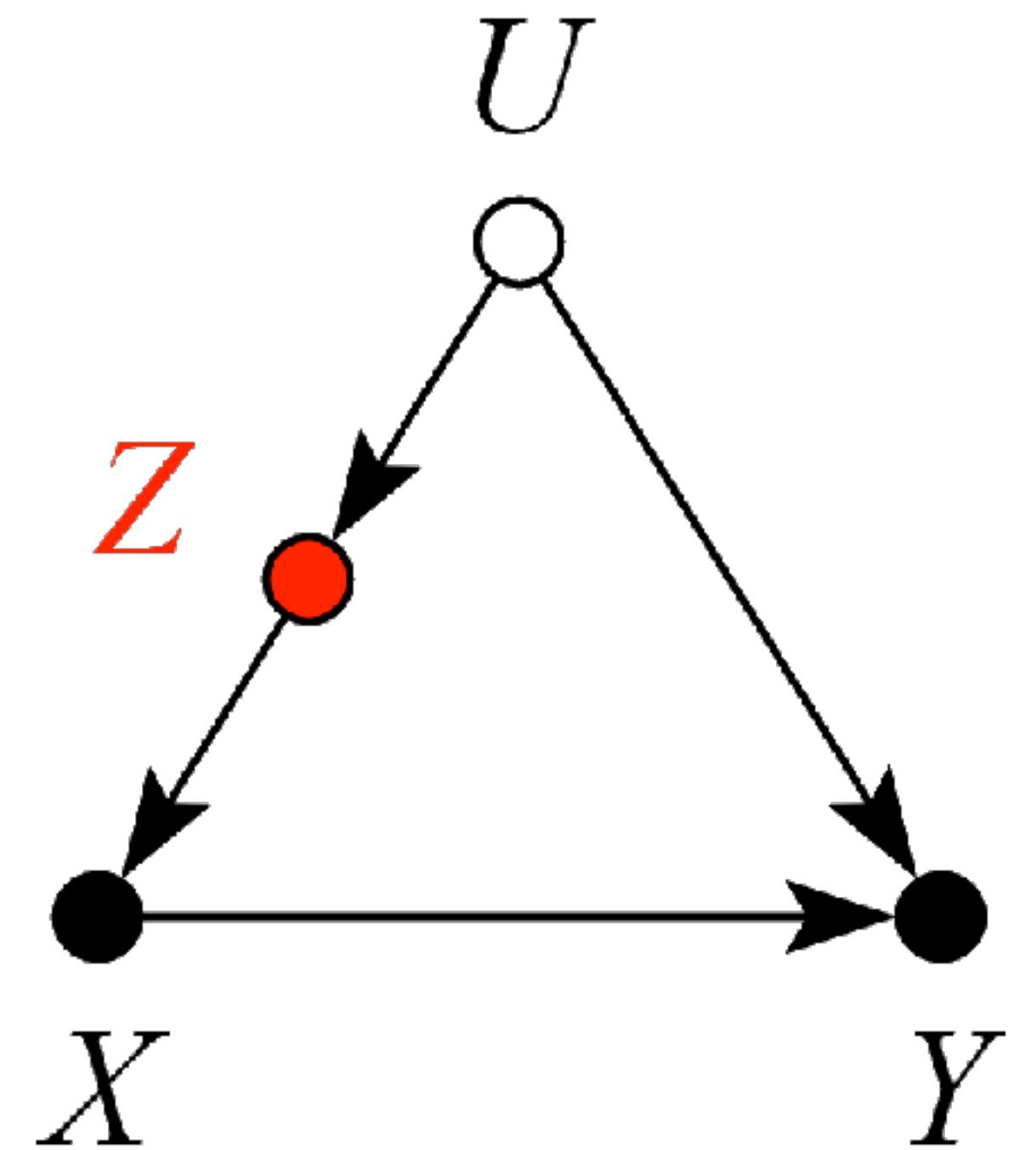
**Backdoor Criterion** is a shortcut to applying (some) results of do-calculus

Can be performed with your eyeballs

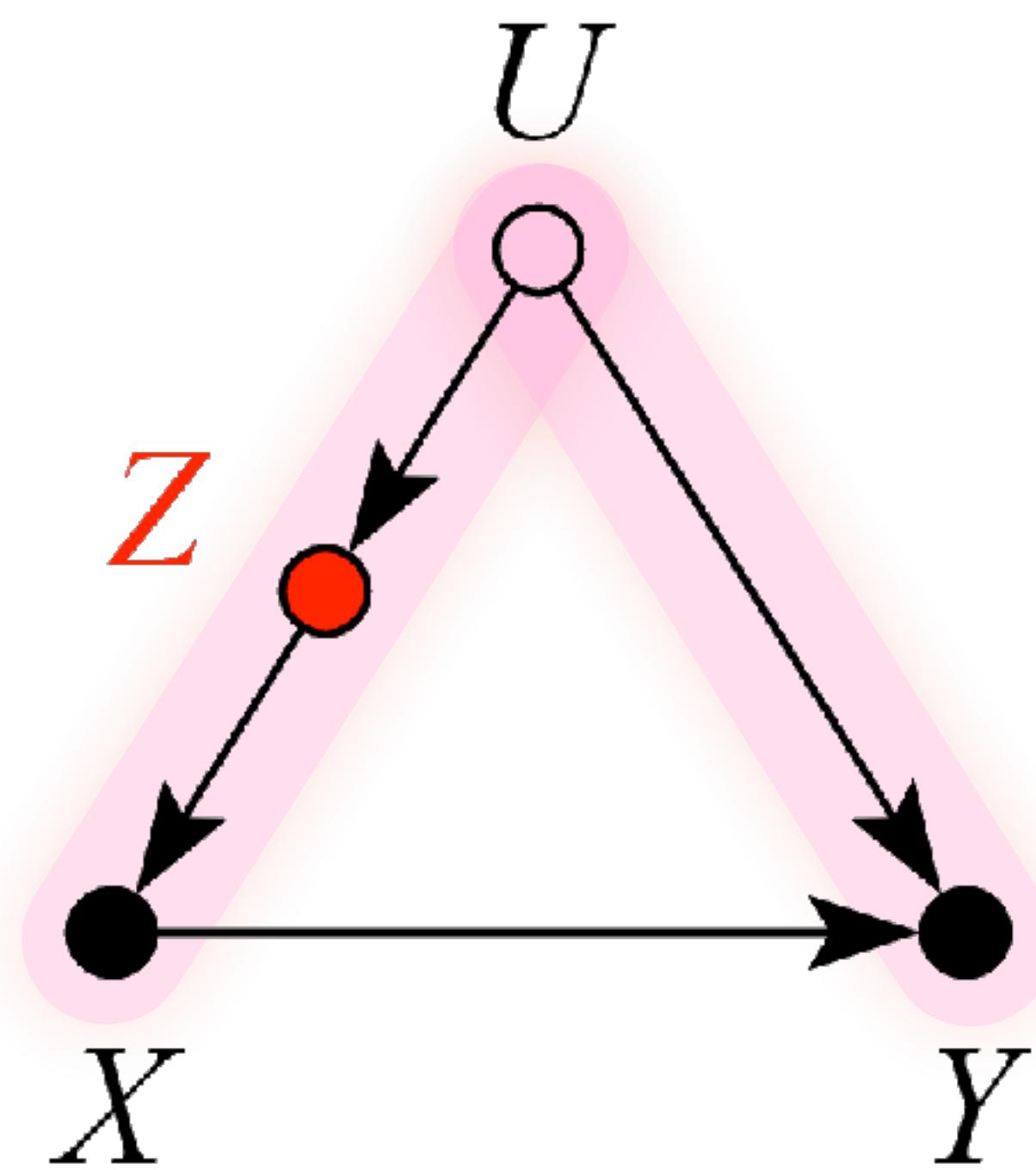
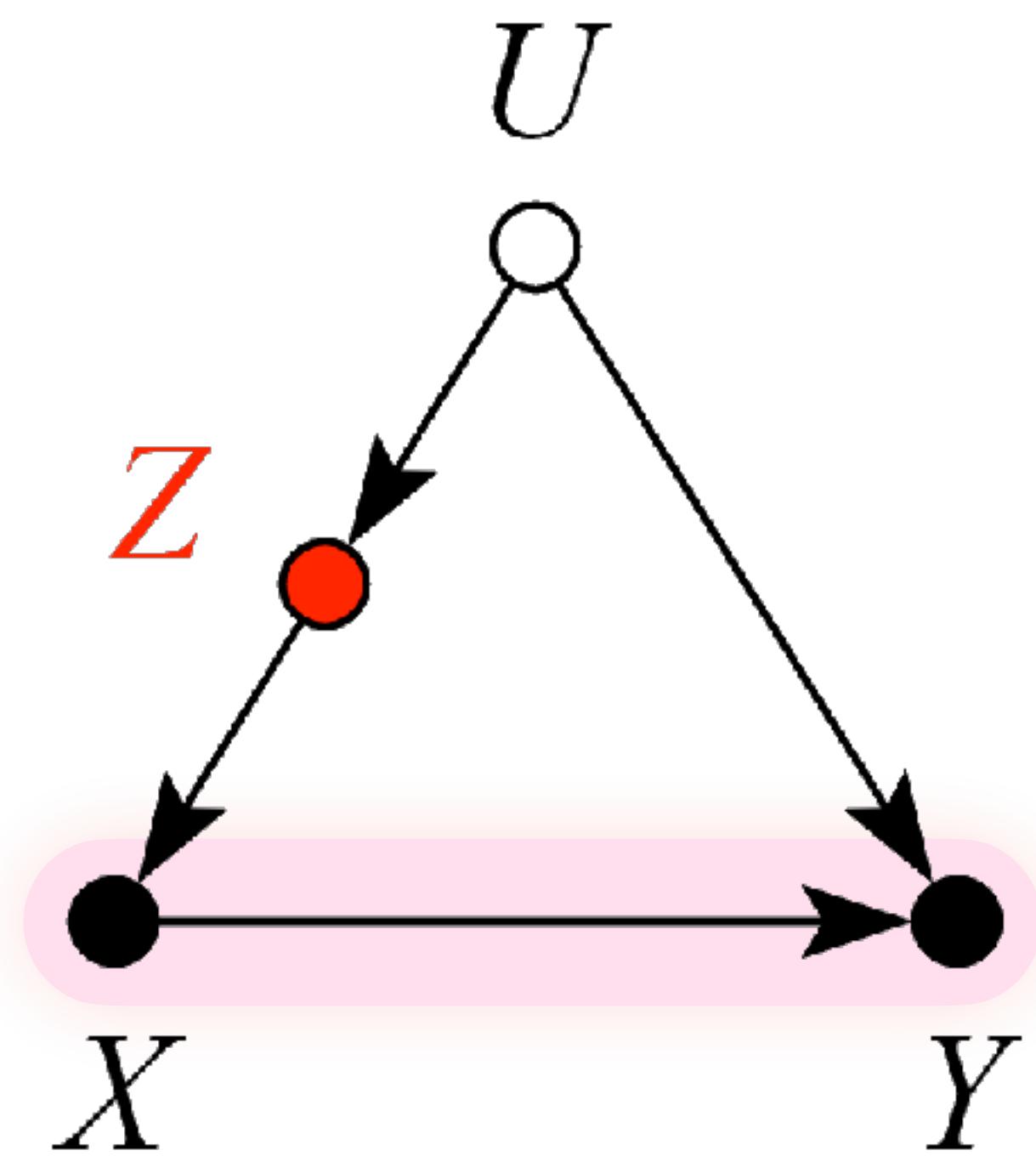


**Backdoor Criterion:** Rule to find a set of variables to stratify by to yield  $P(Y|\text{do}(X))$

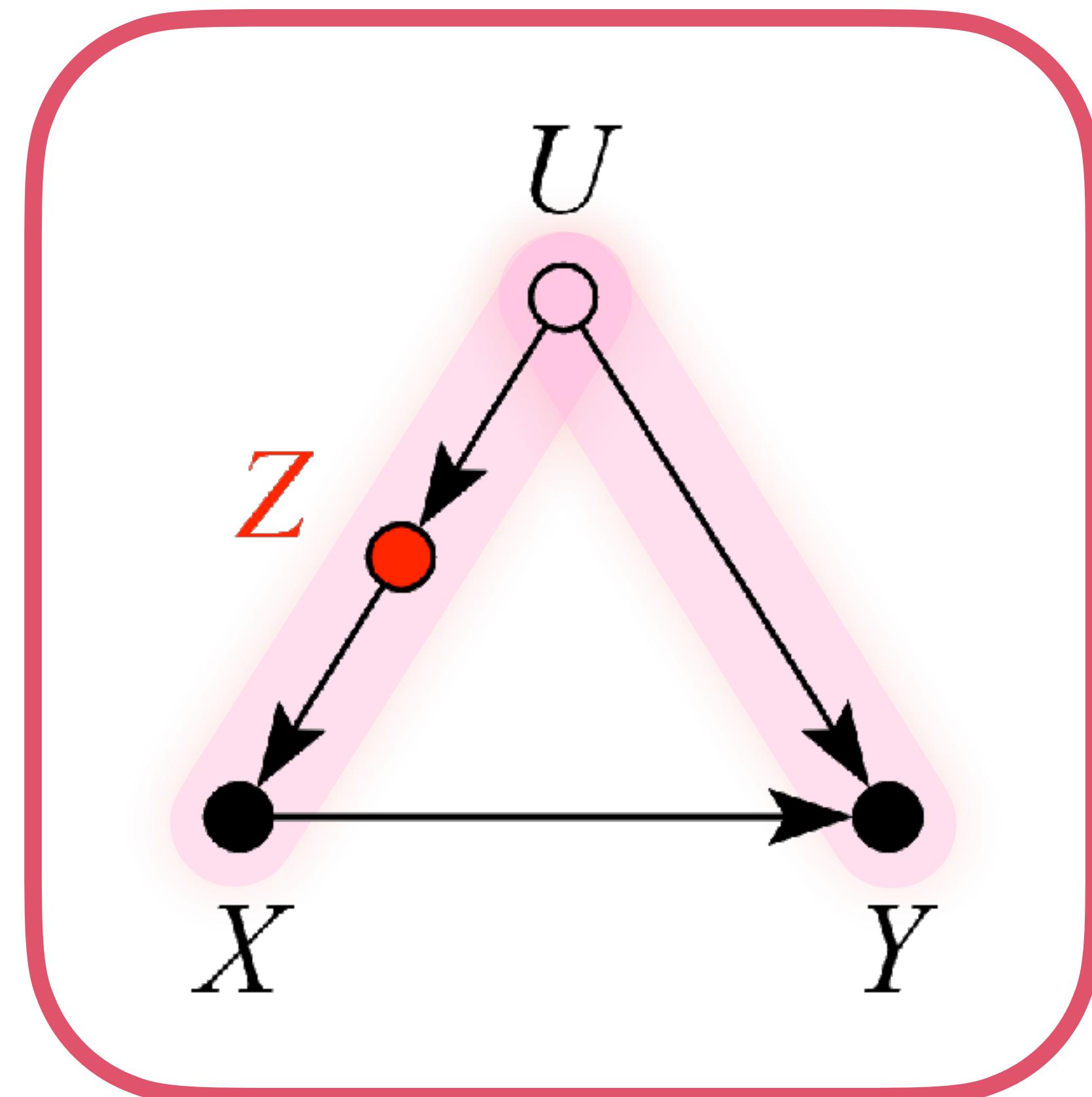
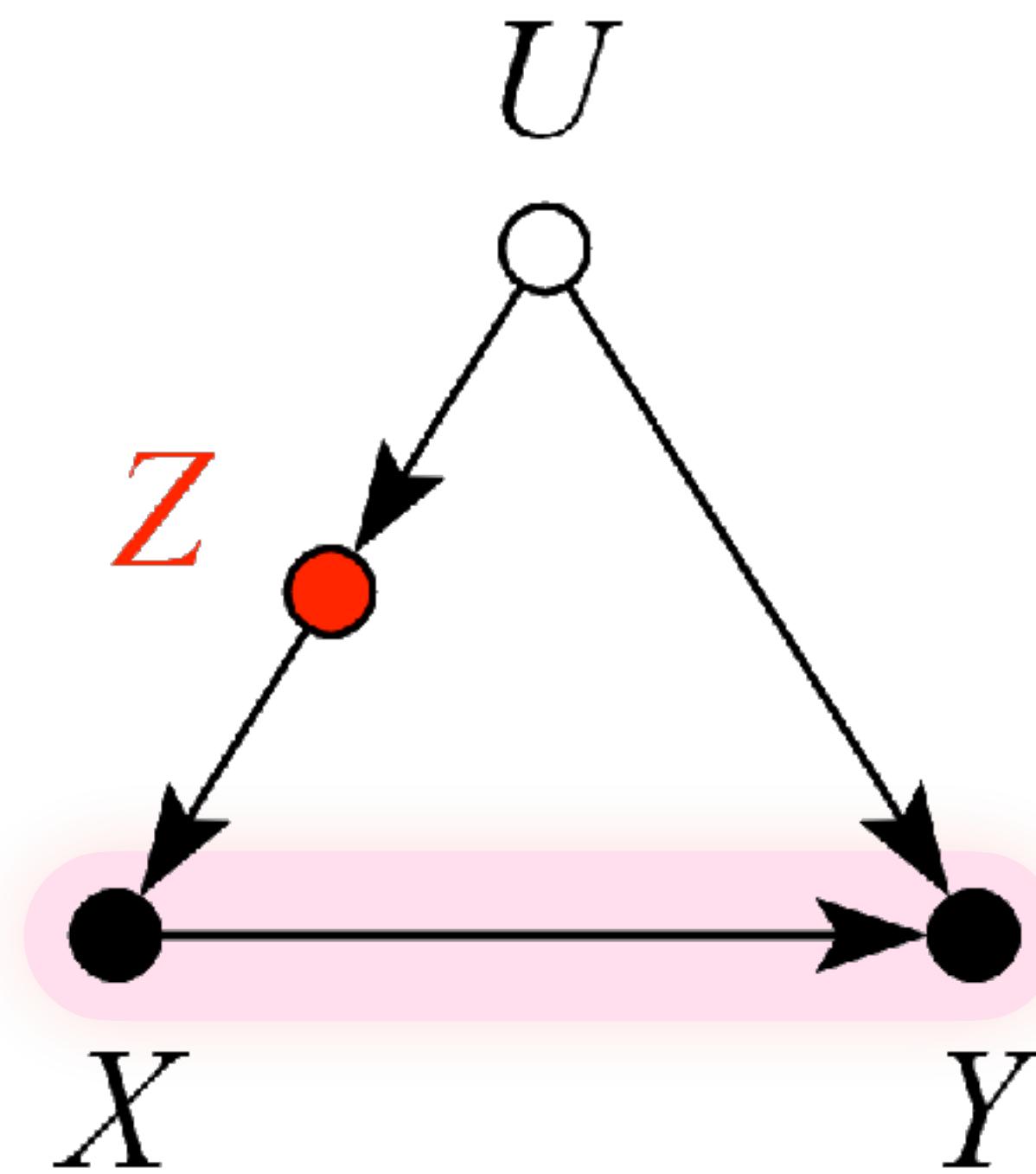
- (1) Identify all **paths** connecting the treatment ( $X$ ) to the outcome ( $Y$ )
- (2) Paths with arrows **entering**  $X$  are backdoor paths (non-causal paths)
- (3) Find **adjustment set** that closes/blocks all backdoor paths



(1) Identify all **paths** connecting the treatment ( $X$ ) to the outcome ( $Y$ )



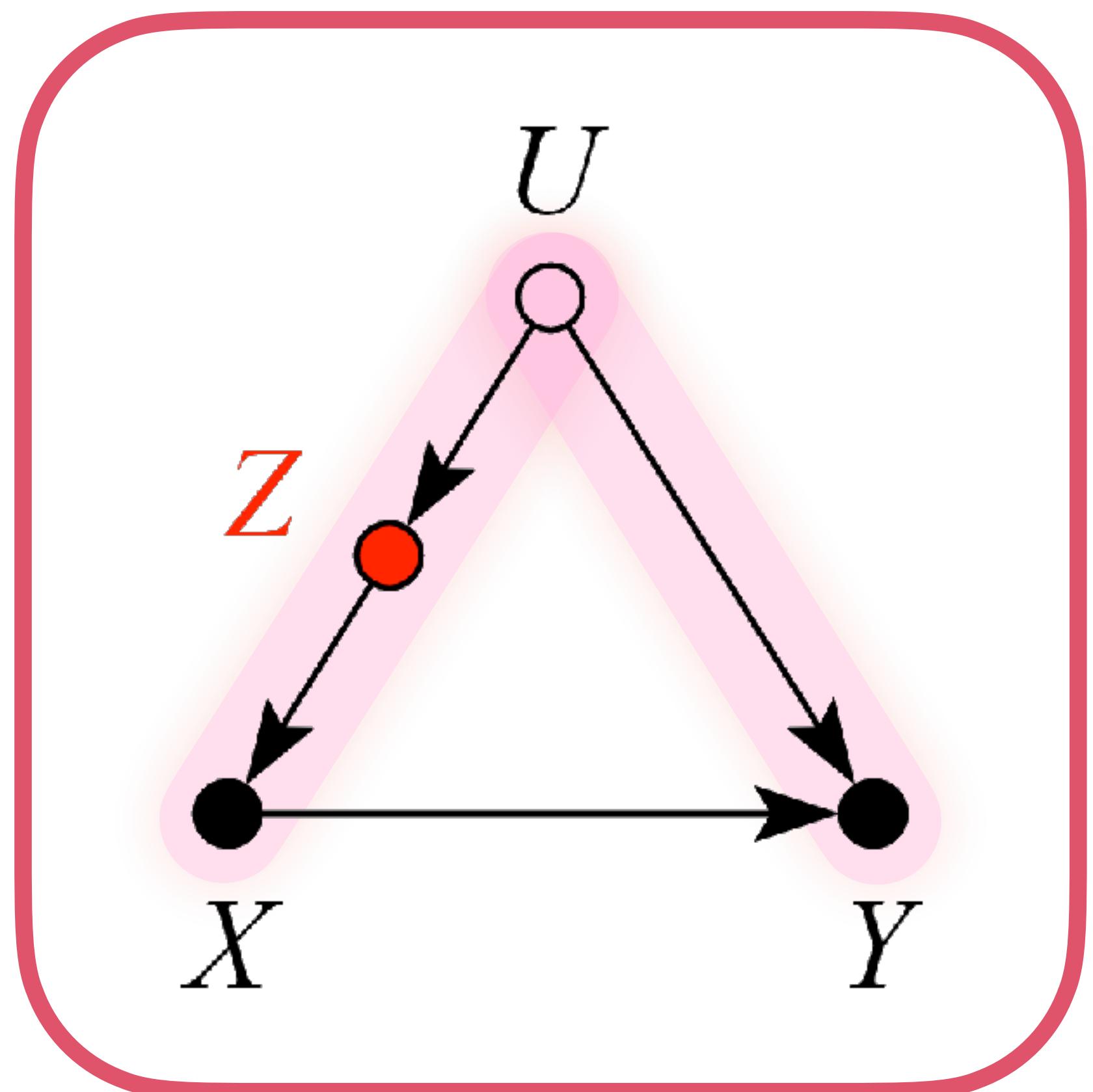
(2) Paths with arrows **entering**  $X$  are backdoor paths (confounding paths)



(3) Find a set of control variables that close/block all backdoor paths

Block the pipe:  $X \perp\!\!\!\perp U | Z$

Z “knows” all of the association between  $X, Y$  that is due to  $U$



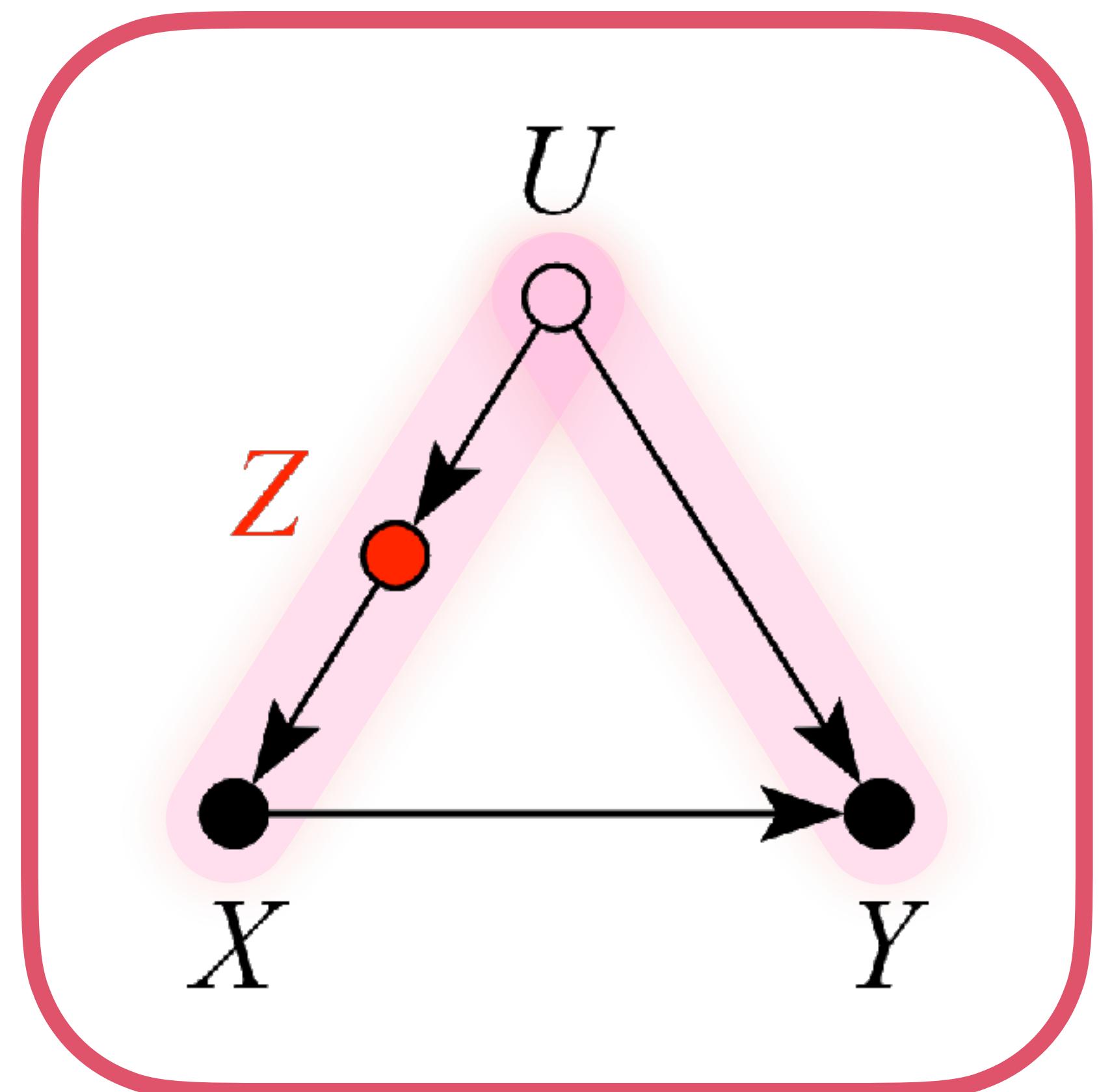
(3) Find a set of control variables that close/block all backdoor paths

Block the pipe:  $X \perp\!\!\!\perp U | Z$

$$P(Y | \text{do}(X)) = \sum_z P(Y | X, Z)P(Z = z)$$

$$Y_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_X X_i + \beta_Z Z_i$$

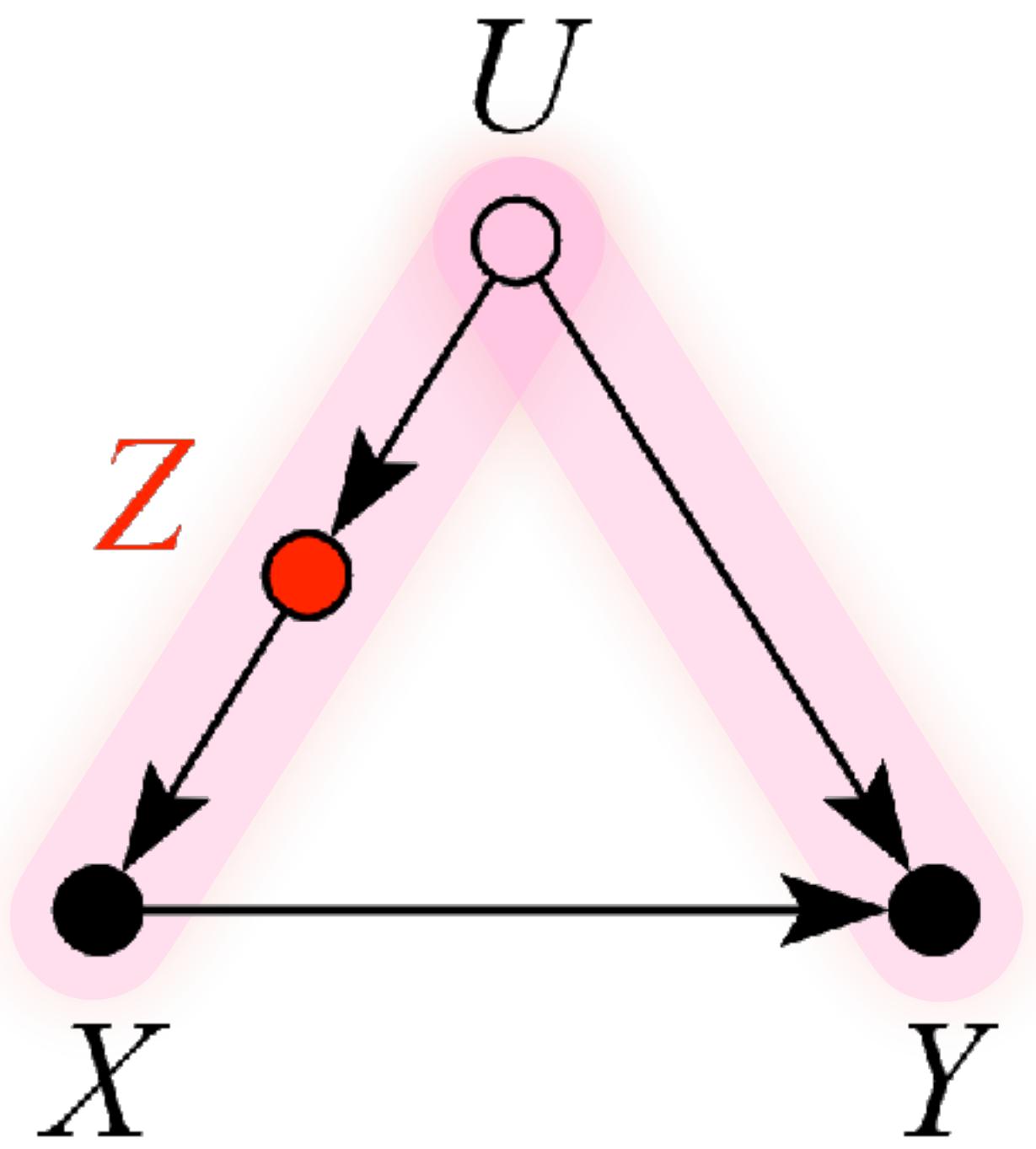


```

# simulate confounded Y
N <- 200
b_XY <- 0
b_UY <- -1
b_UZ <- -1
b_ZX <- 1

set.seed(10)
U <- rbern(N)
Z <- rnorm(N,b_UZ*U)
X <- rnorm(N,b_ZX*Z)
Y <- rnorm(N,b_XY*X+b_UY*U)
d <- list(Y=Y,X=X,Z=Z)

```



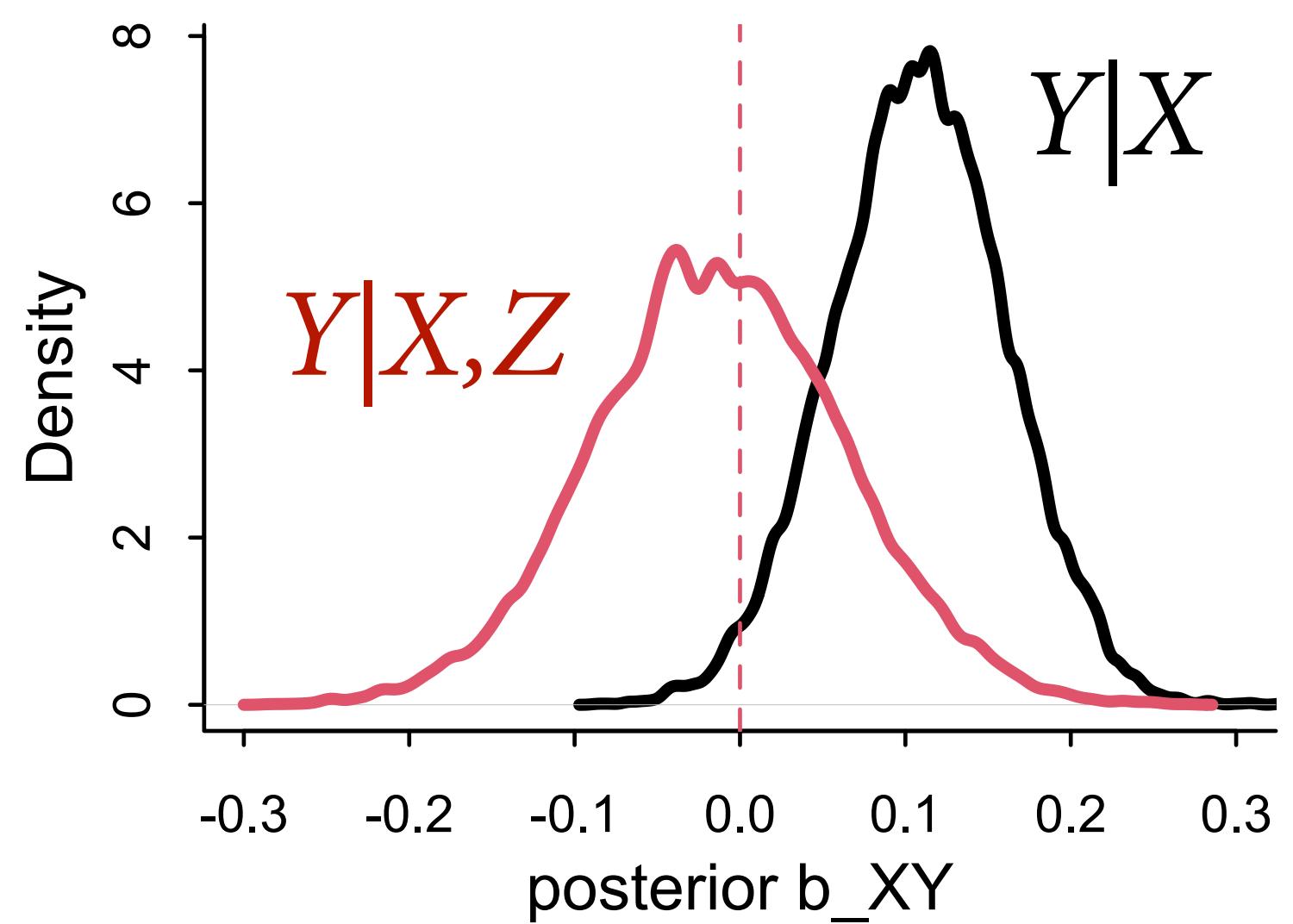
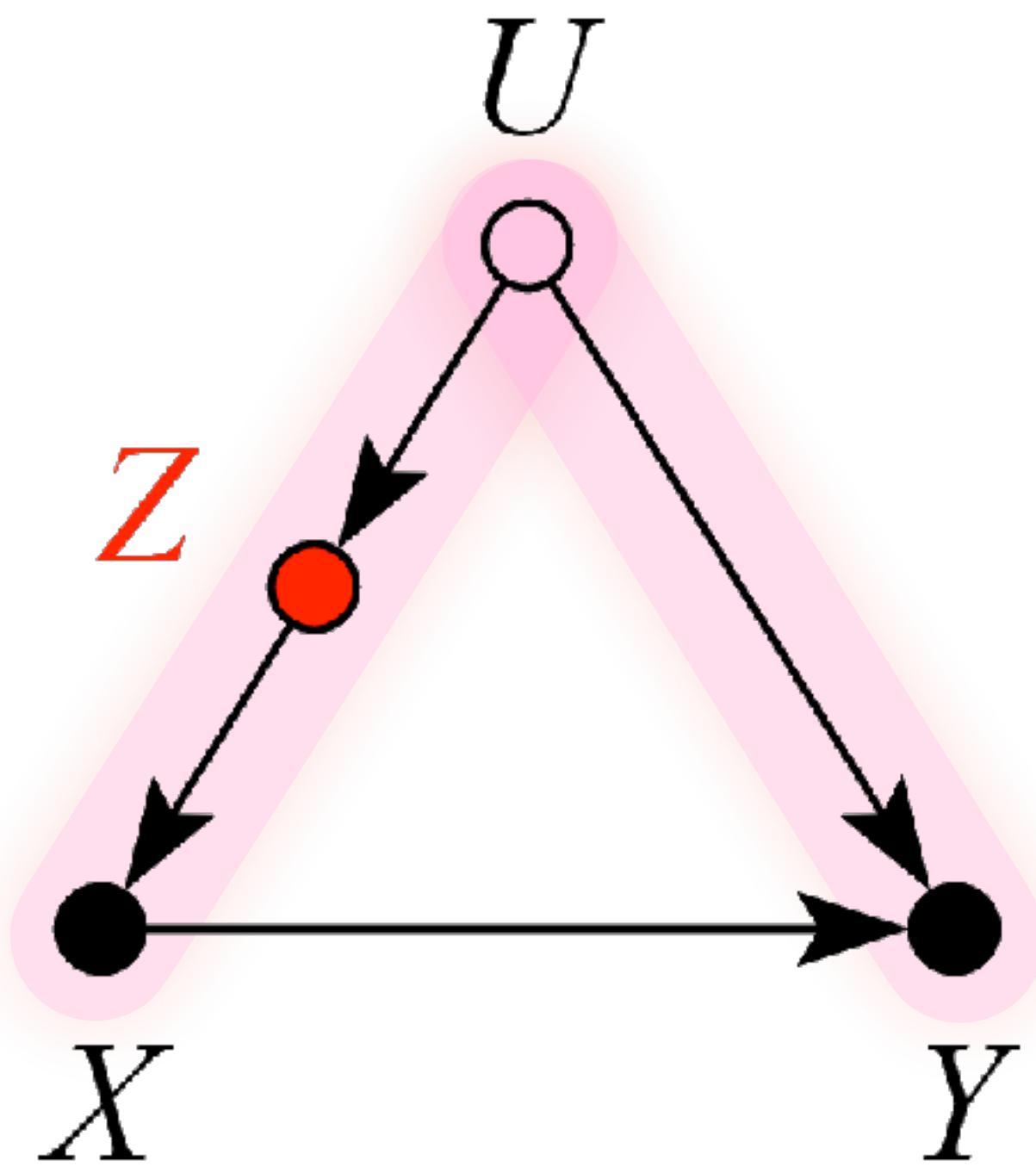
```

# ignore U,Z
m_YX <- quap(
  alist(
    Y ~ dnorm( mu , sigma ),
    mu <- a + b_XY*X,
    a ~ dnorm( 0 , 1 ),
    b_XY ~ dnorm( 0 , 1 ),
    sigma ~ dexp( 1 )
  ), data=d )

# stratify by Z
m_YXZ <- quap(
  alist(
    Y ~ dnorm( mu , sigma ),
    mu <- a + b_XY*X + b_Z*Z,
    a ~ dnorm( 0 , 1 ),
    c(b_XY,b_Z) ~ dnorm( 0 , 1 ),
    sigma ~ dexp( 1 )
  ), data=d )

post <- extract.samples(m_YX)
post2 <- extract.samples(m_YXZ)
dens(post$b_XY, lwd=3, col=1, xlab="posterior
b_XY", xlim=c(-0.3,0.3))
dens(post2$b_XY, lwd=3, col=2, add=TRUE)

```



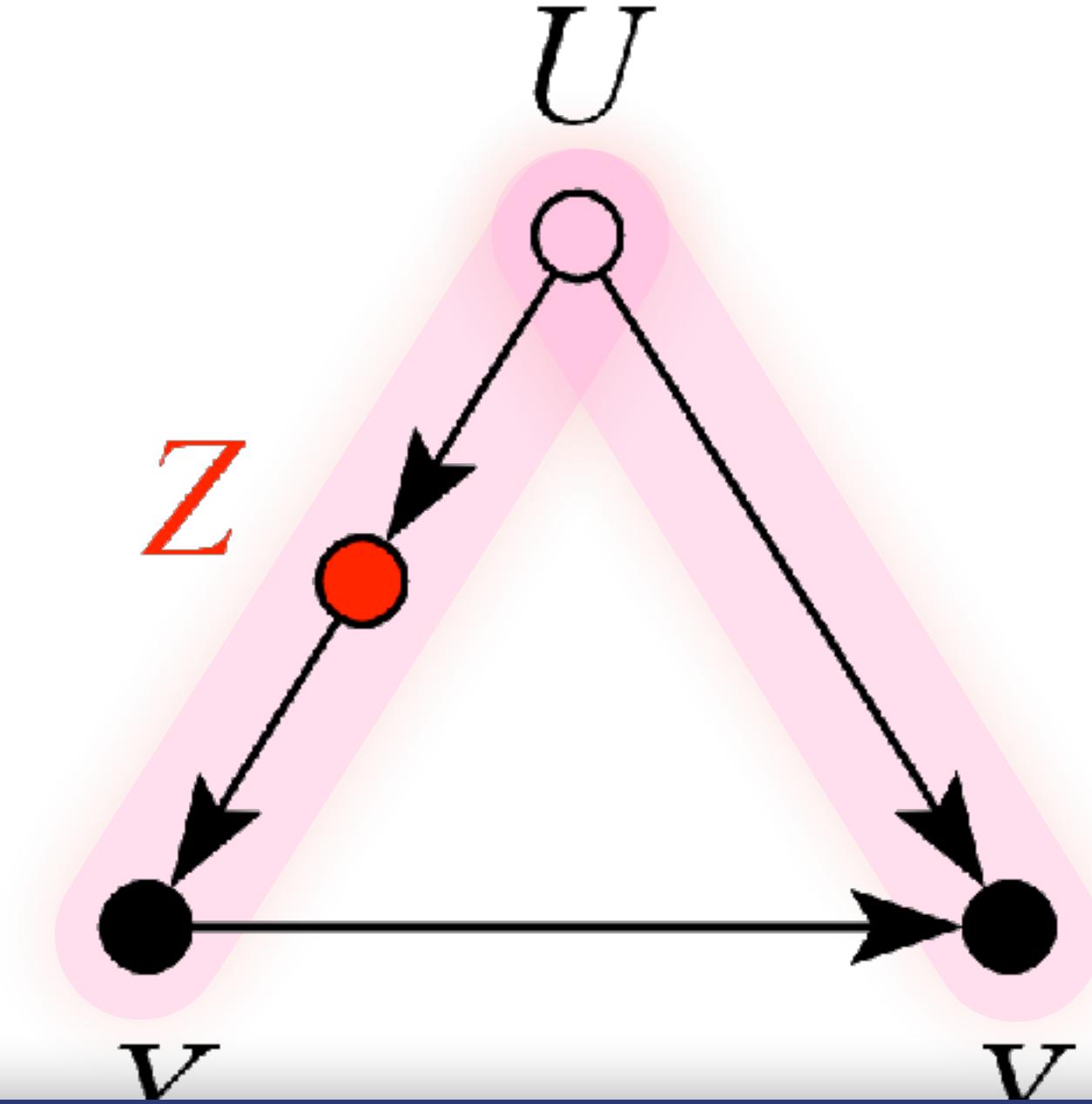
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```



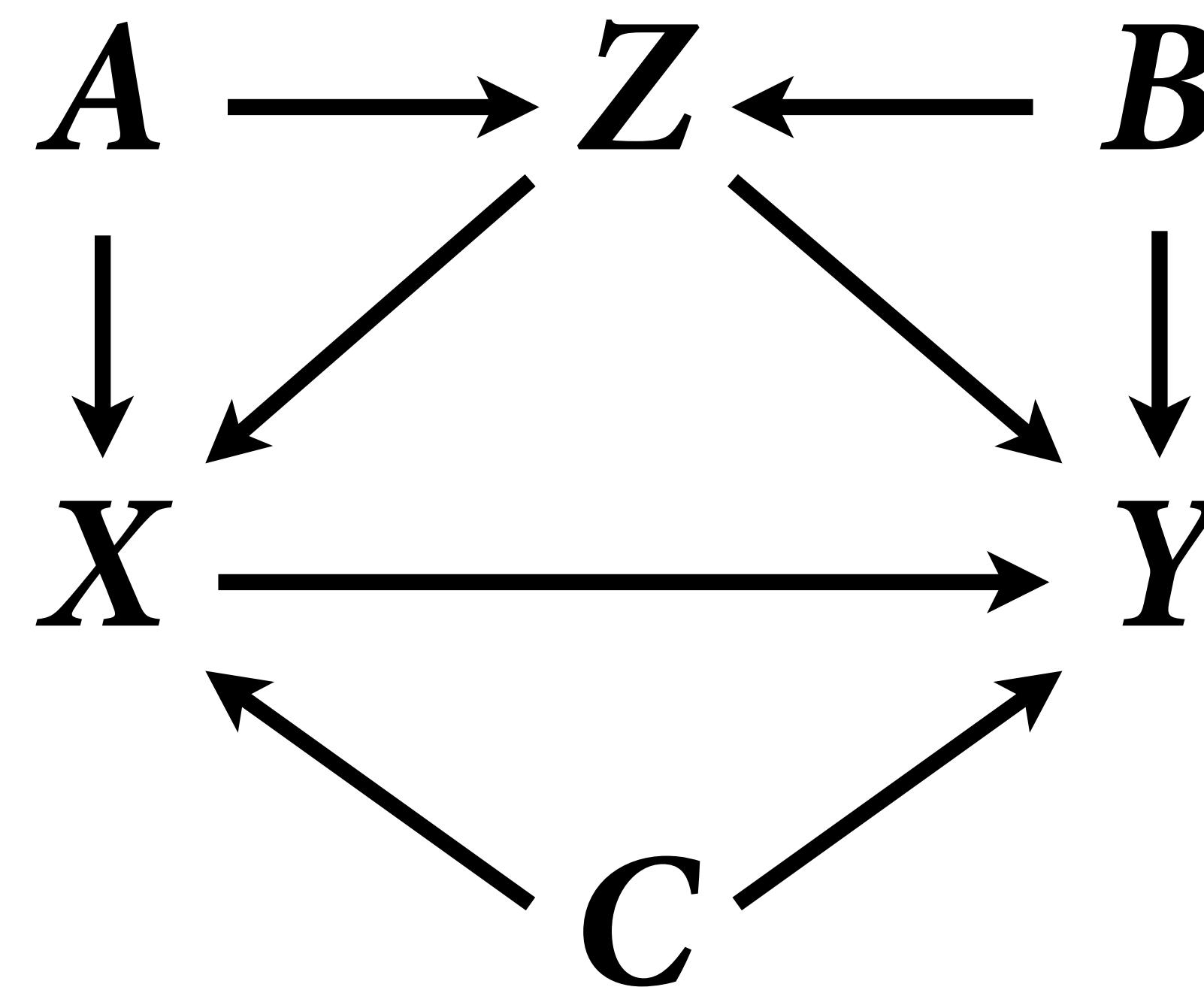
```

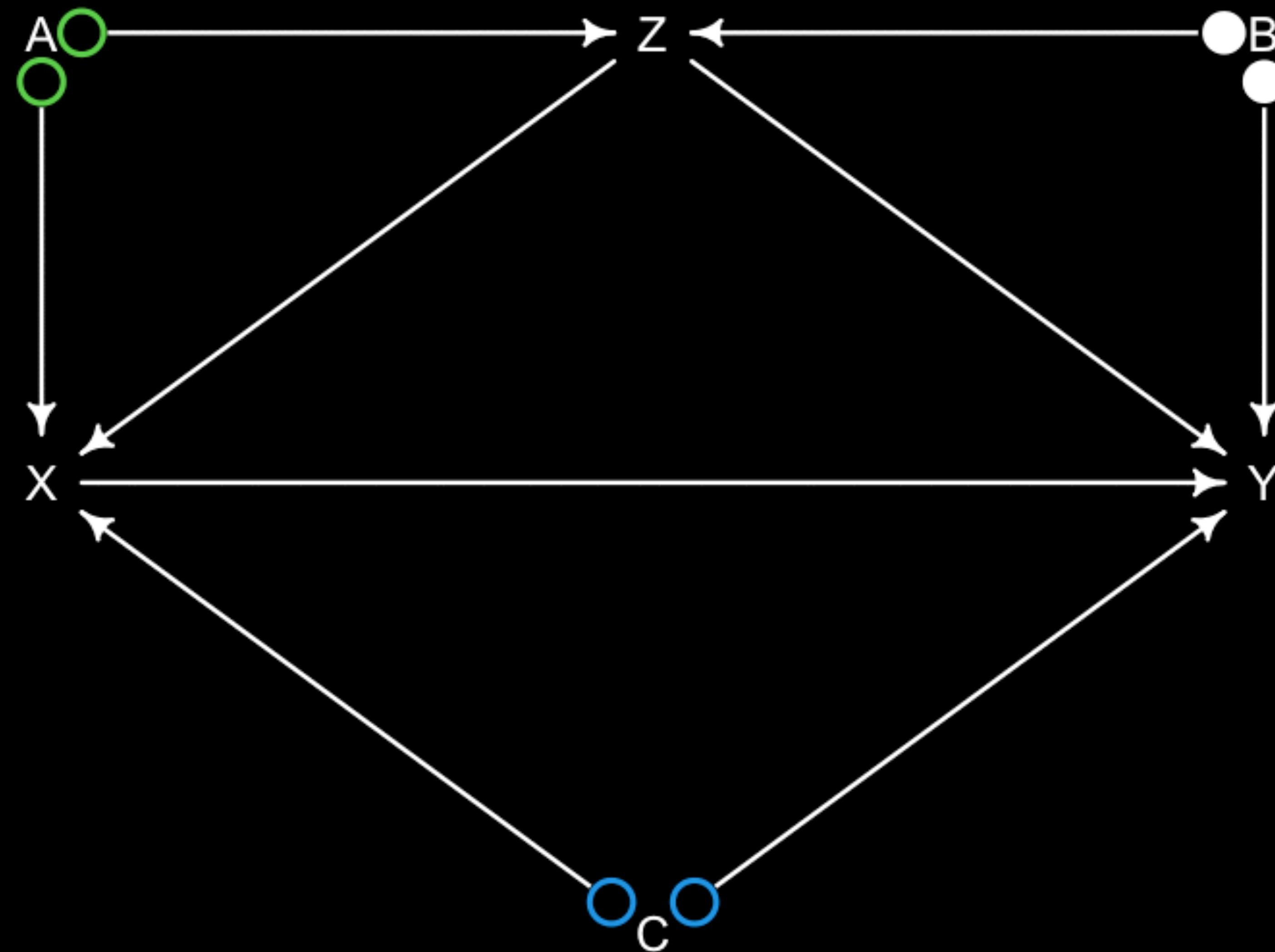
> precis(m_YXZ)
      mean   sd  5.5% 94.5%
a     -0.32 0.09 -0.47 -0.18
b_XY  -0.01 0.08 -0.13  0.11
b_Z    0.24 0.11  0.06  0.42
sigma  1.18 0.06  1.08  1.27

```

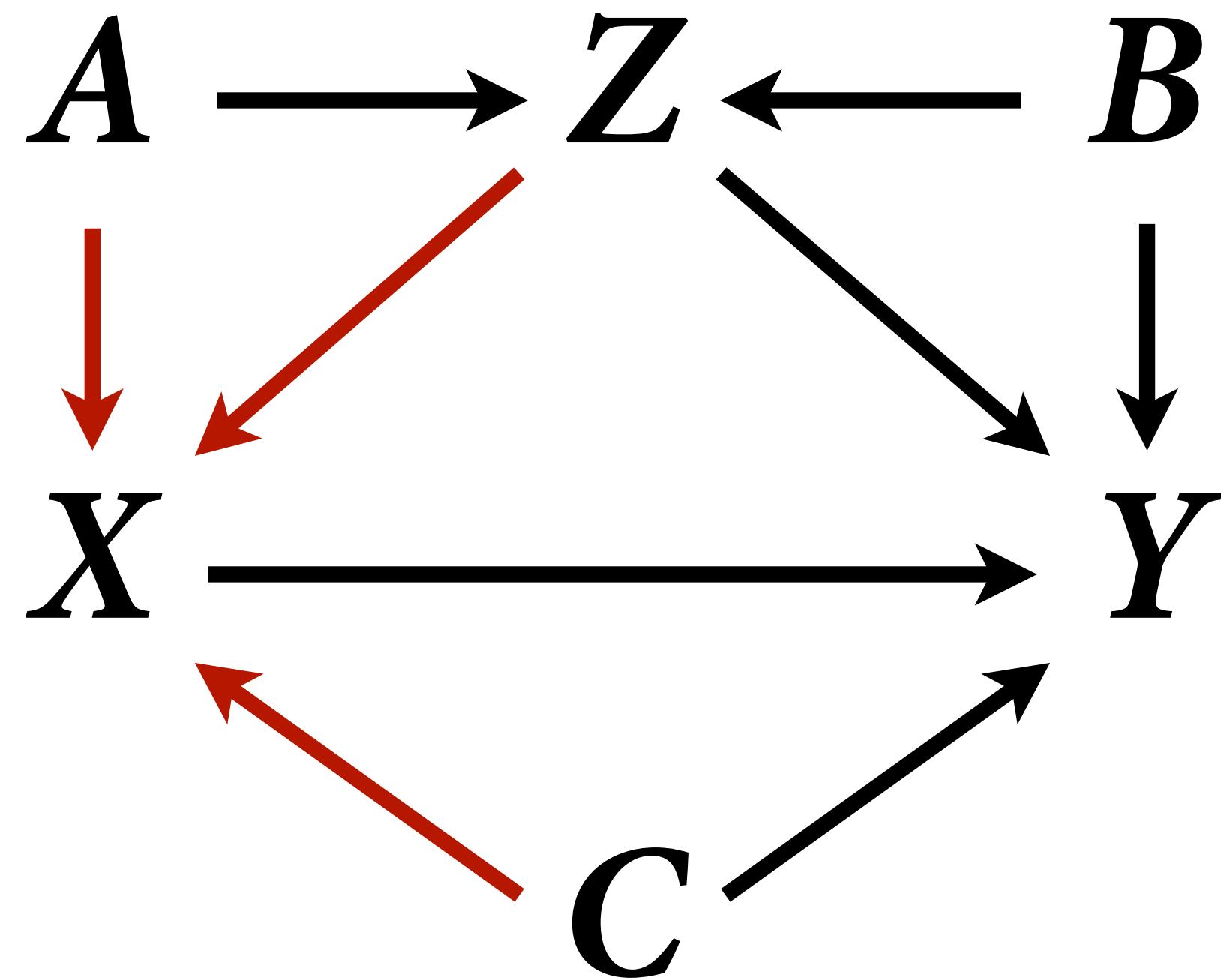
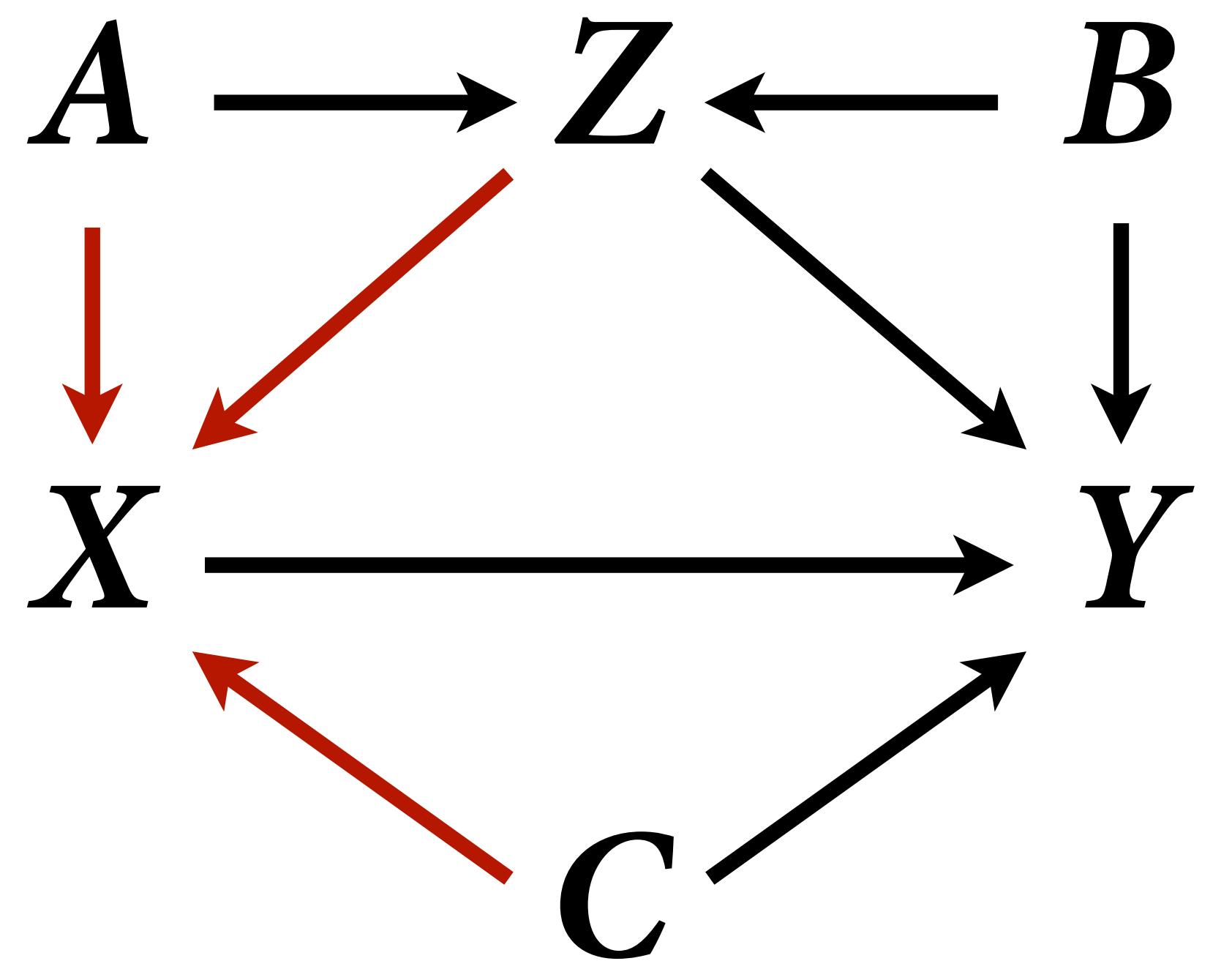
Coefficient on Z means nothing. “Table 2 Fallacy”

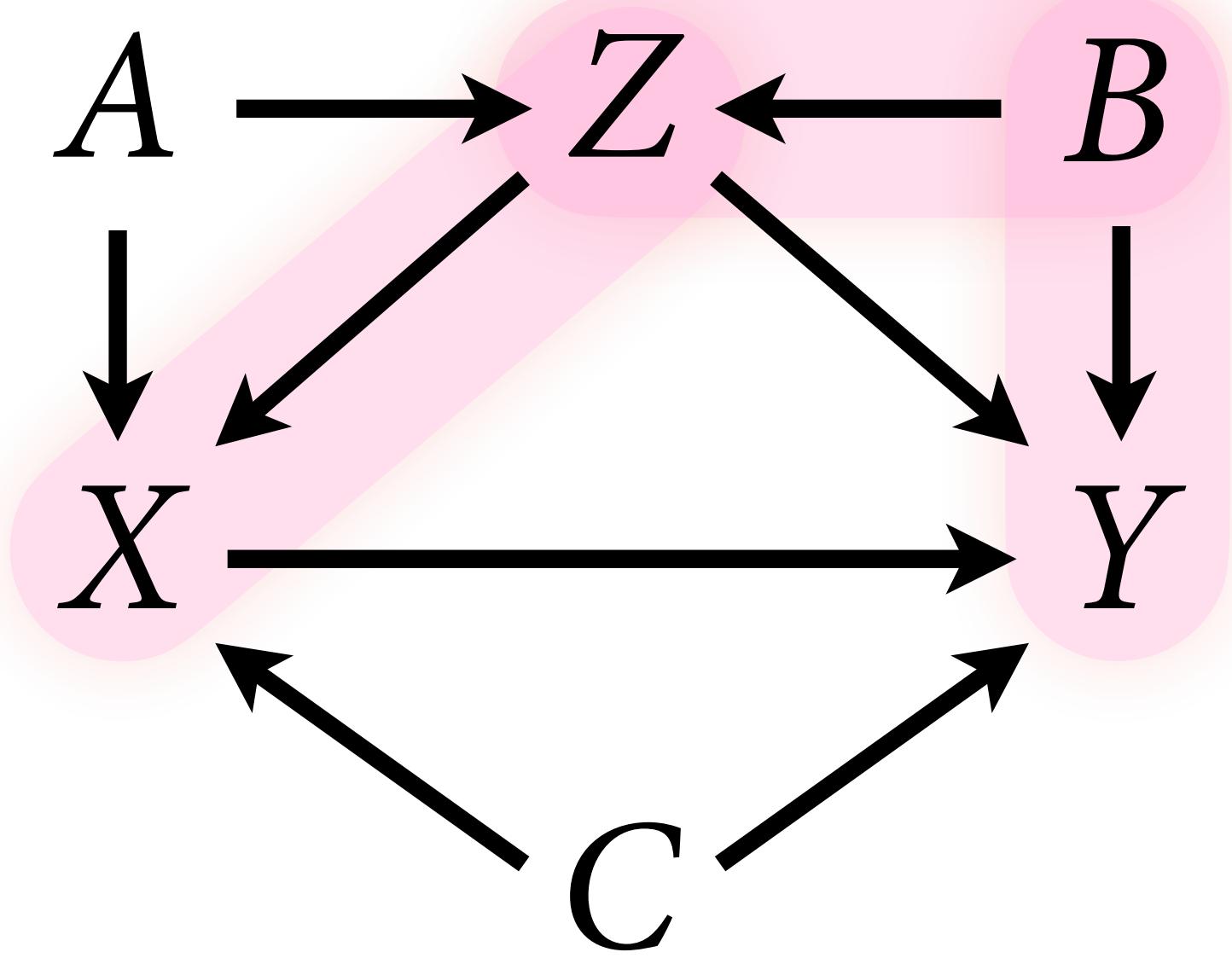
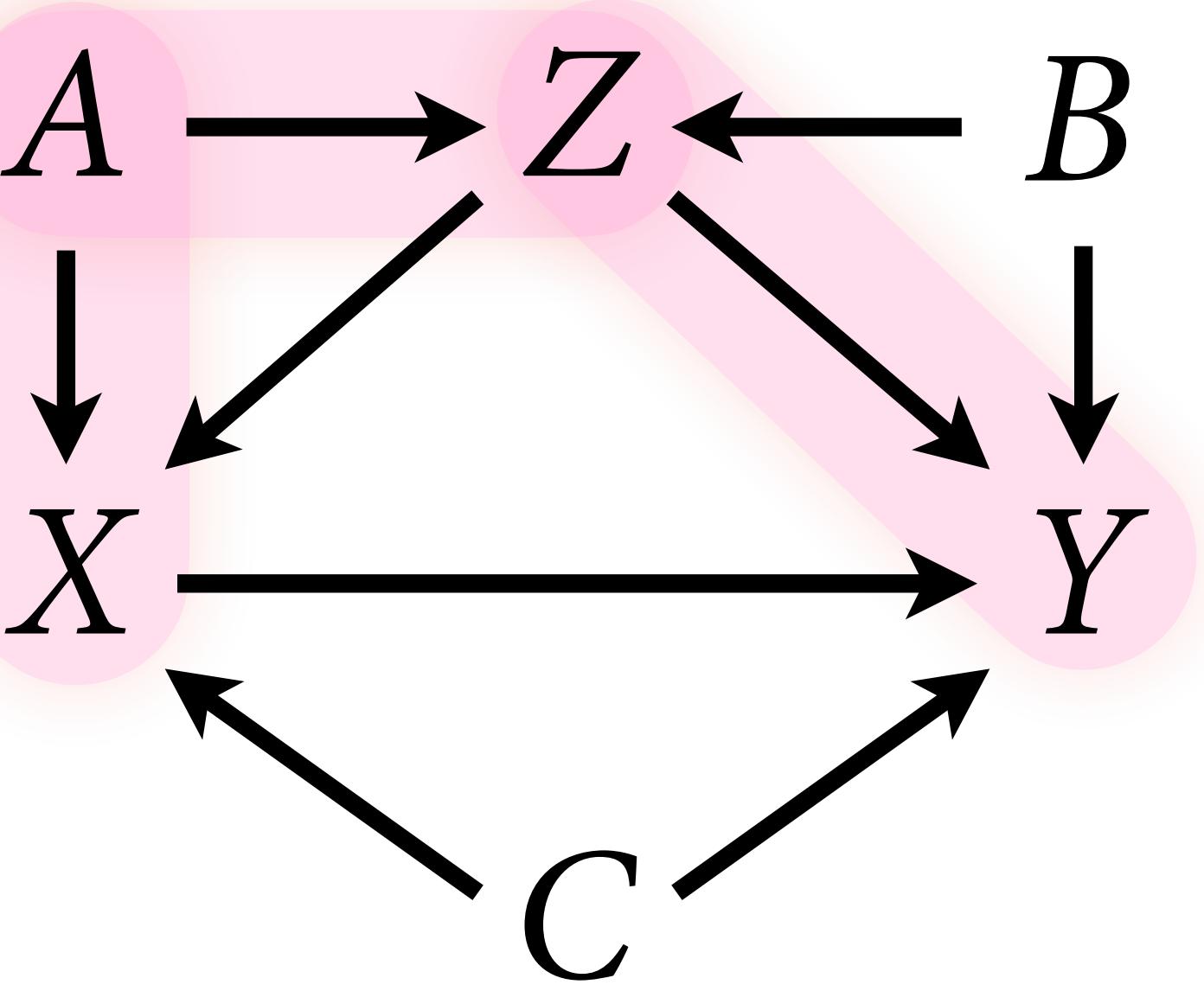
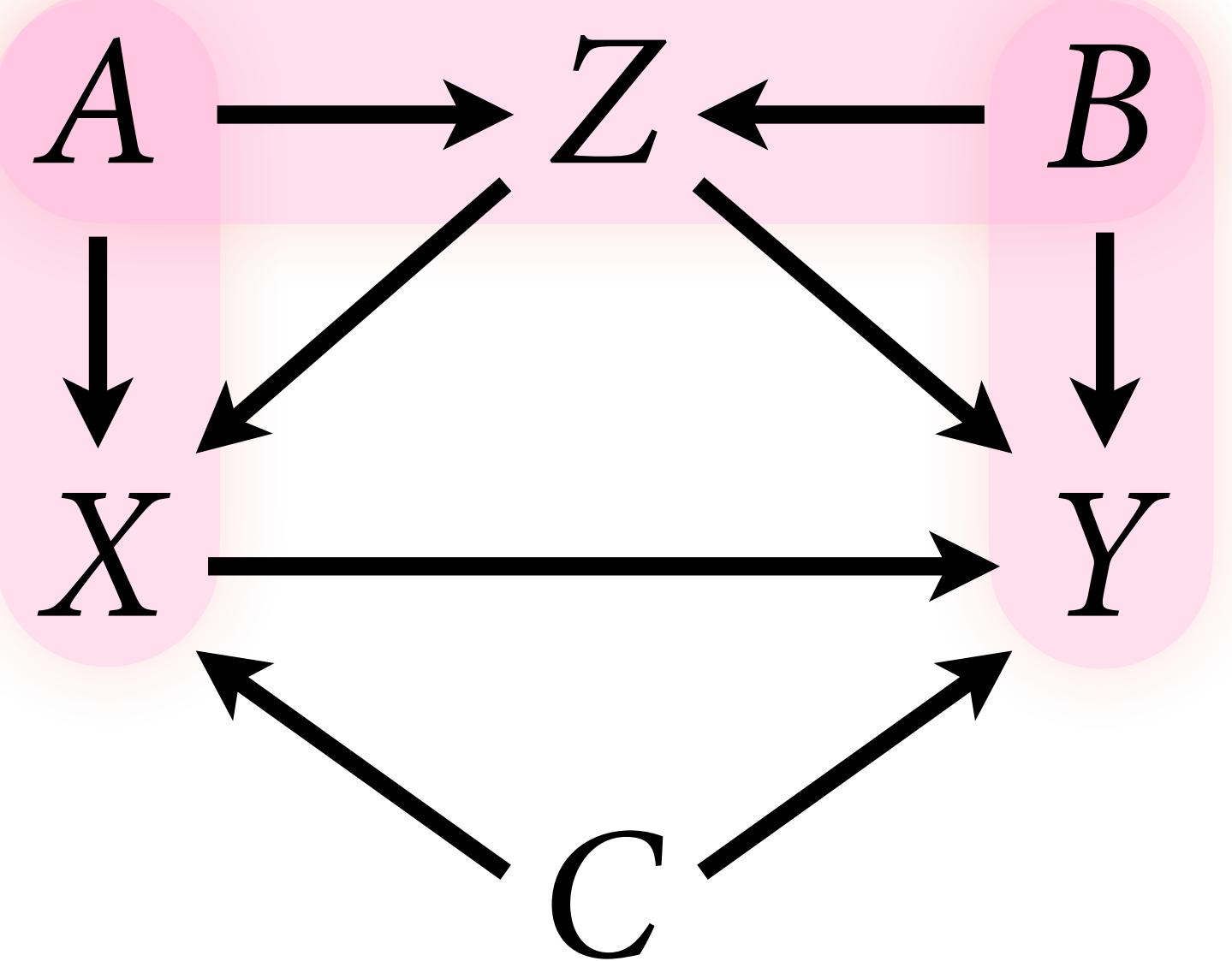
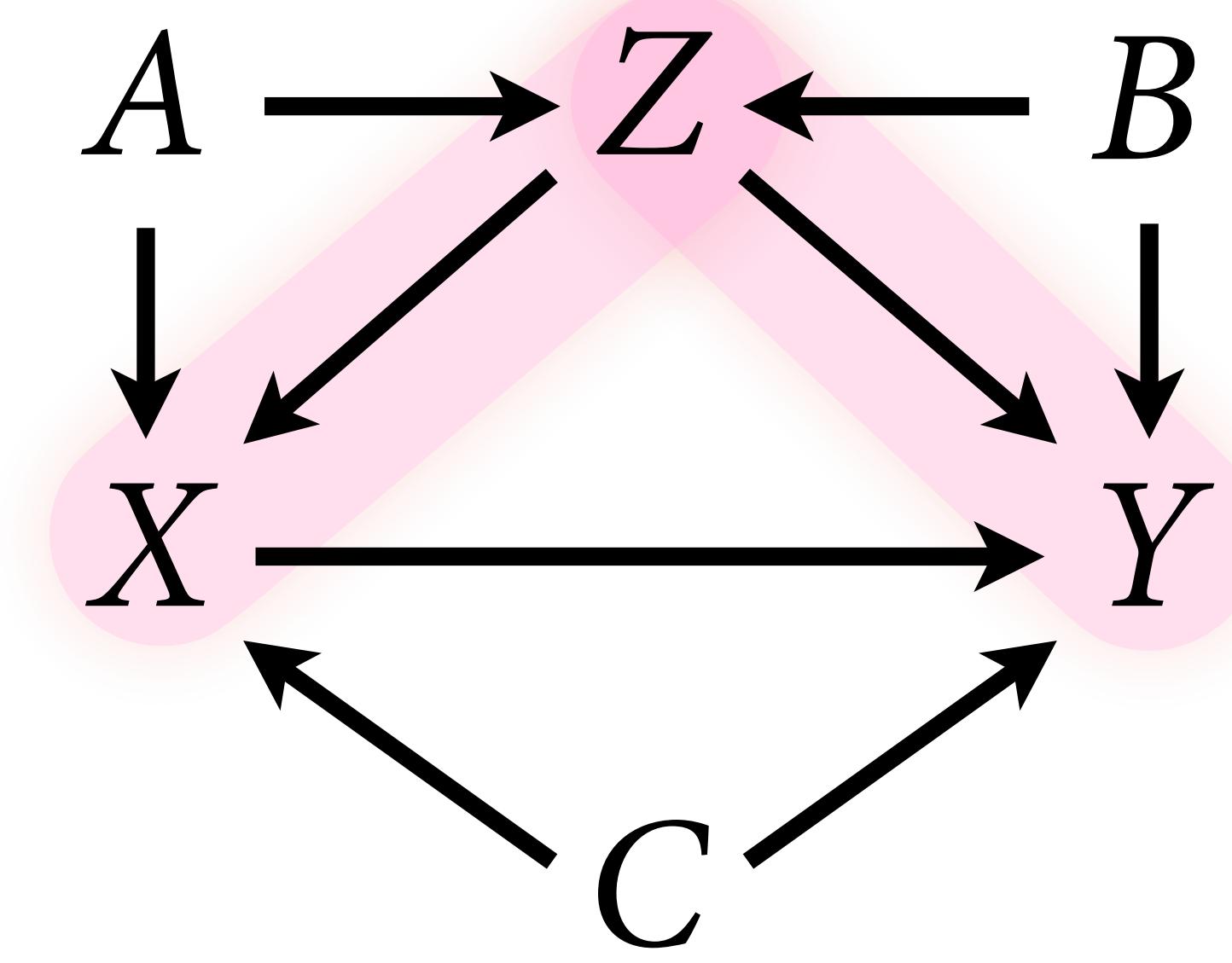
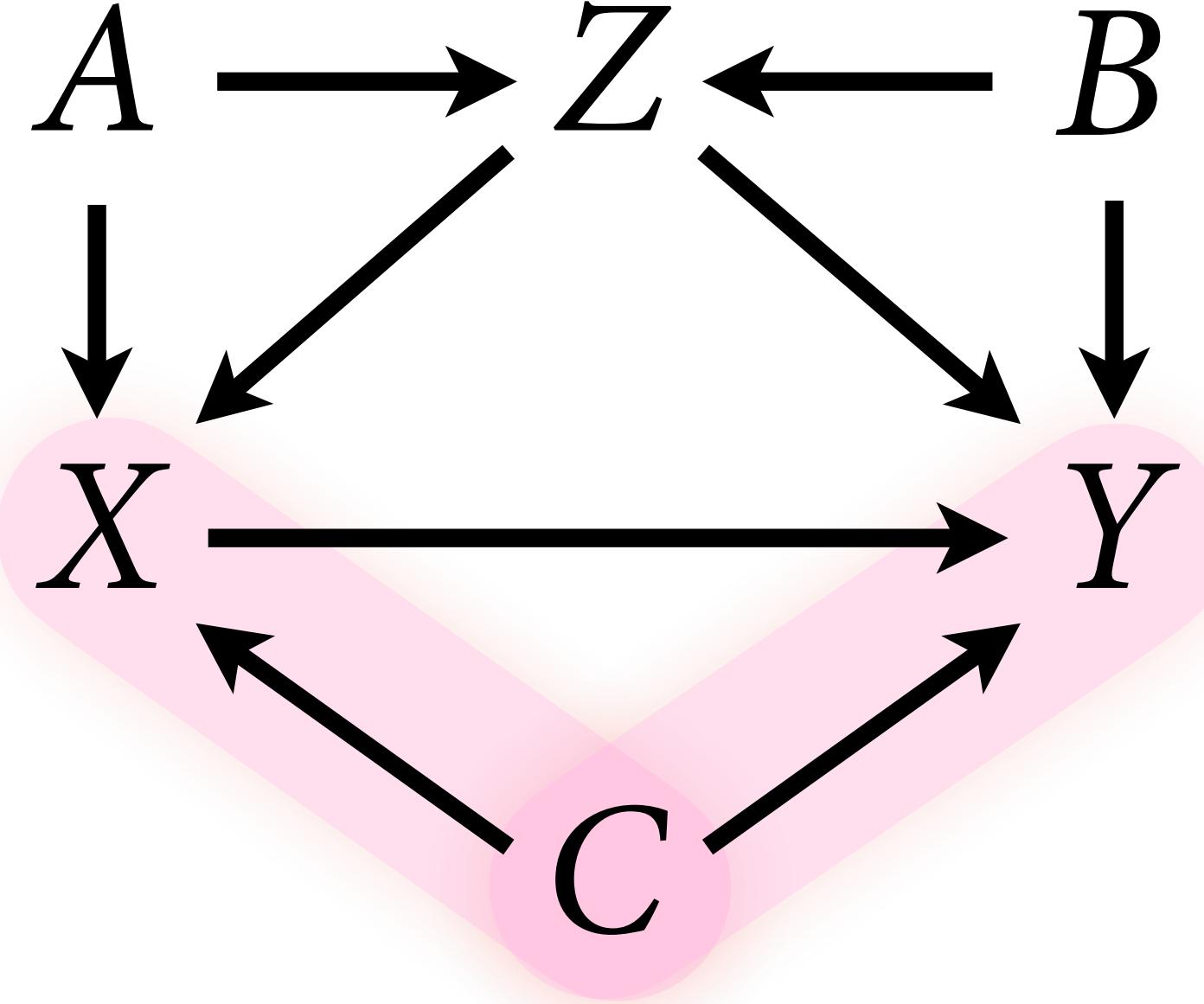
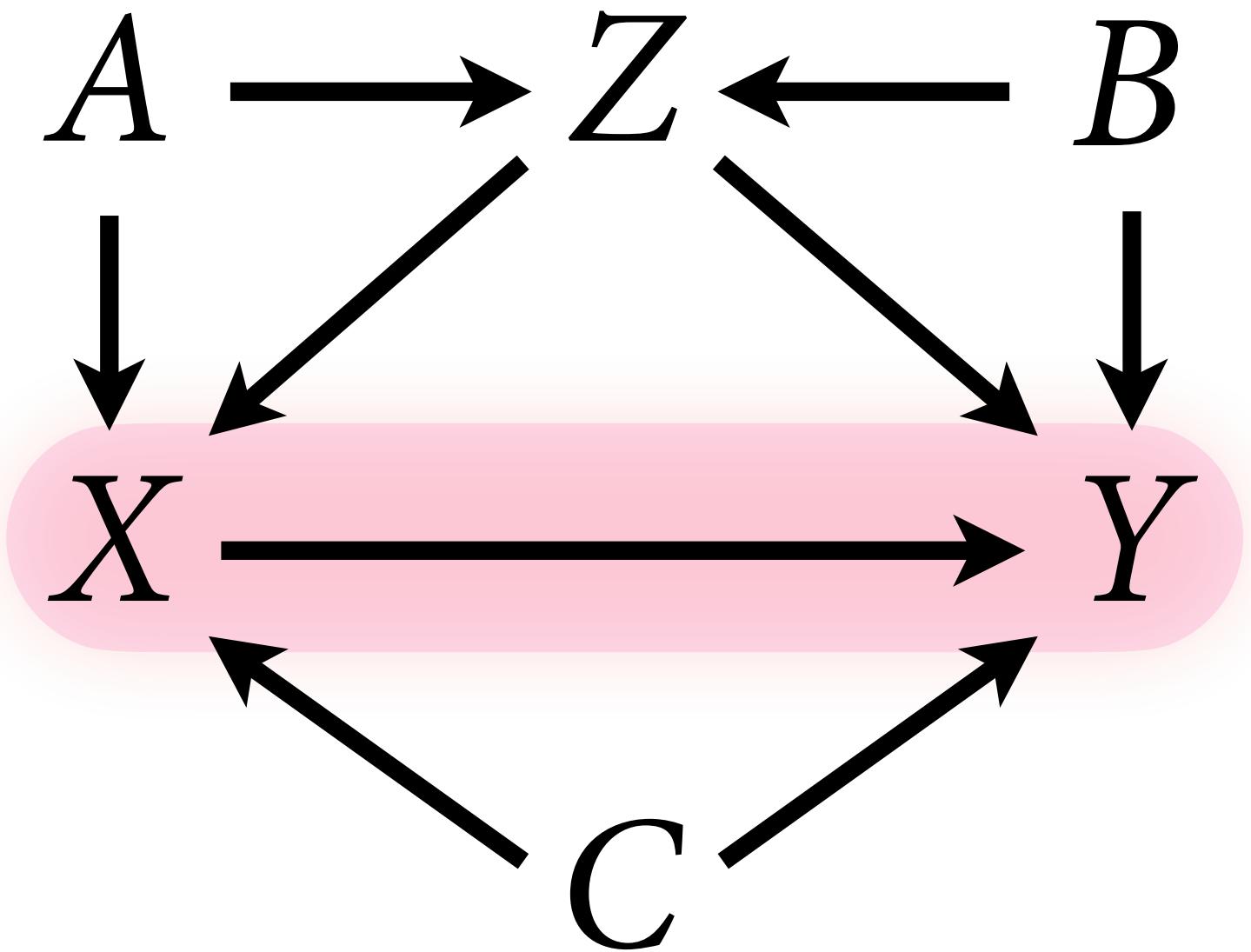
List all the paths connecting X and Y.  
Which need to be closed to estimate  
effect of X on Y?

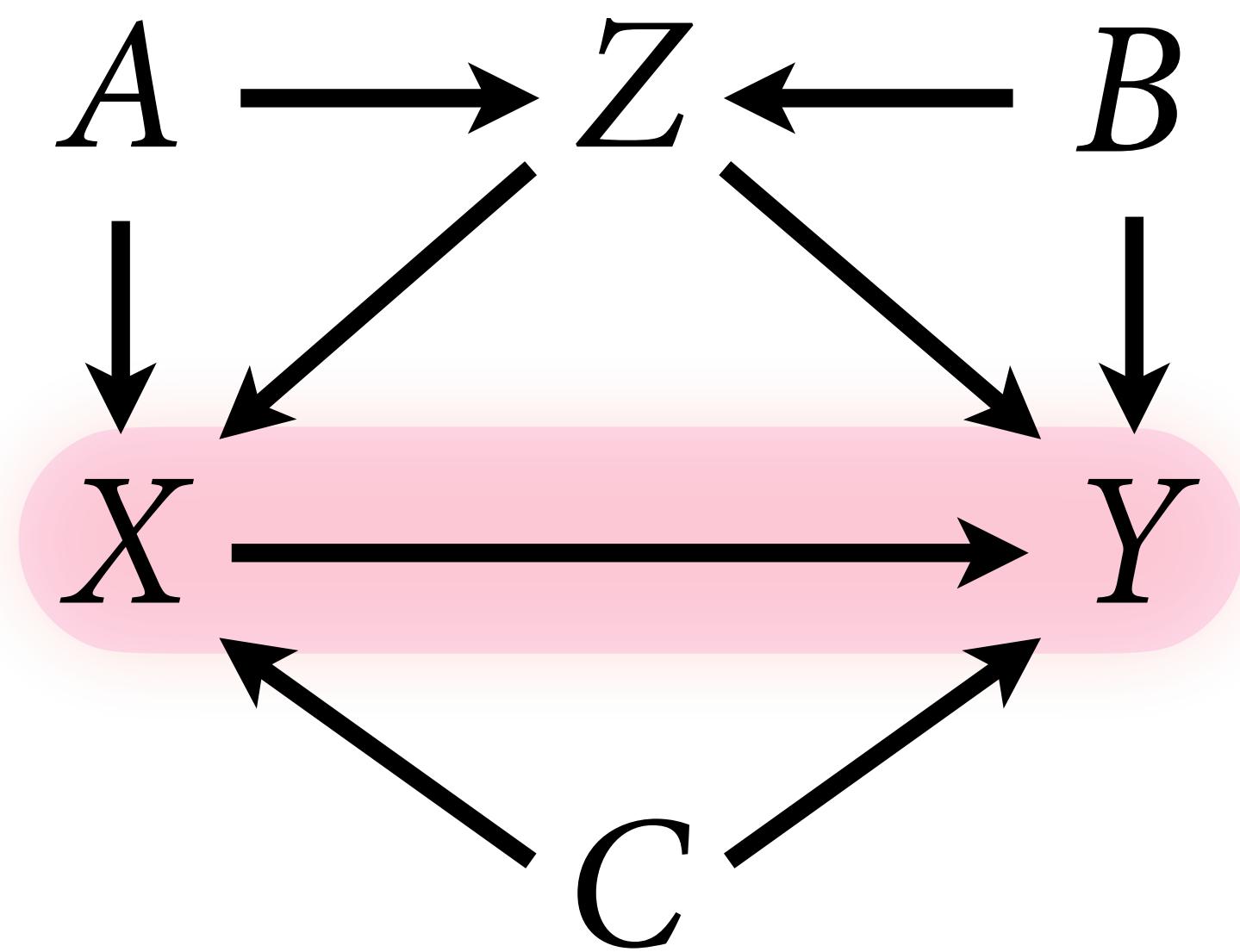




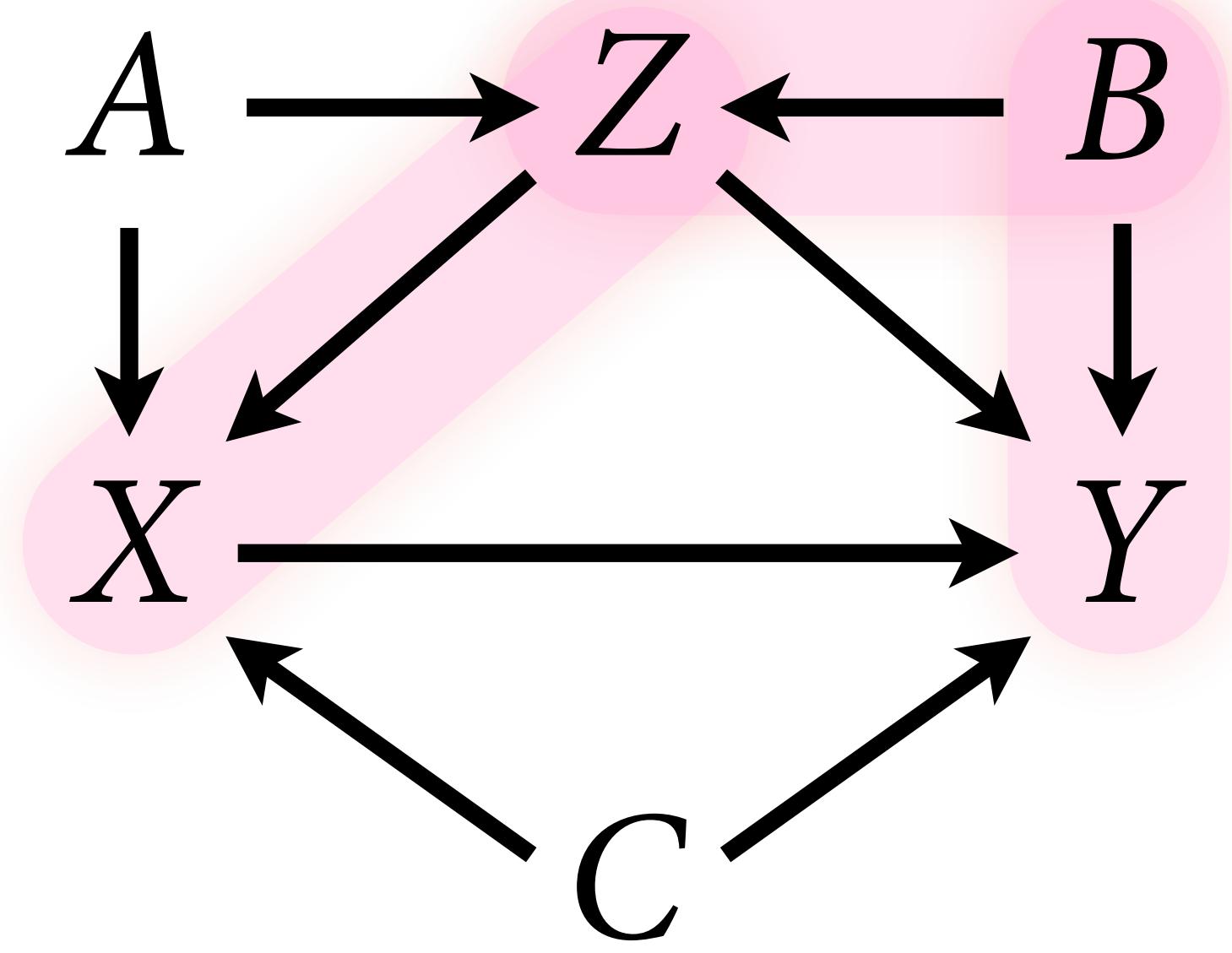
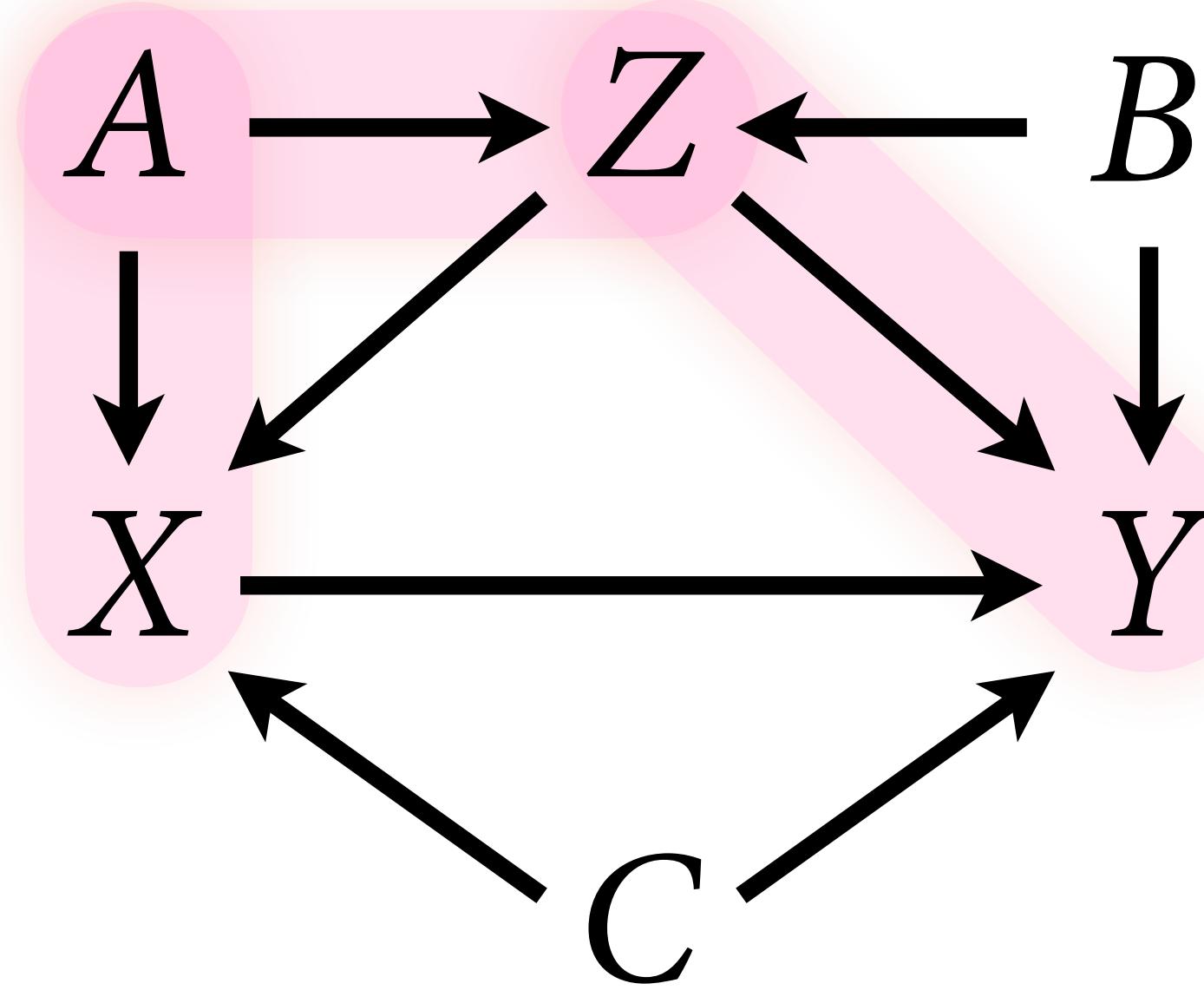
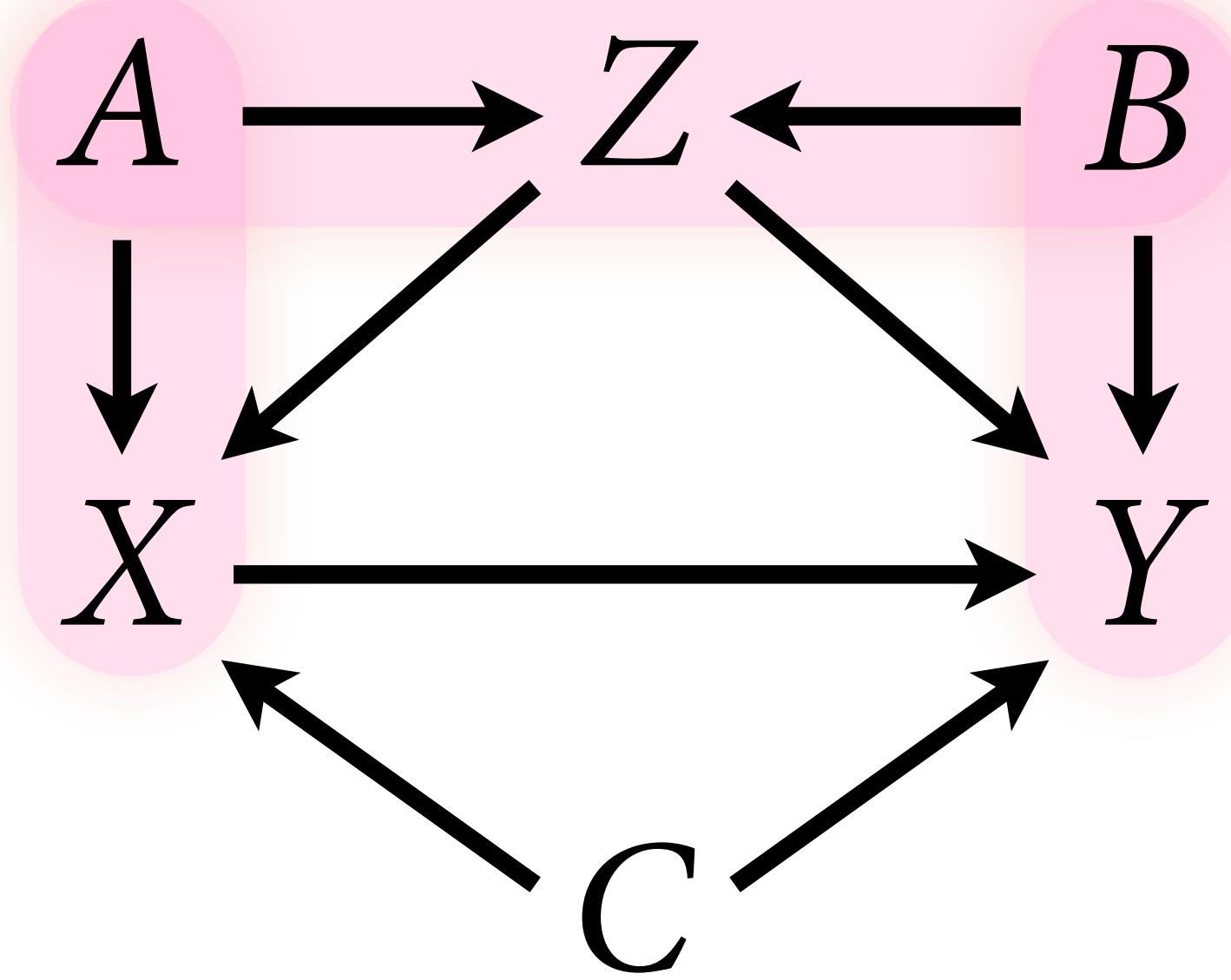
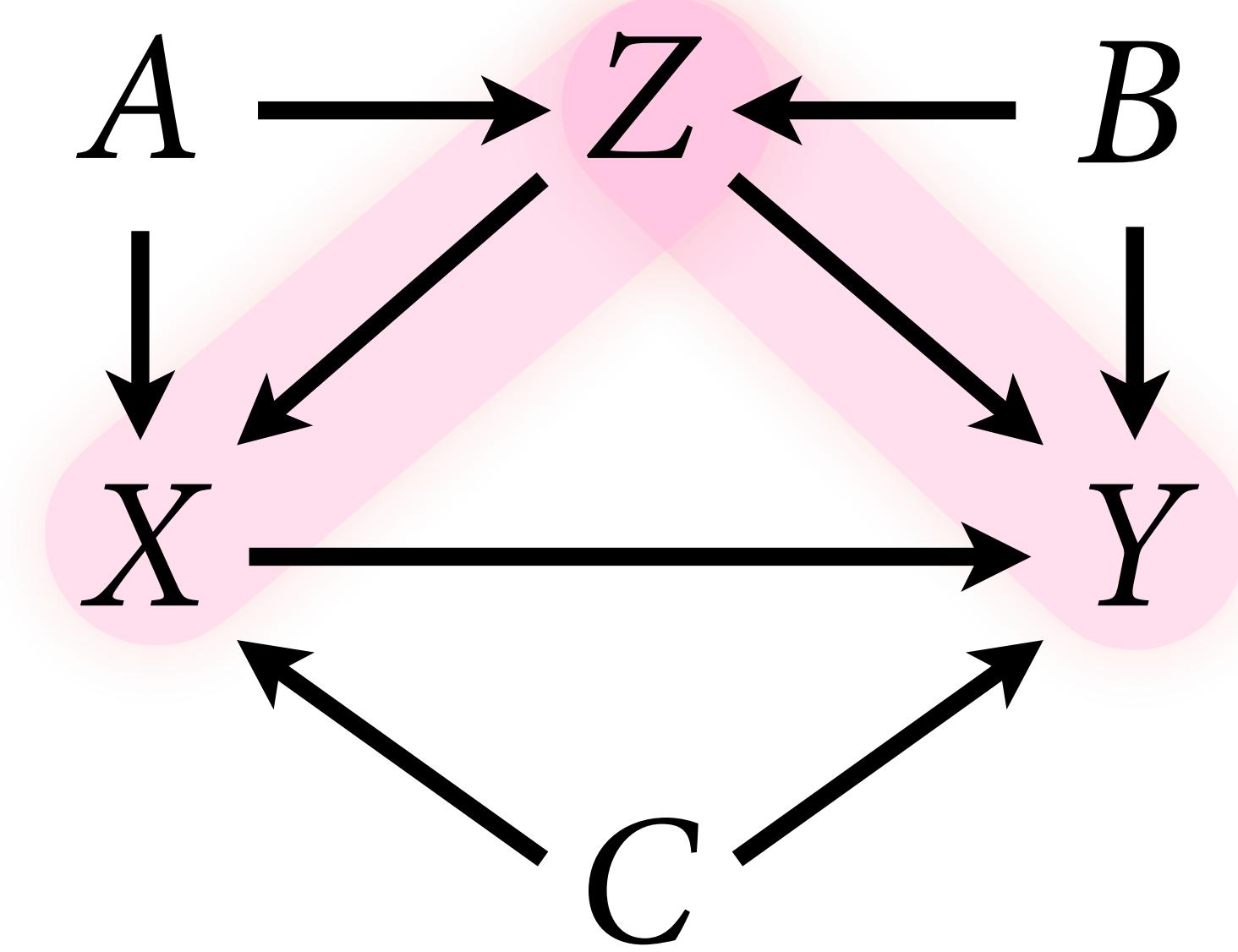
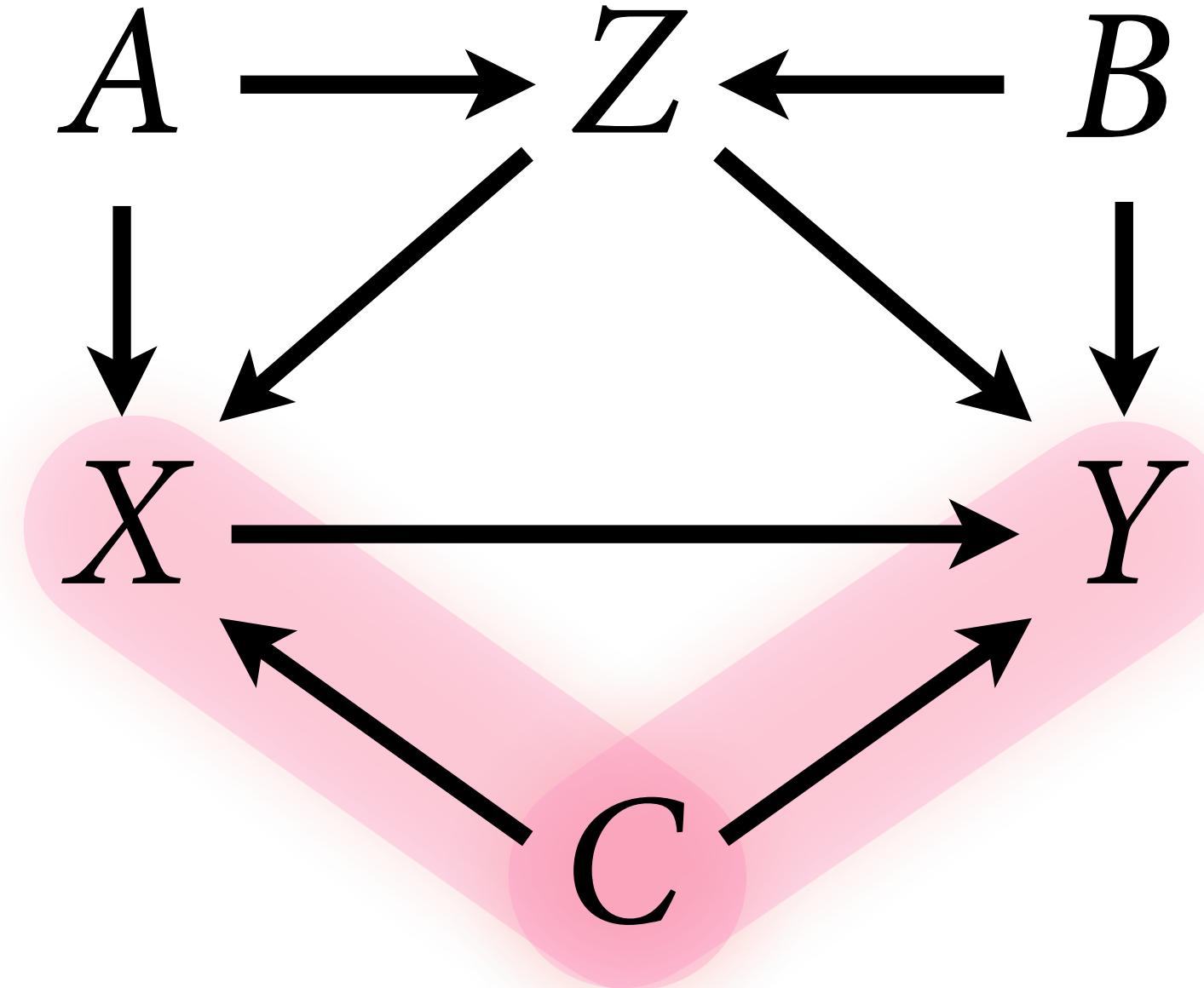
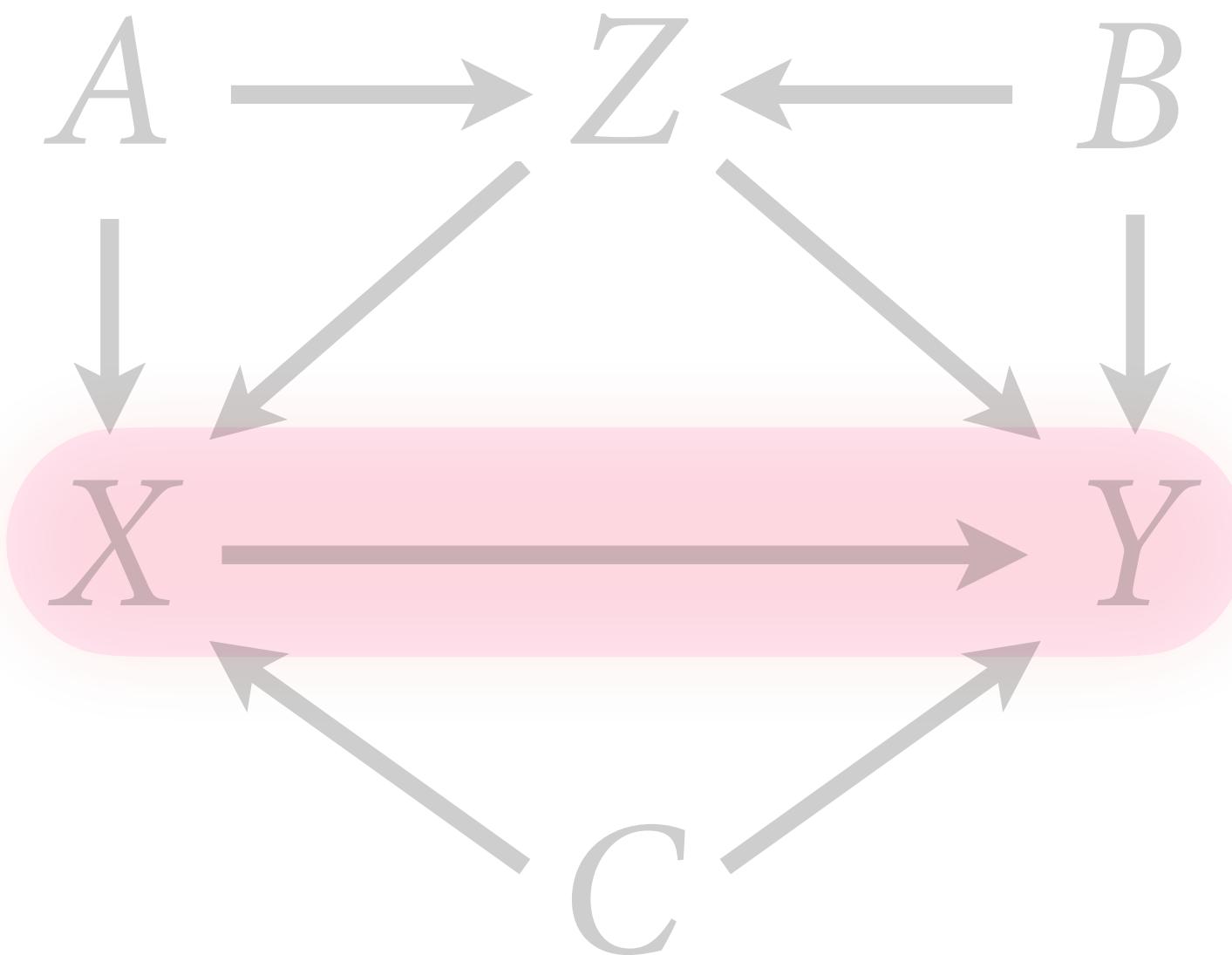
$$P(Y|\text{do}(X))$$

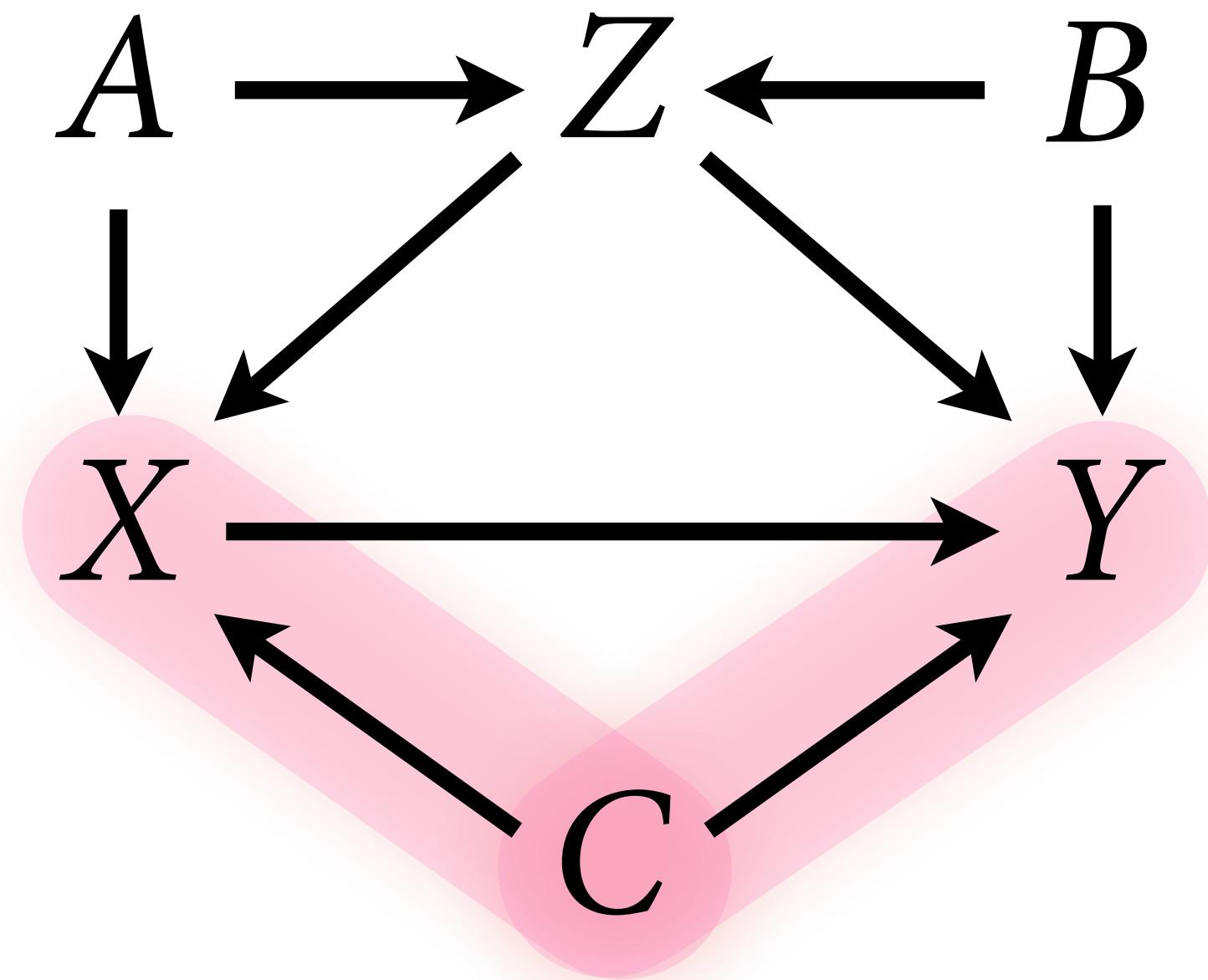




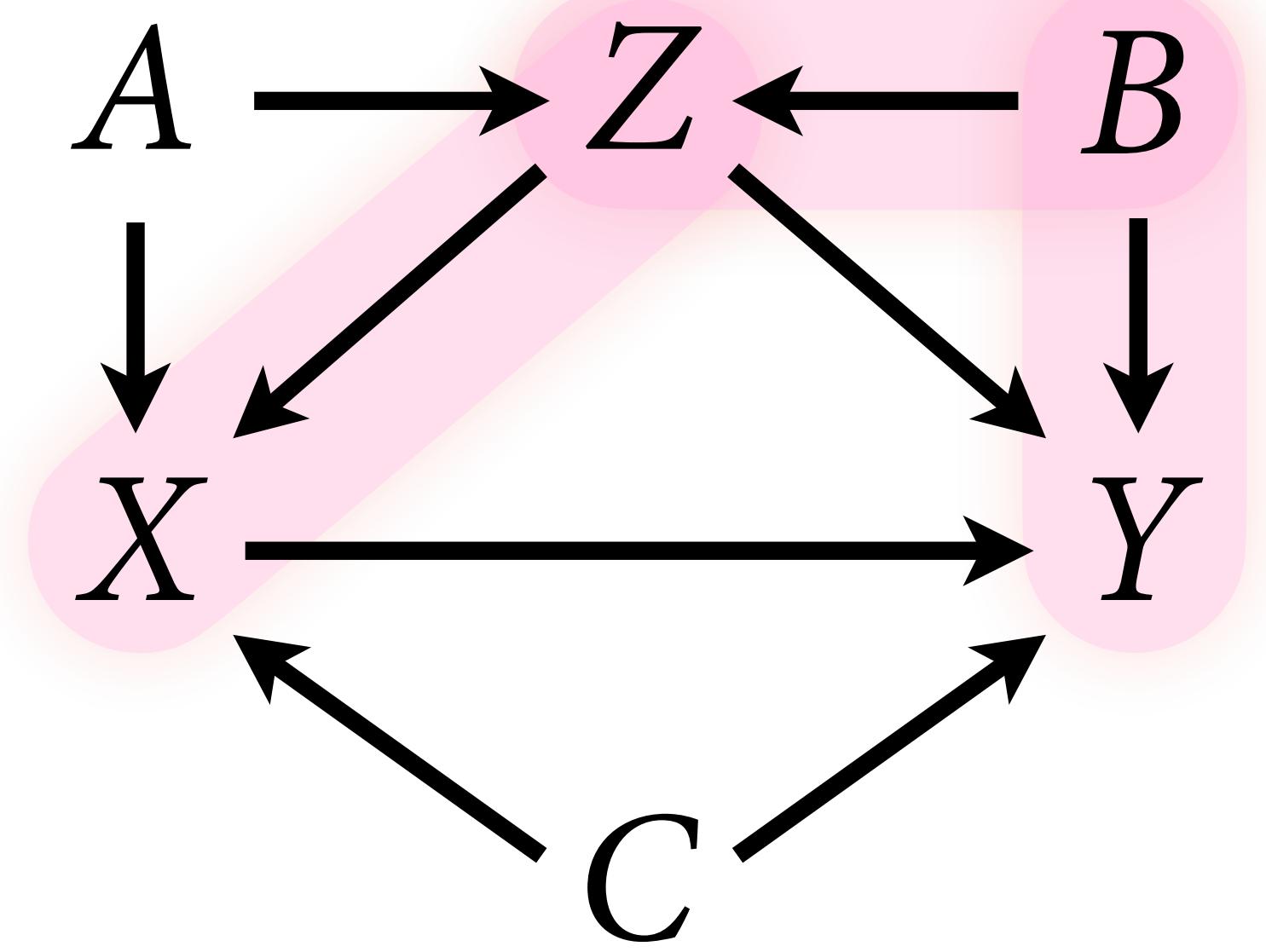
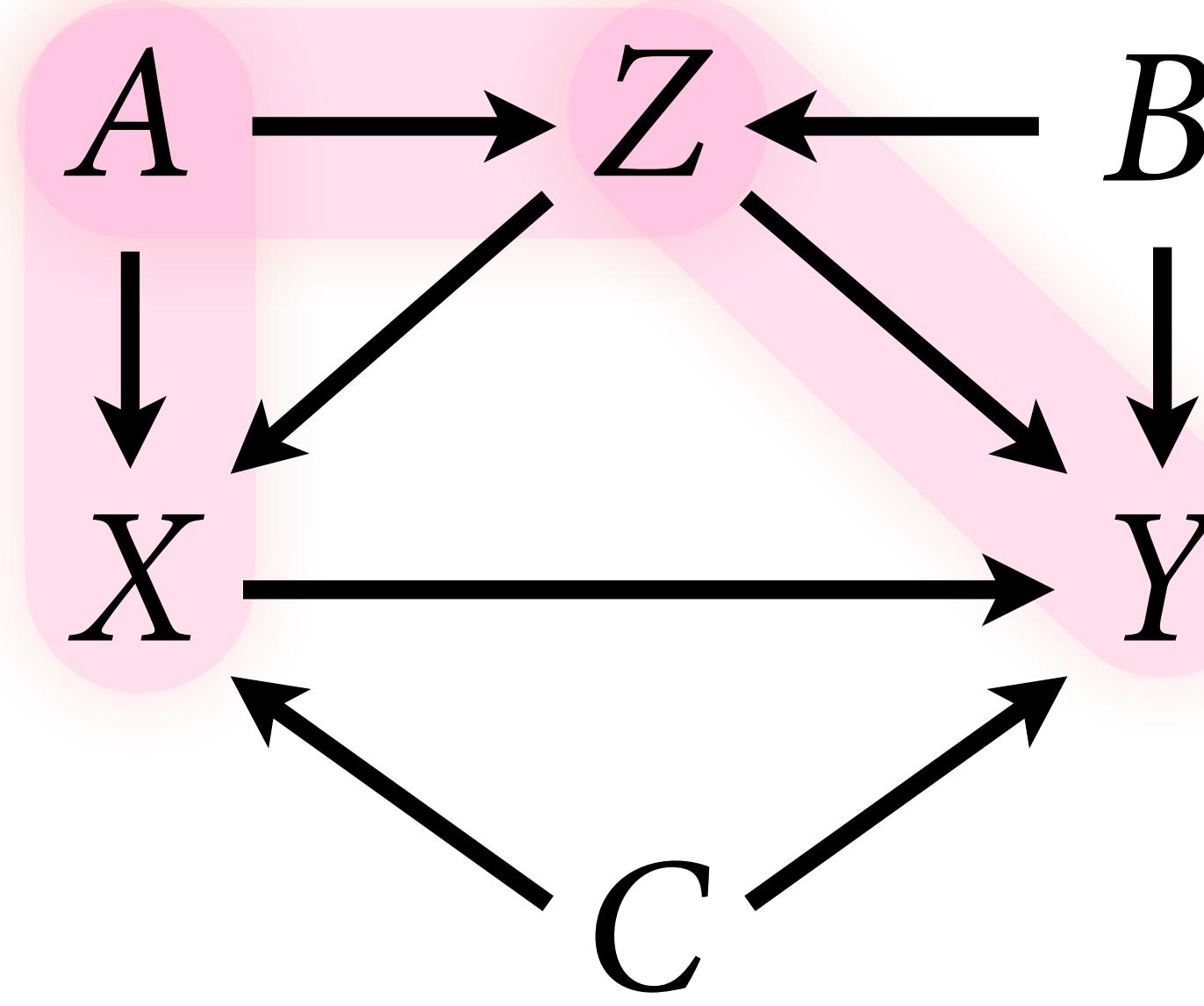
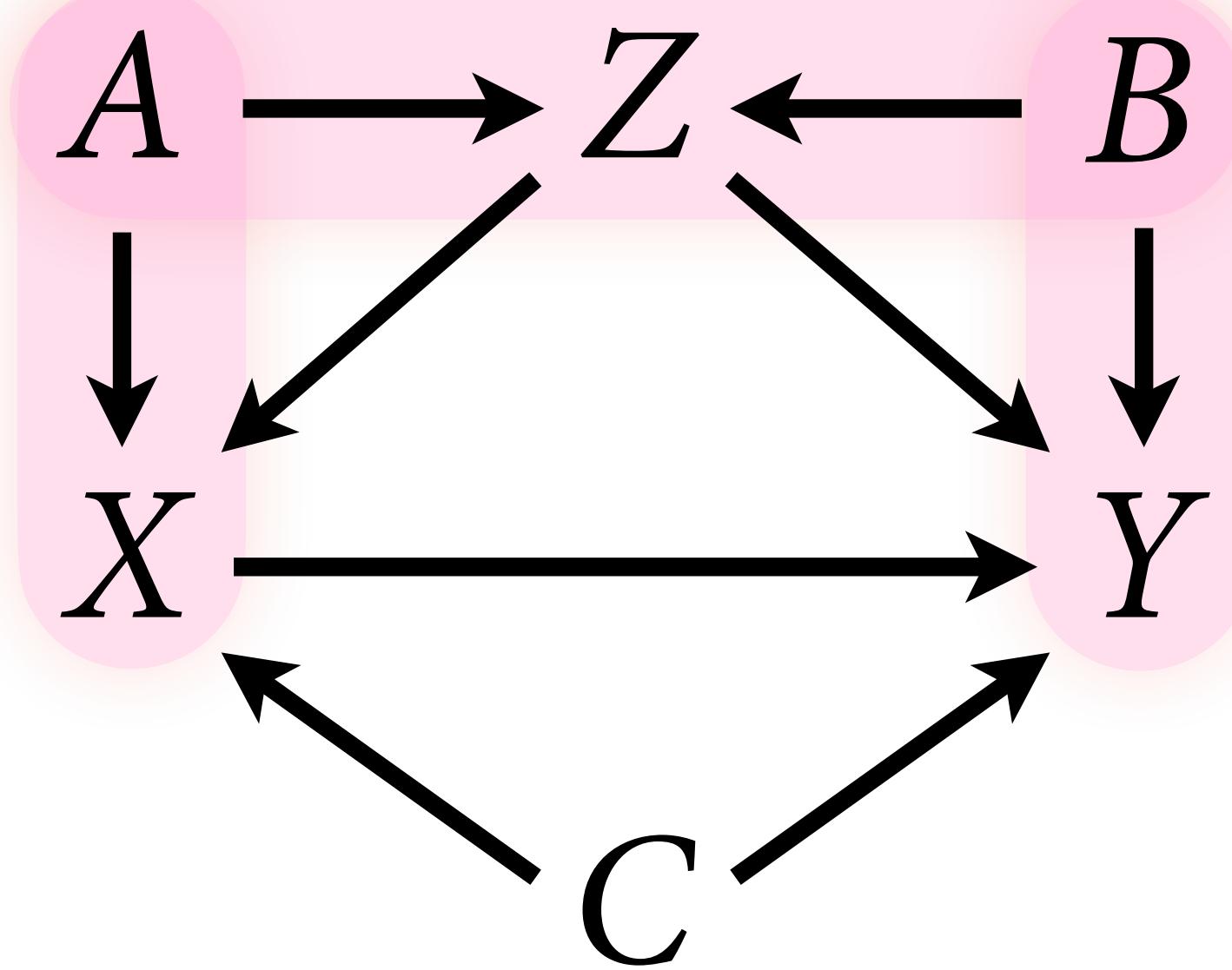
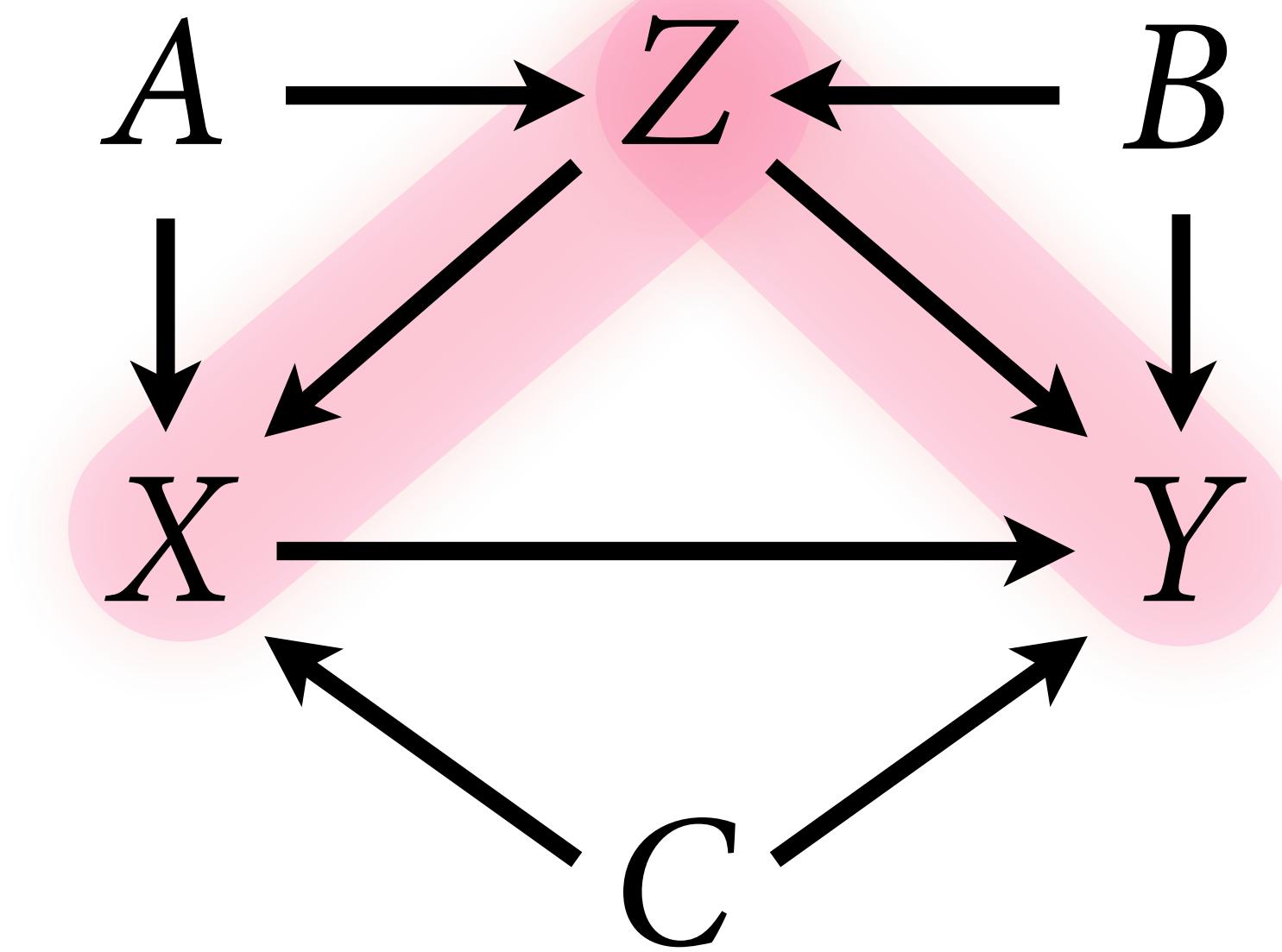
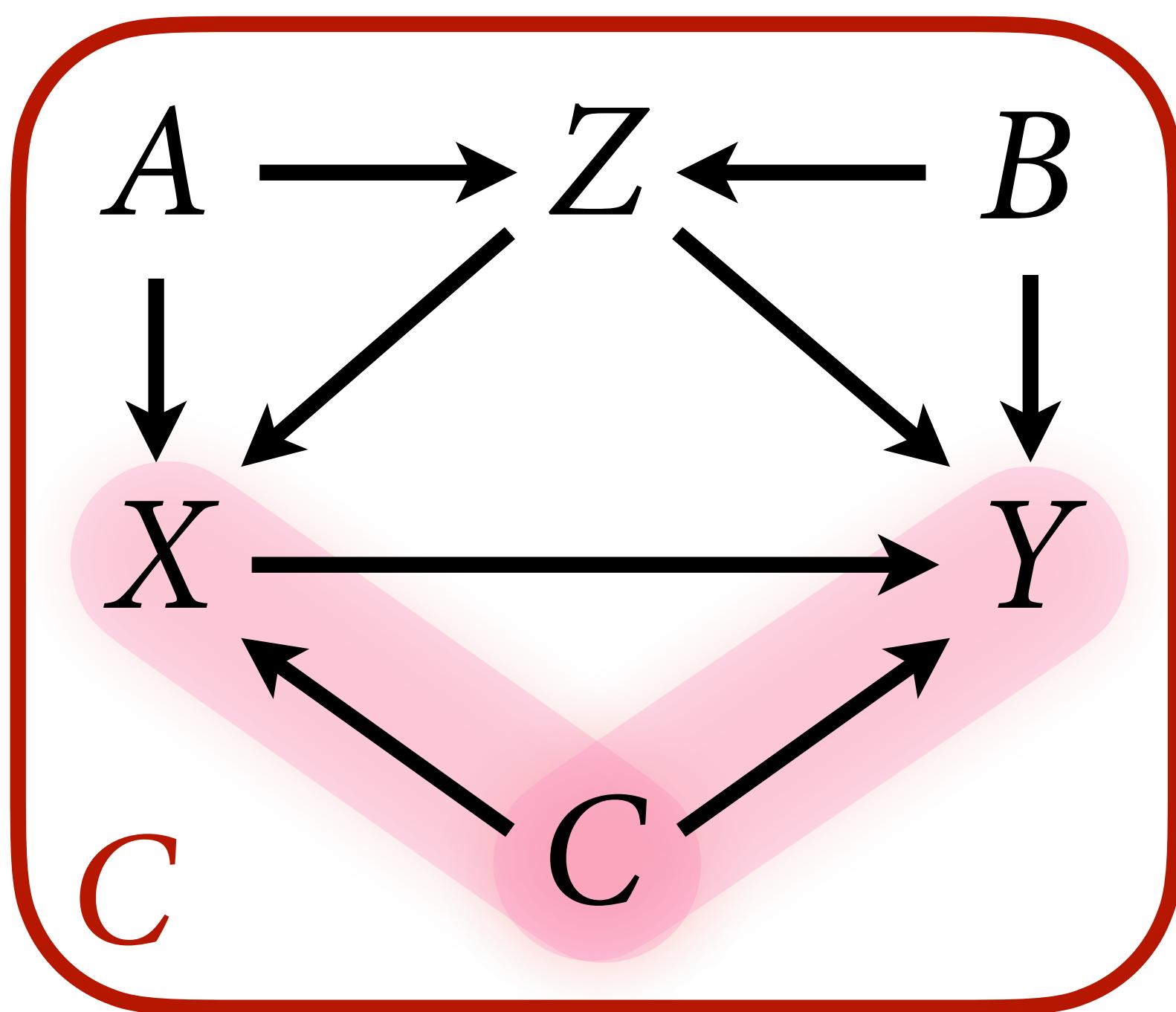
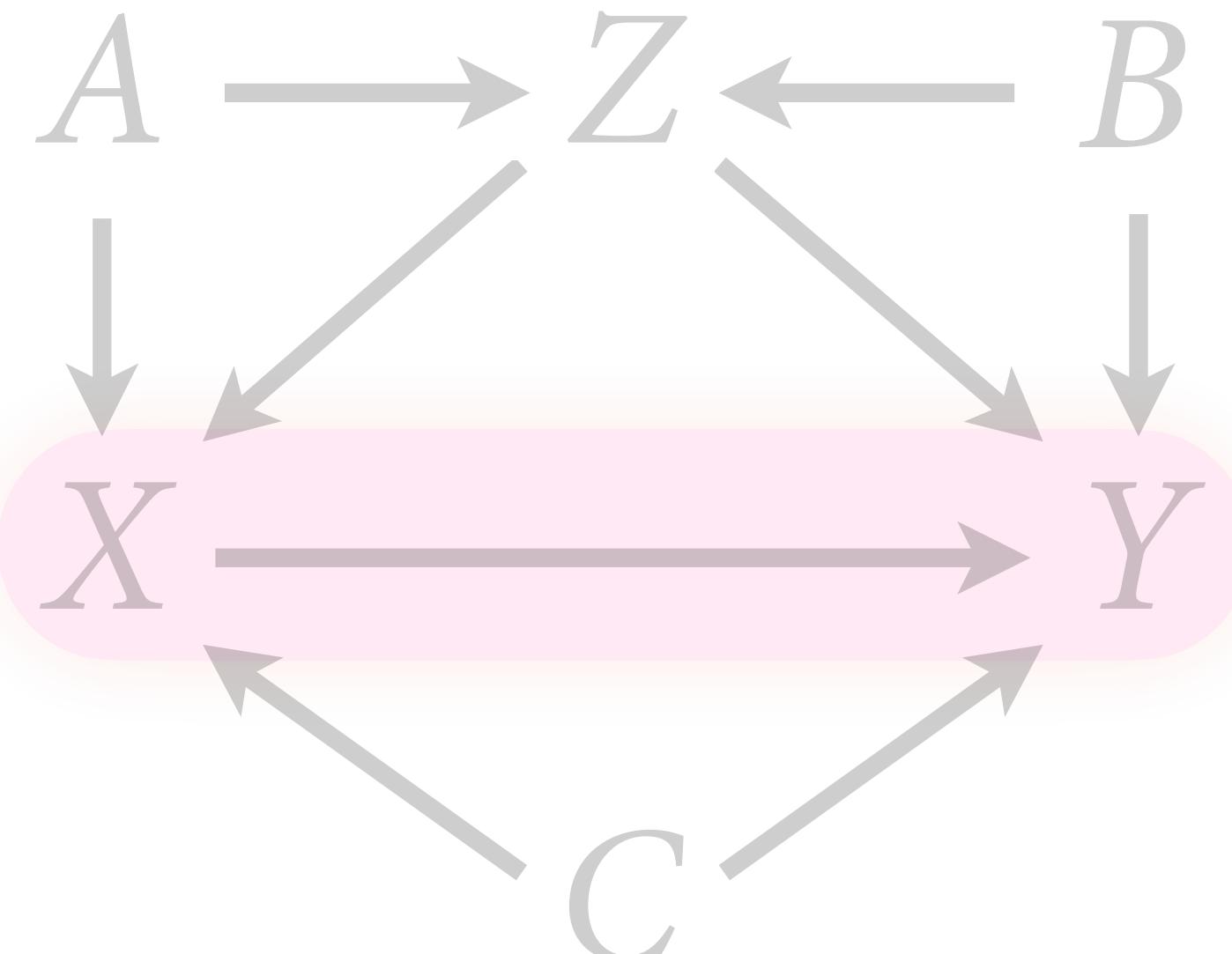


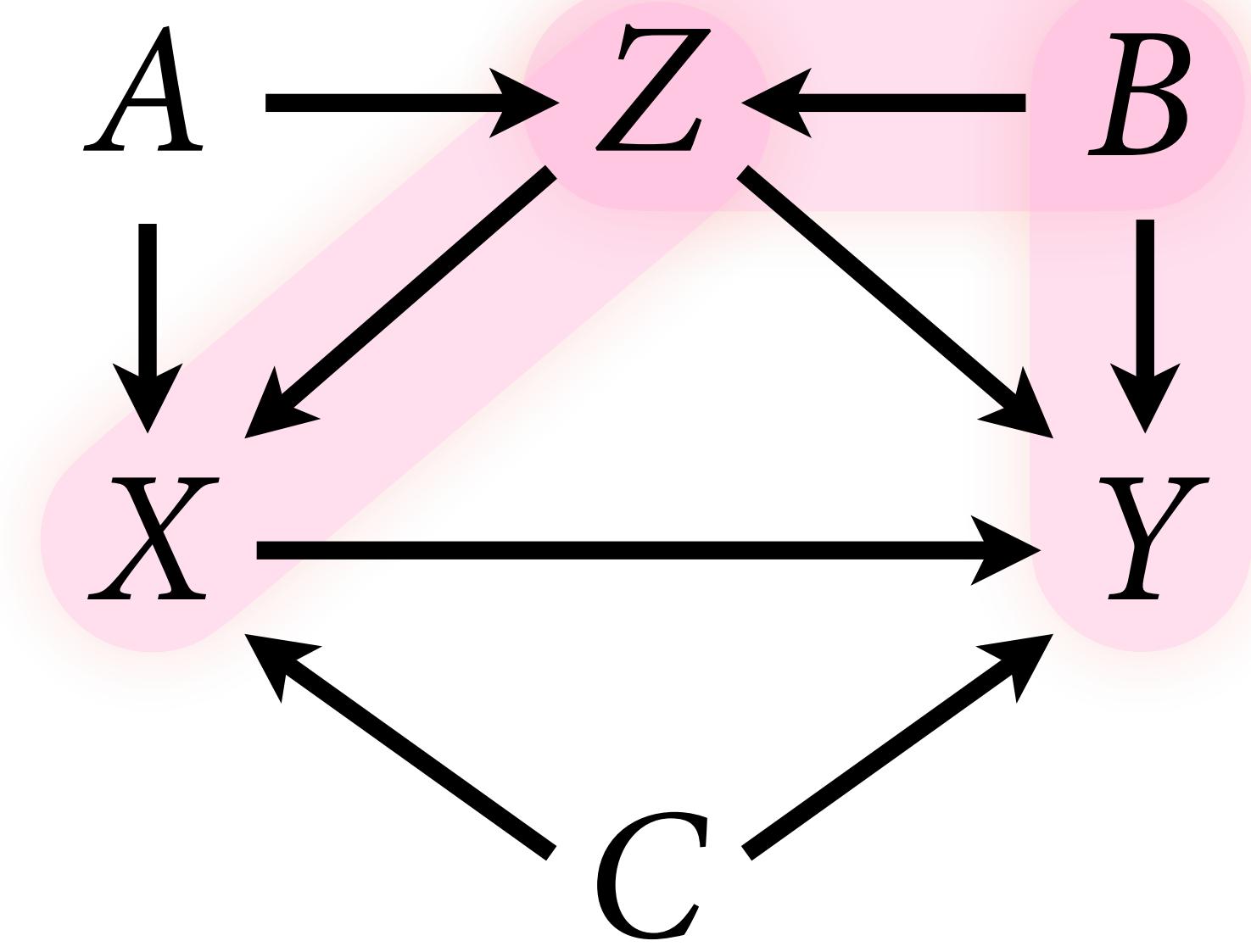
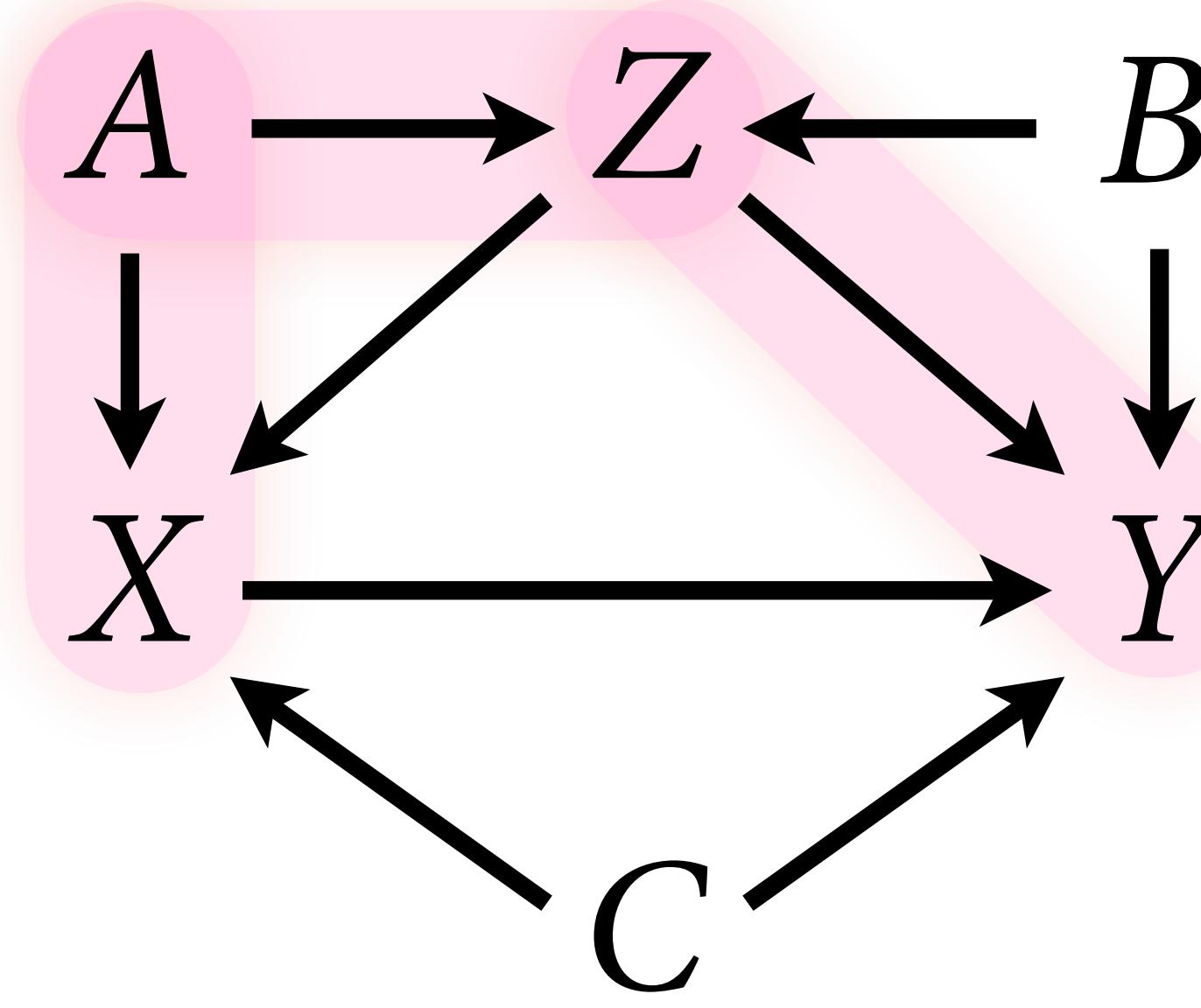
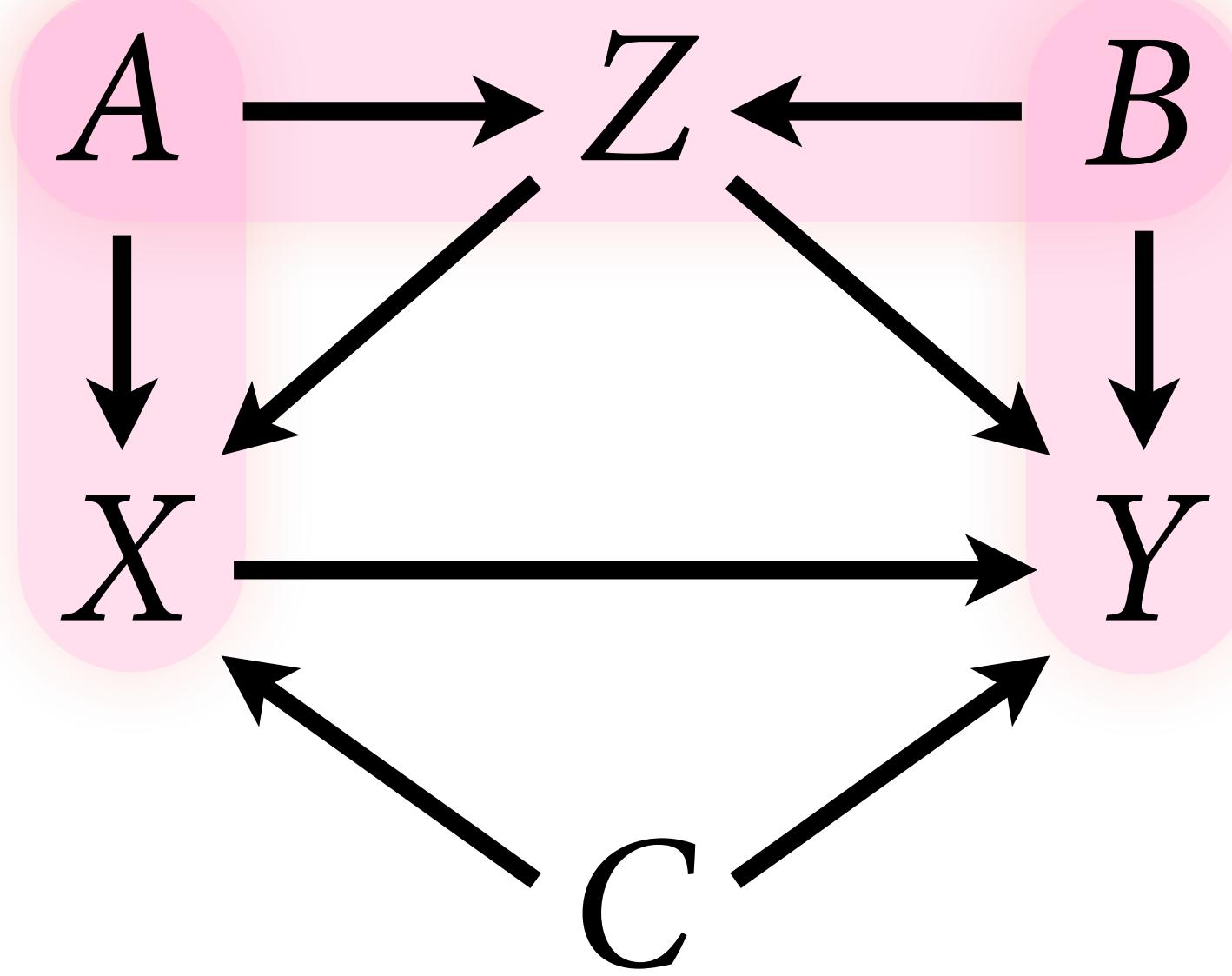
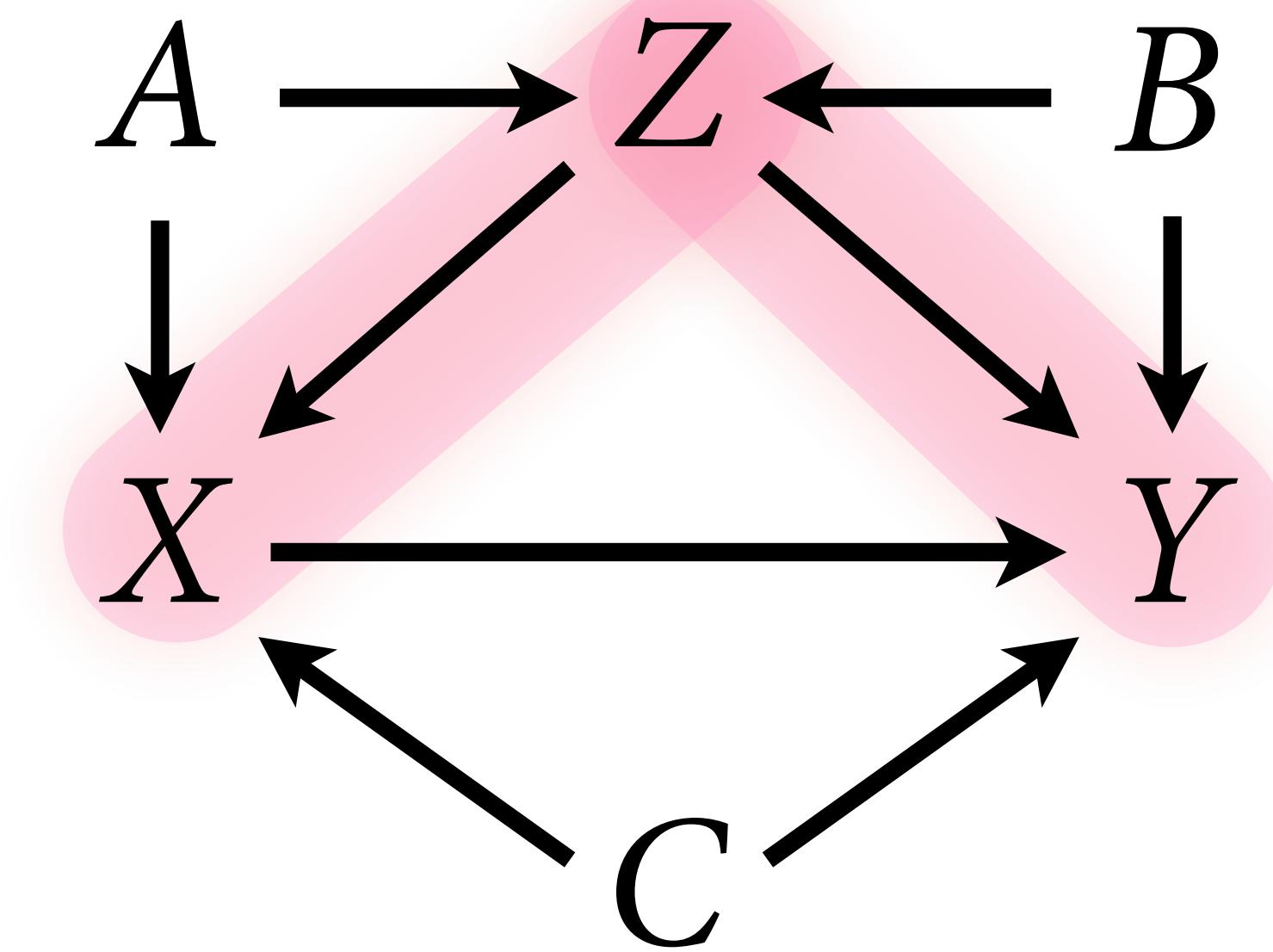
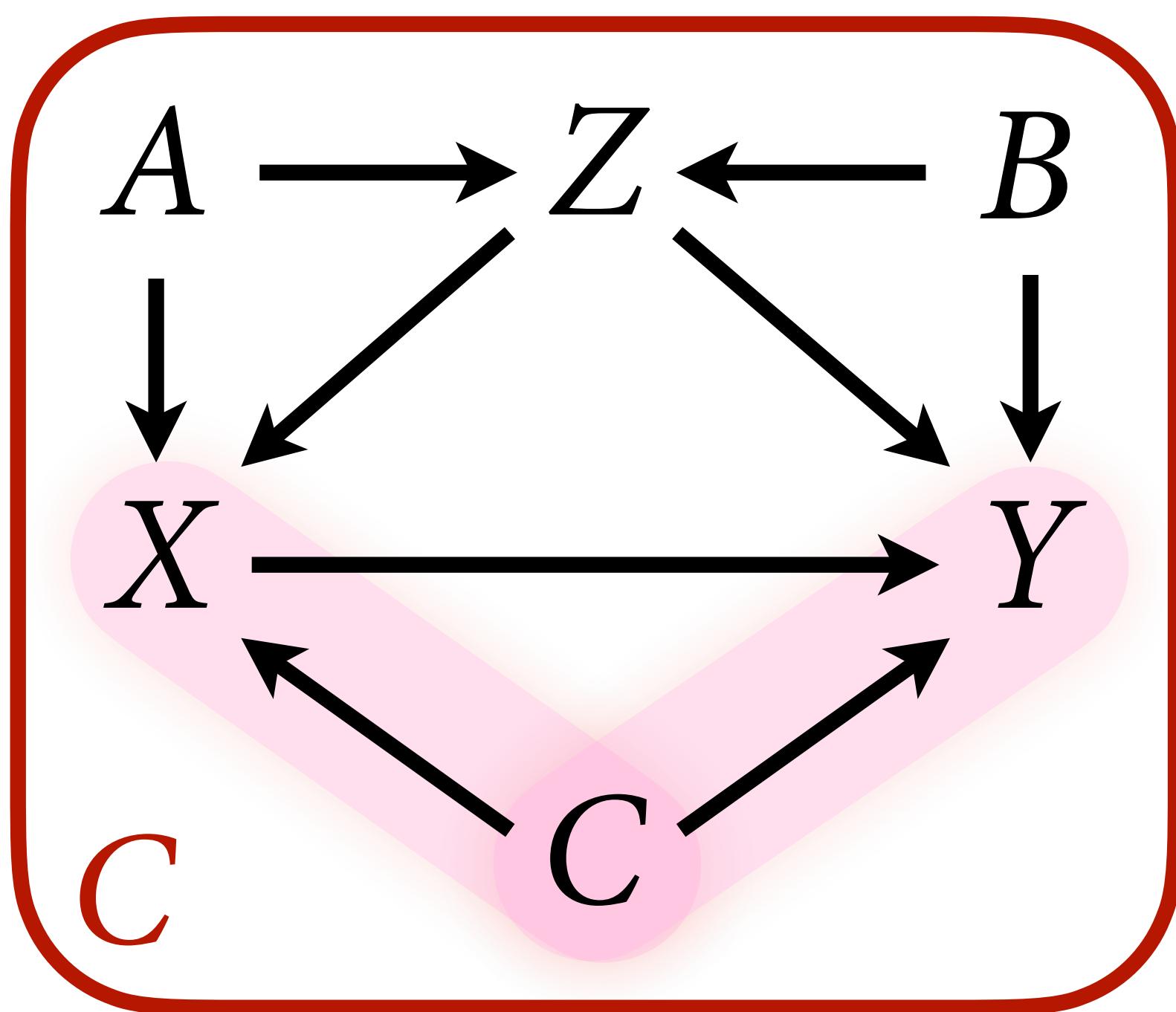
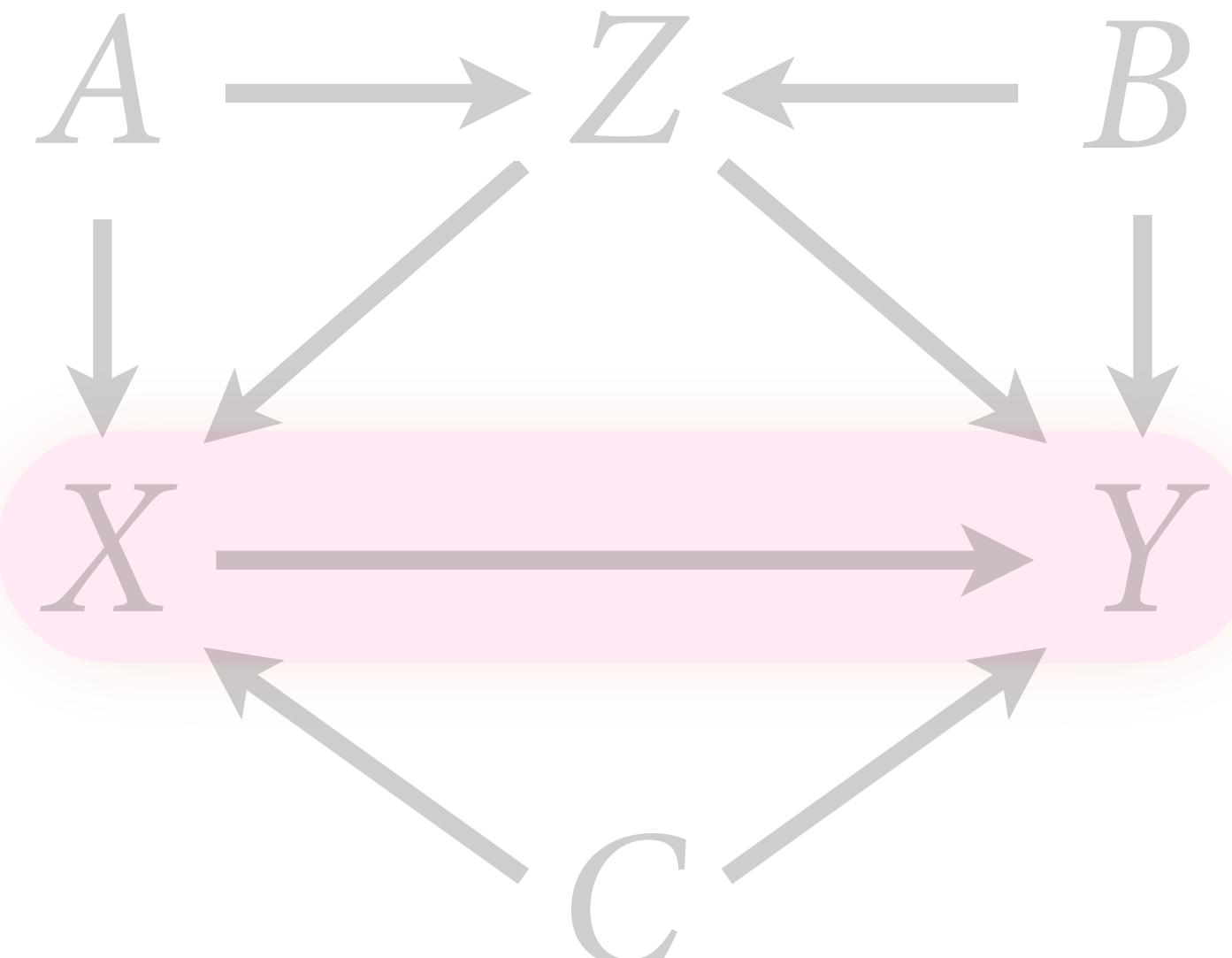
Causal path, open

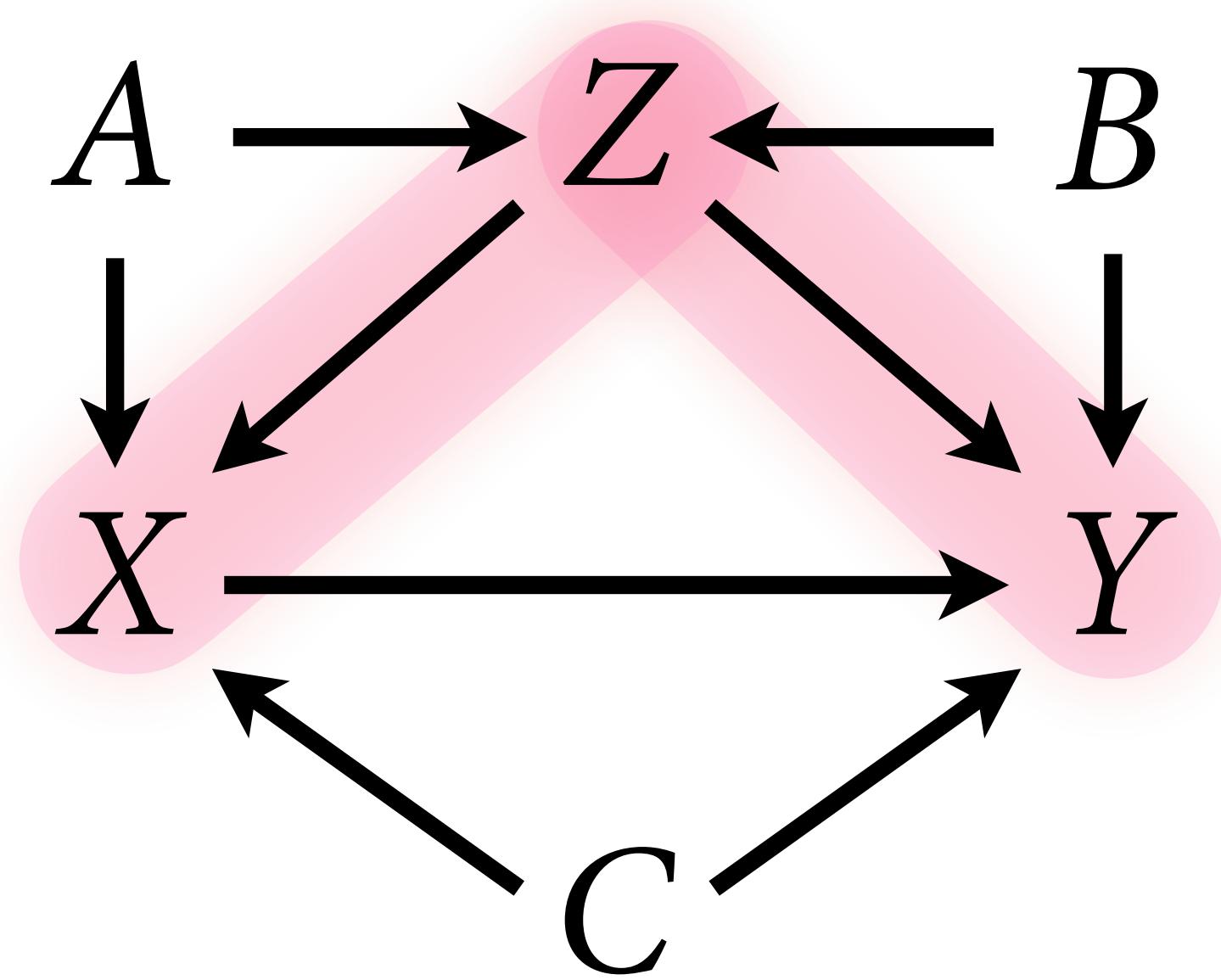




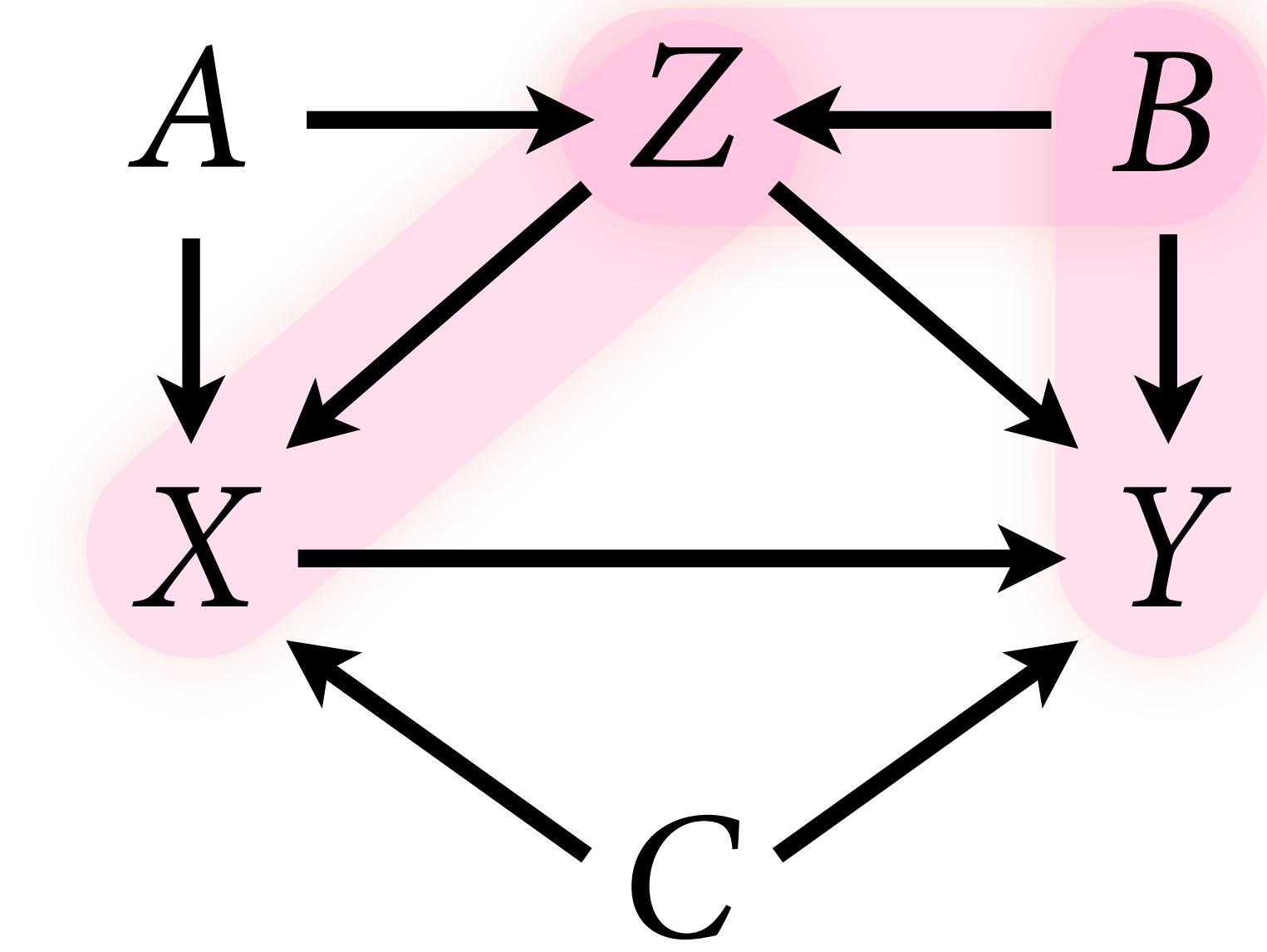
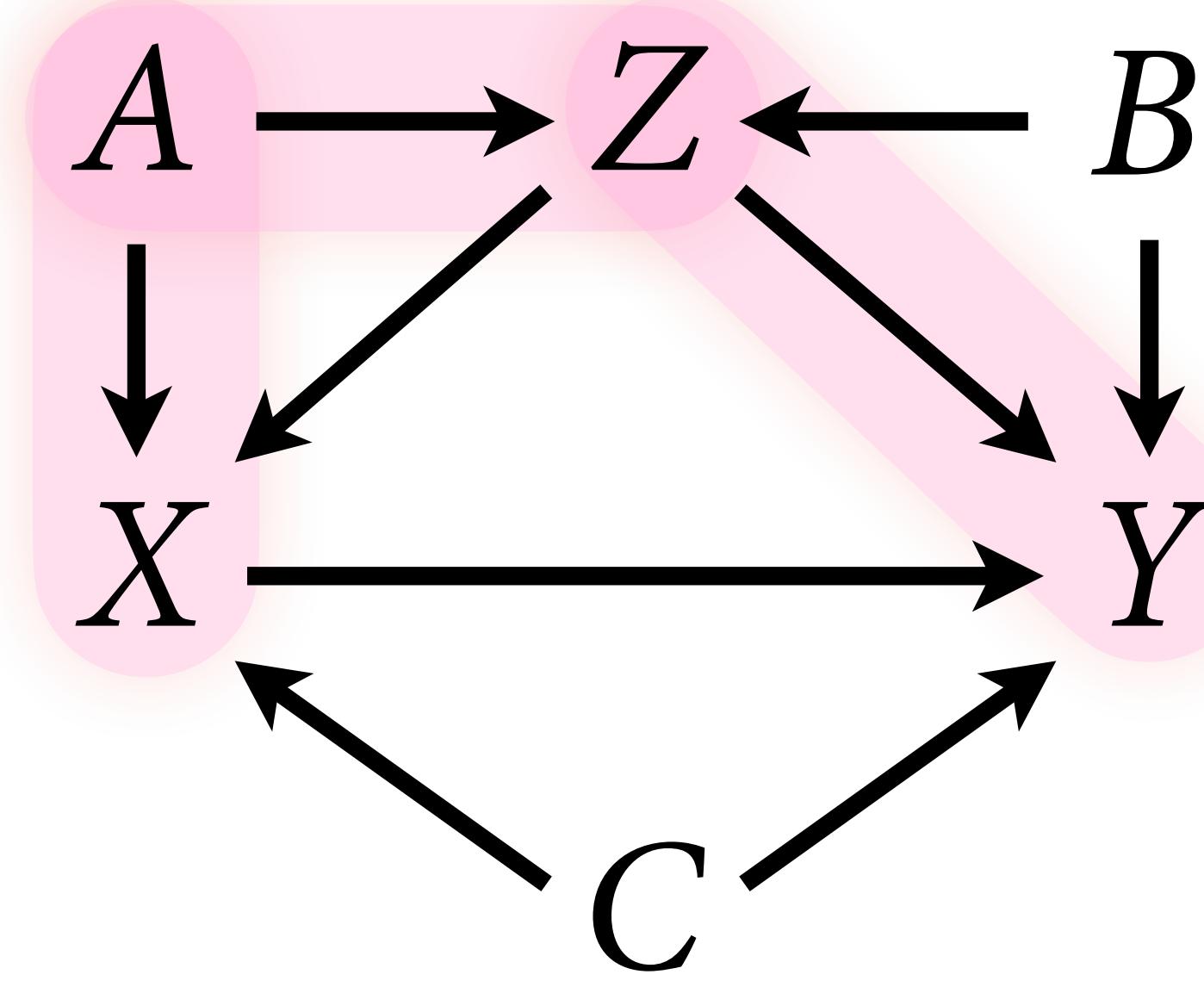
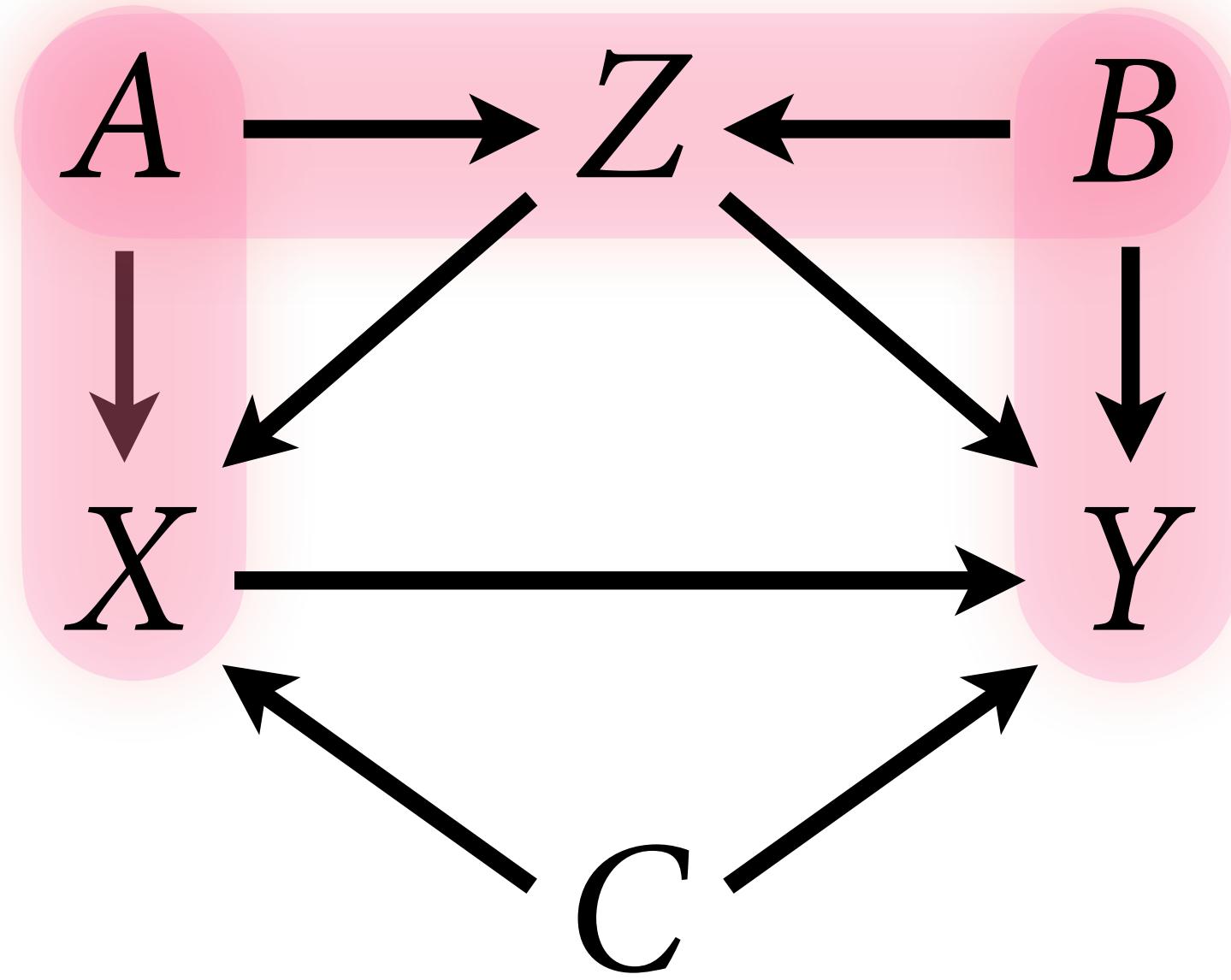
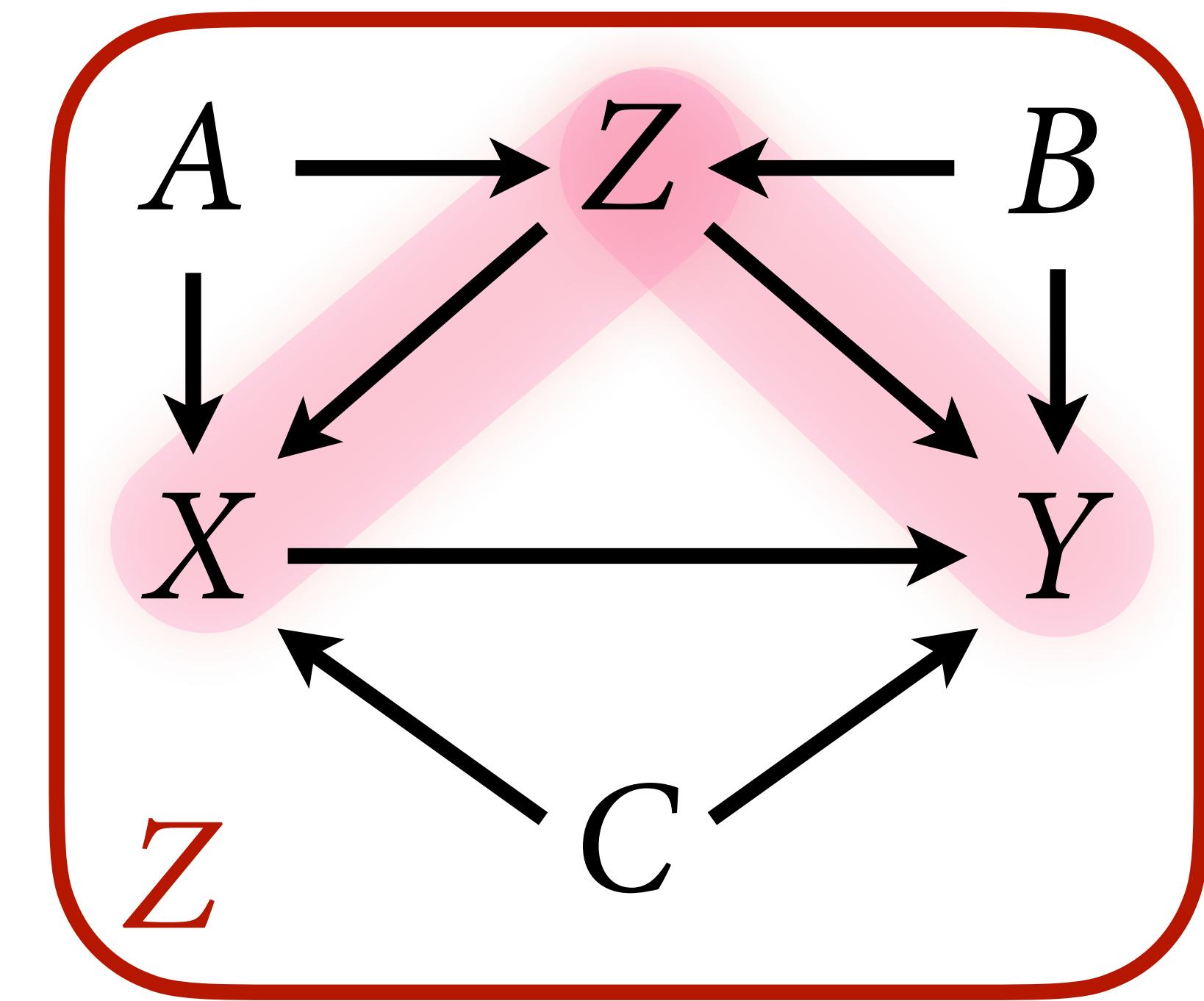
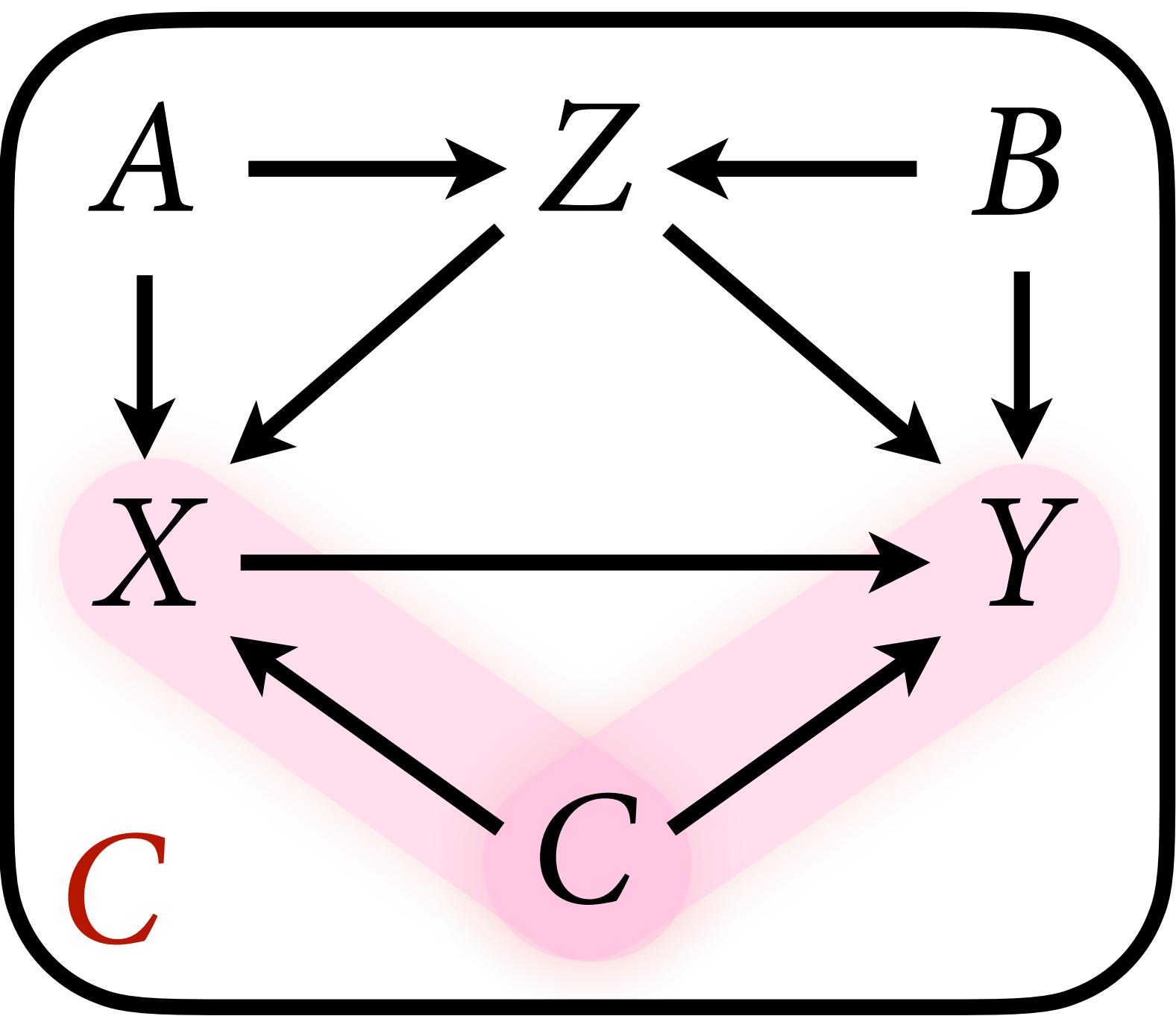
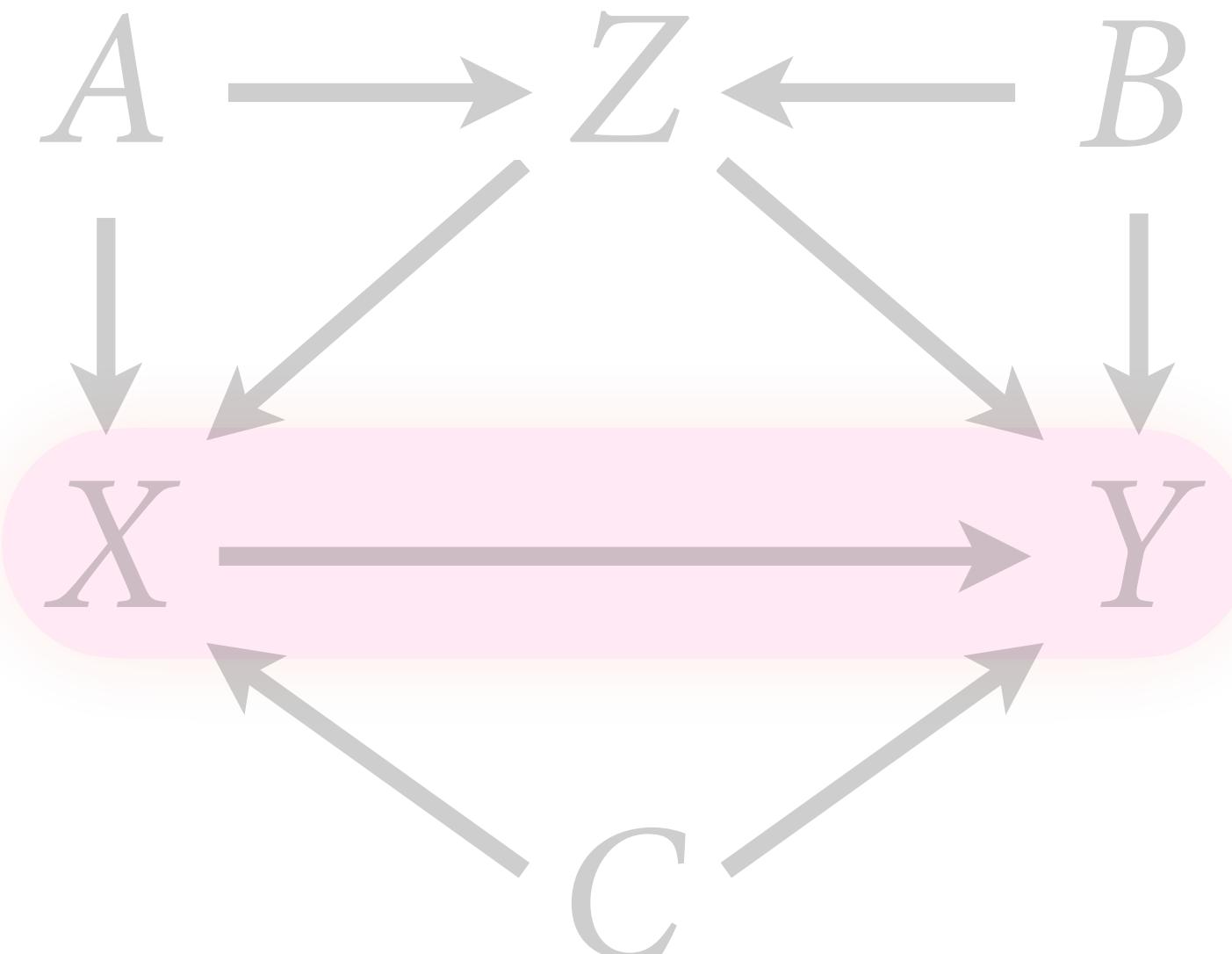
Backdoor path, open  
Close with C

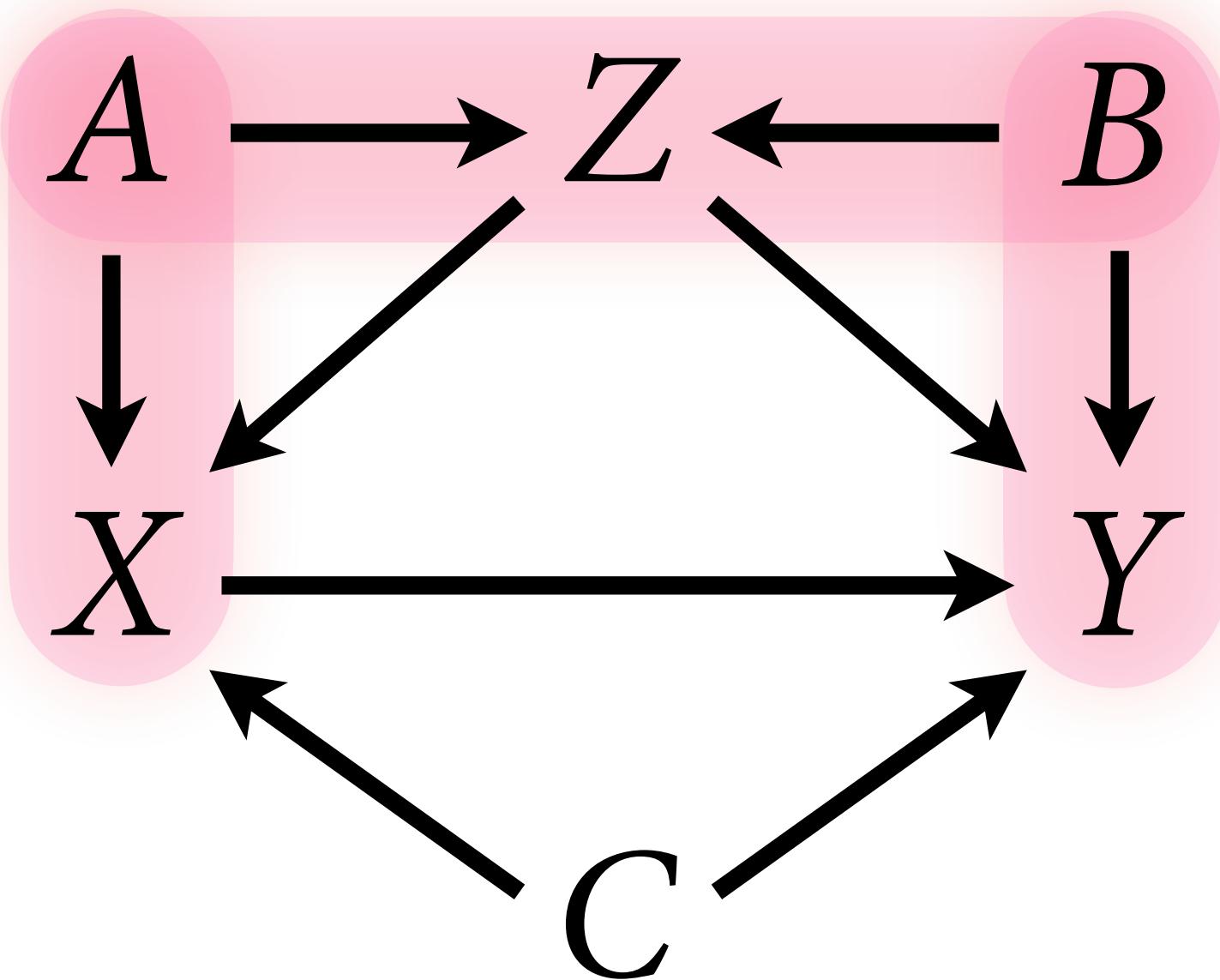




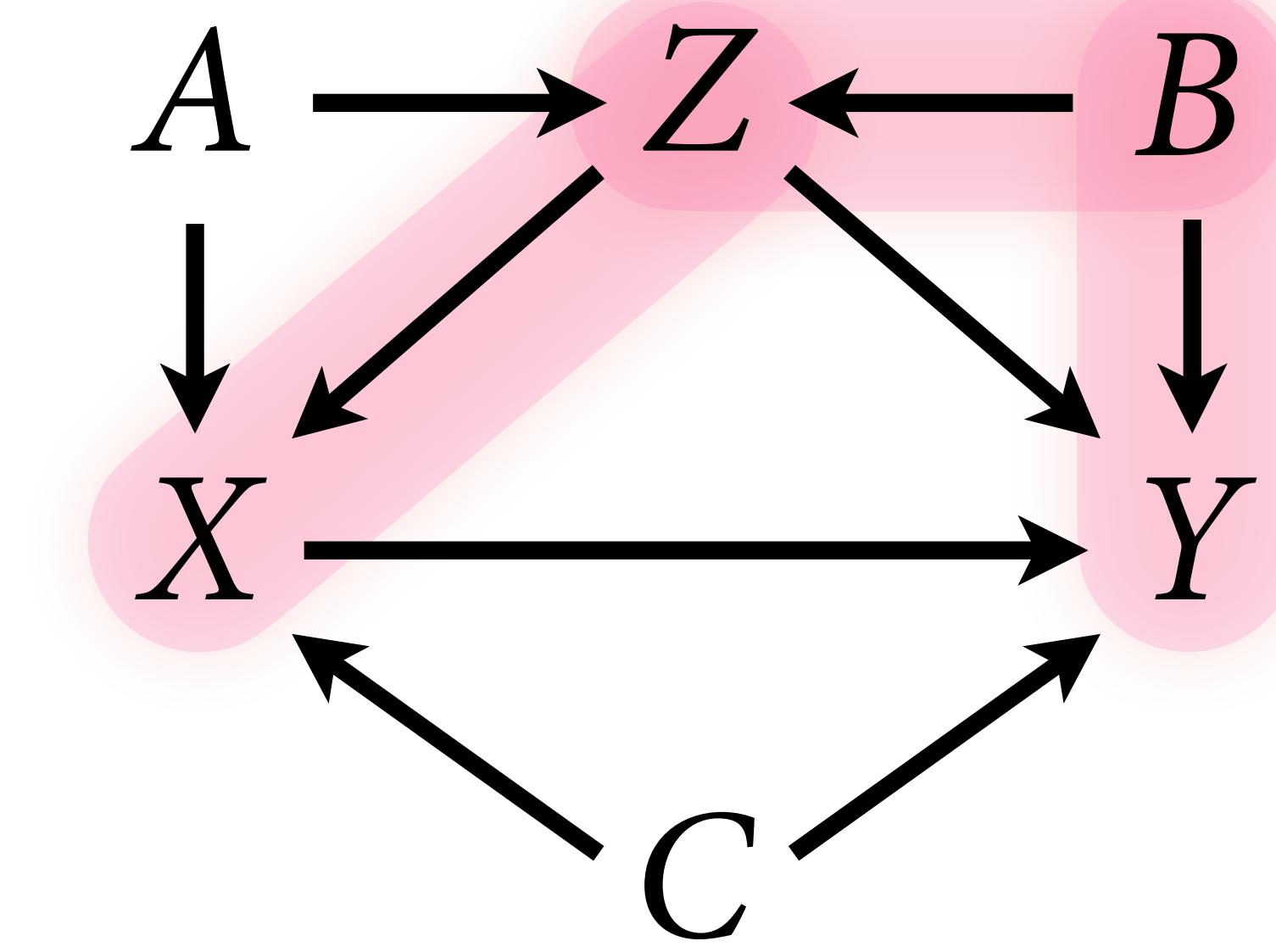
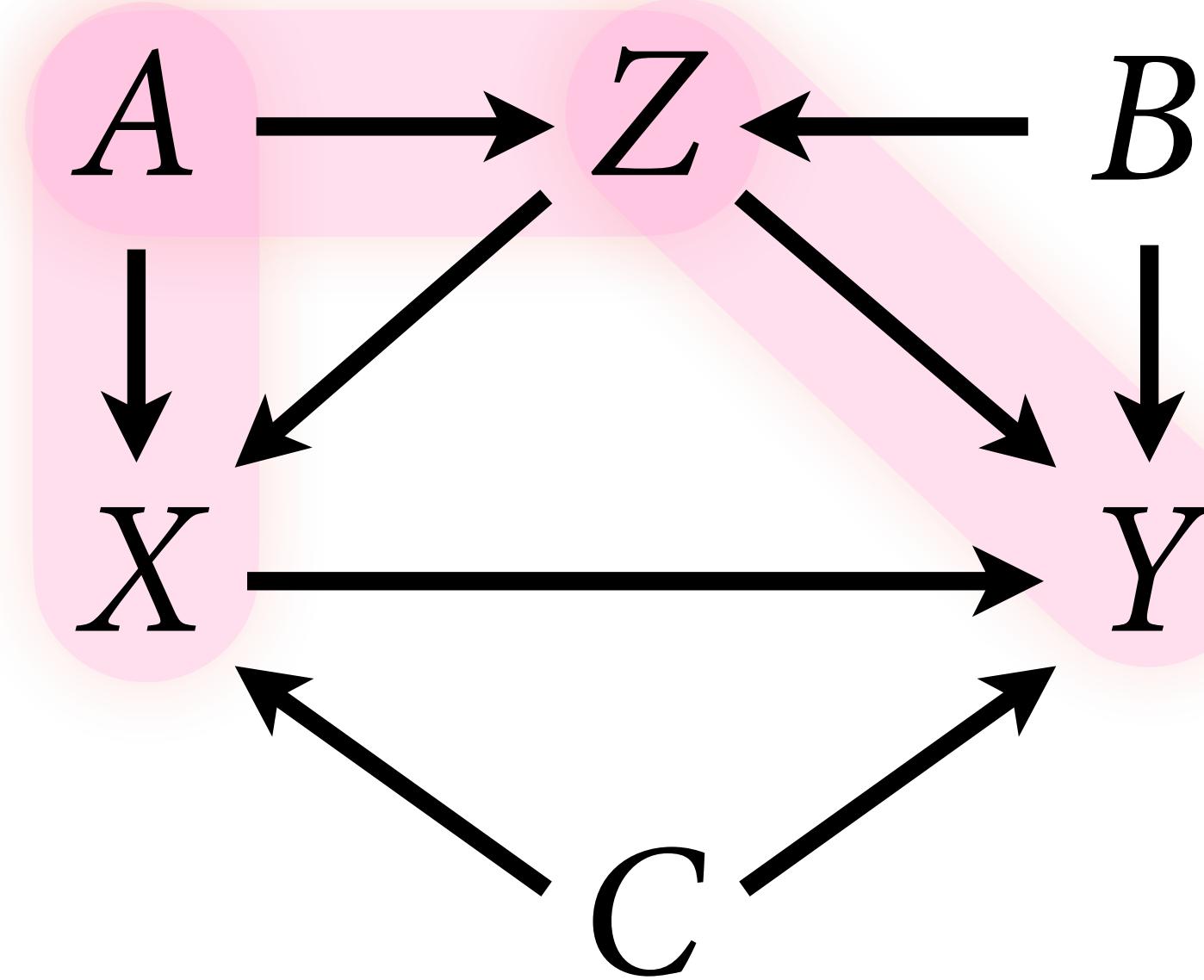
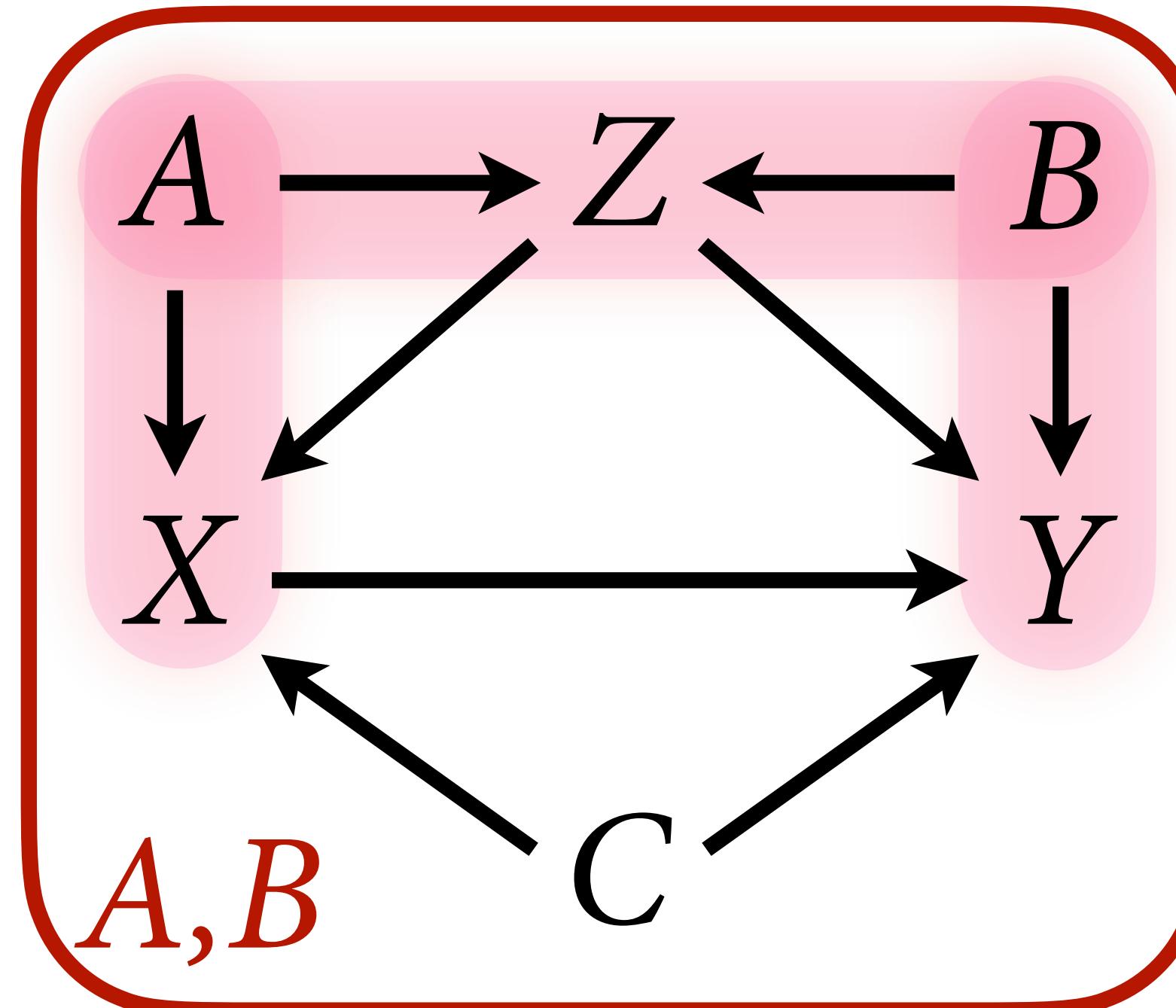
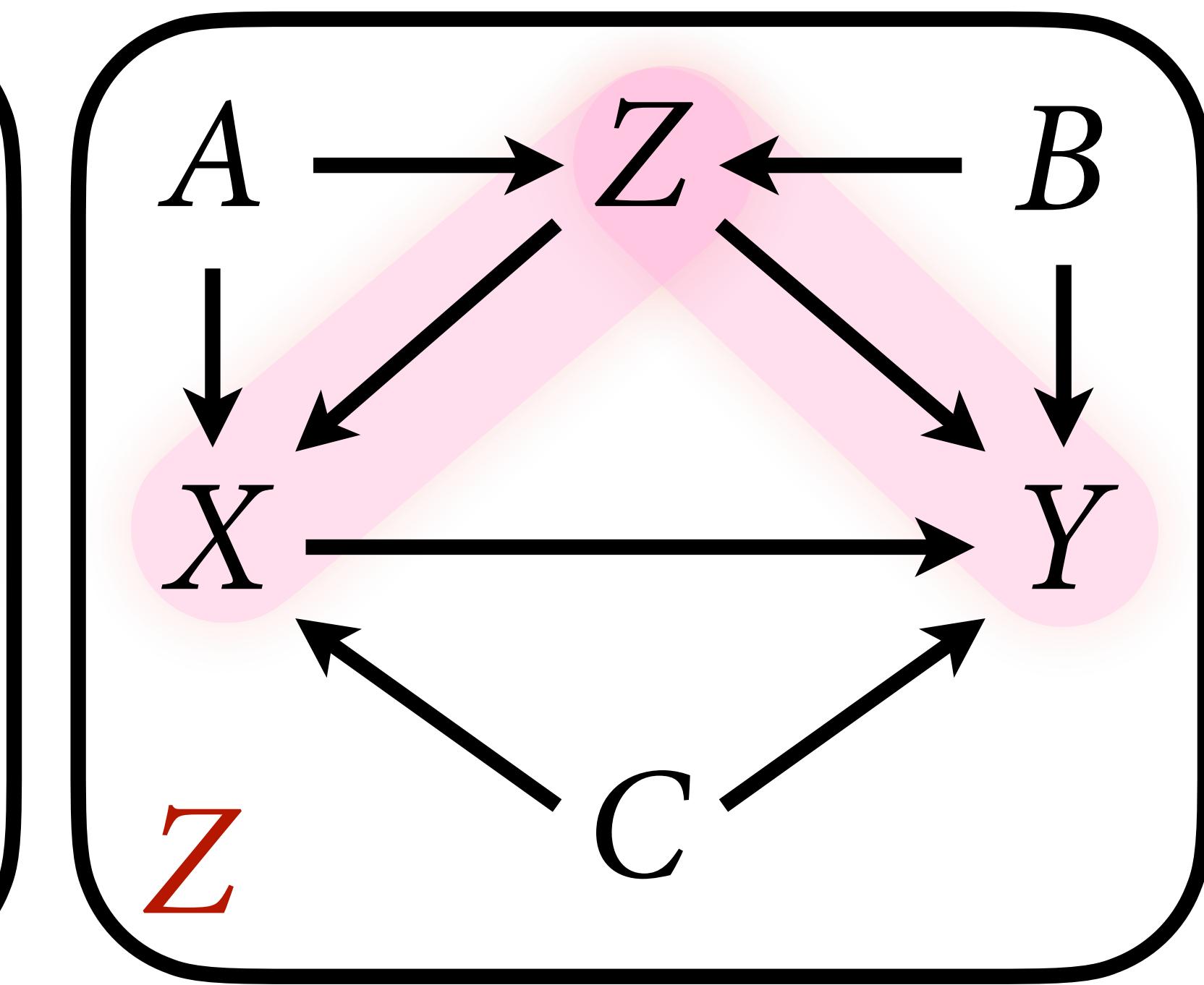
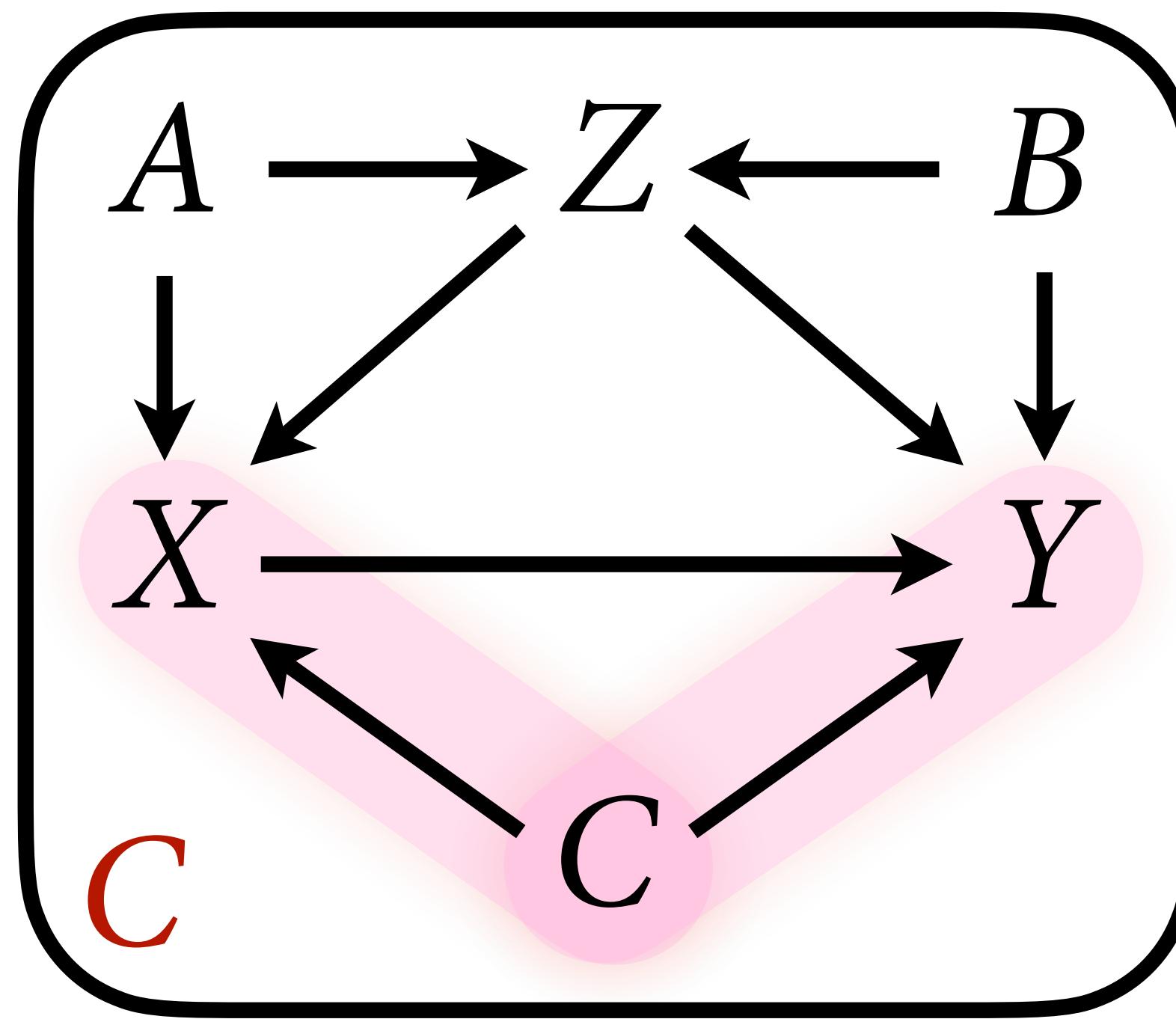
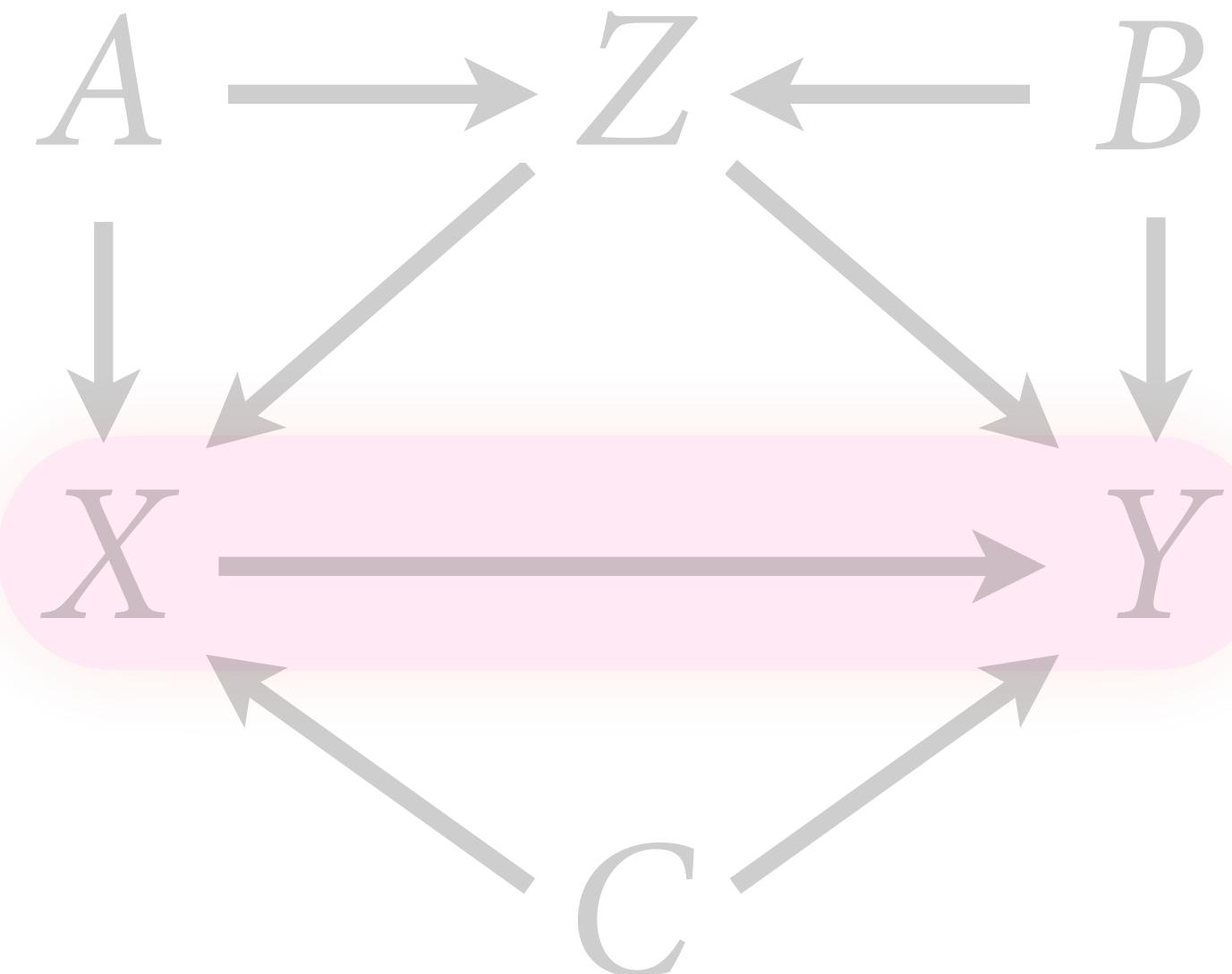


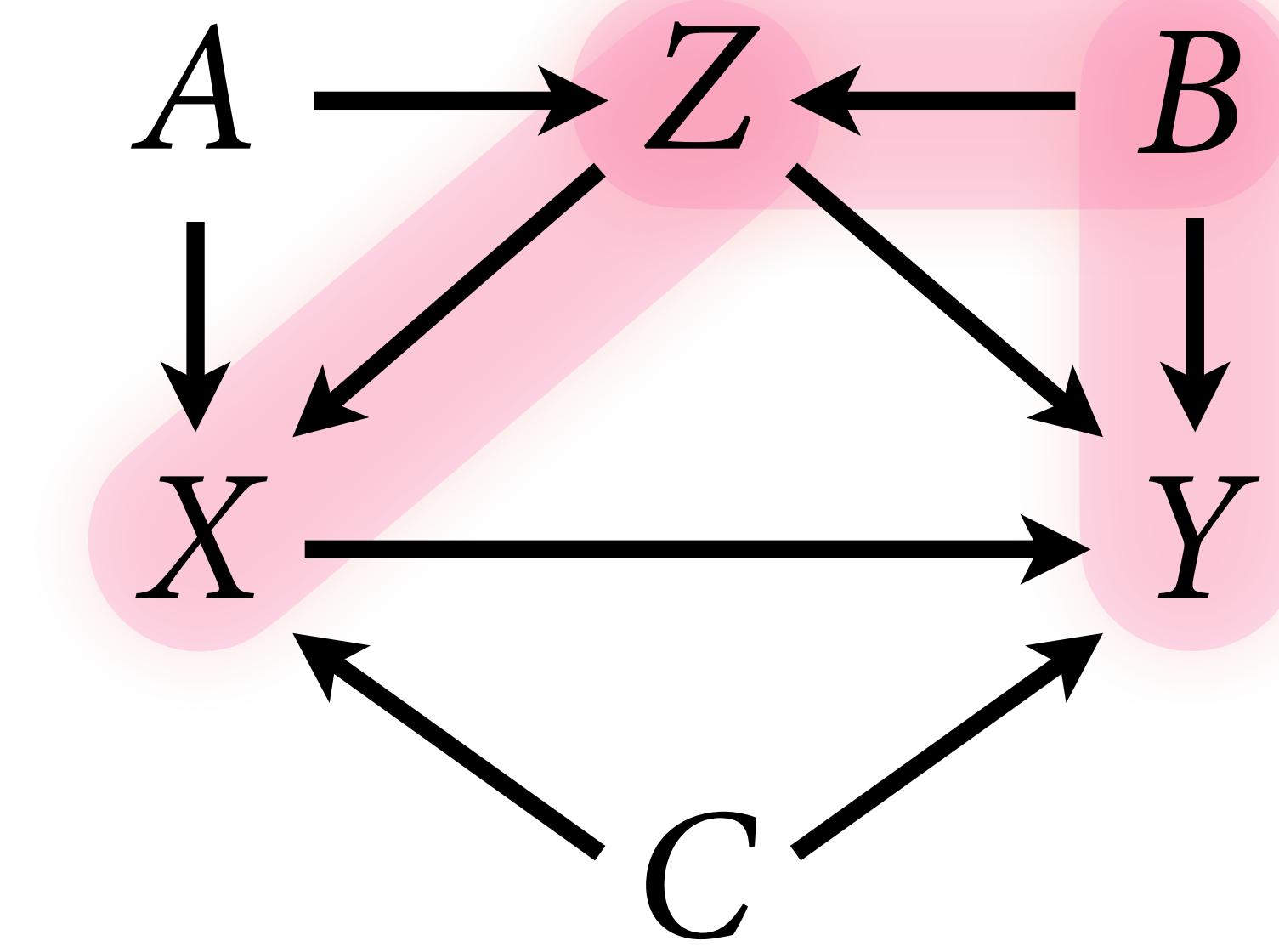
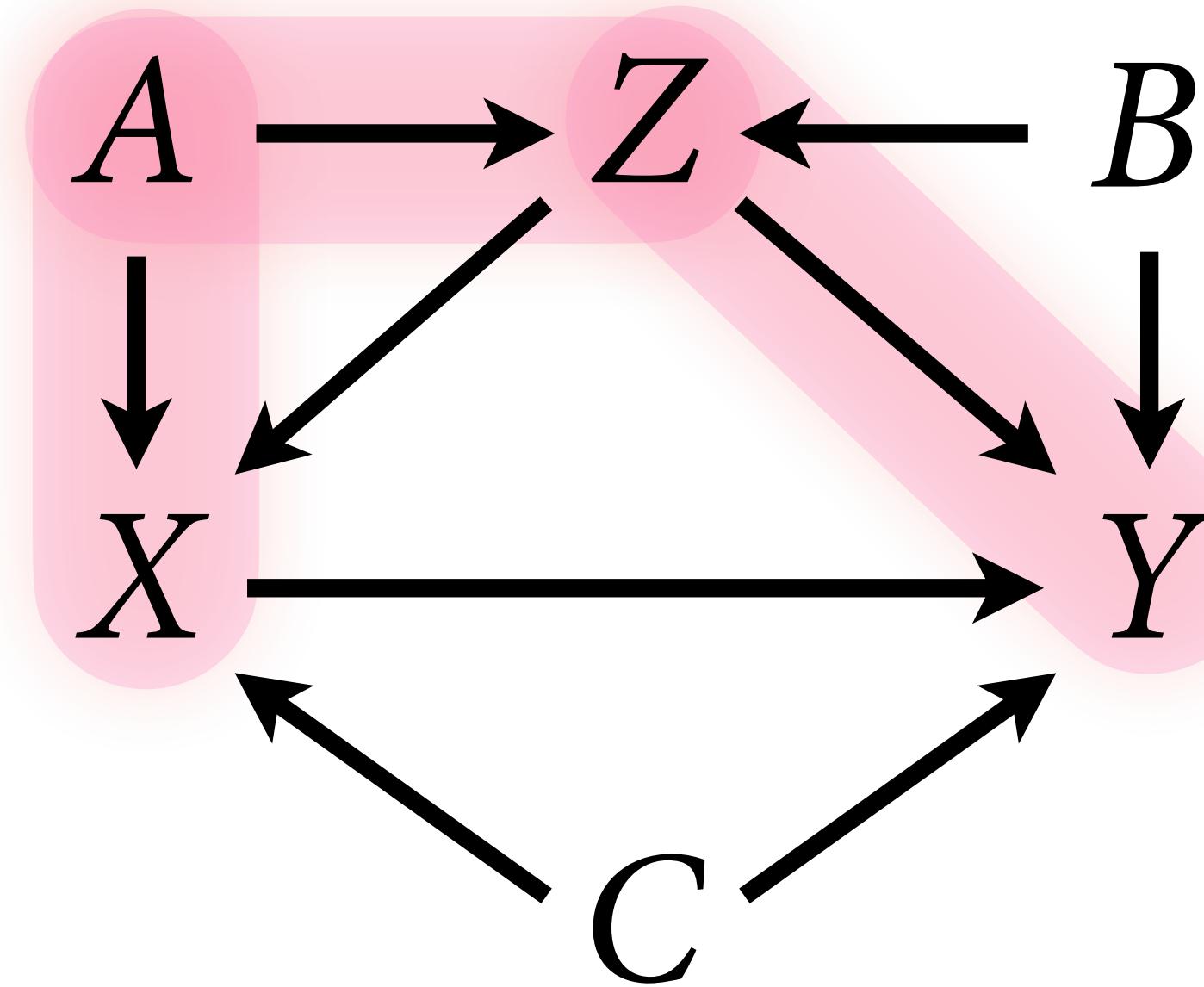
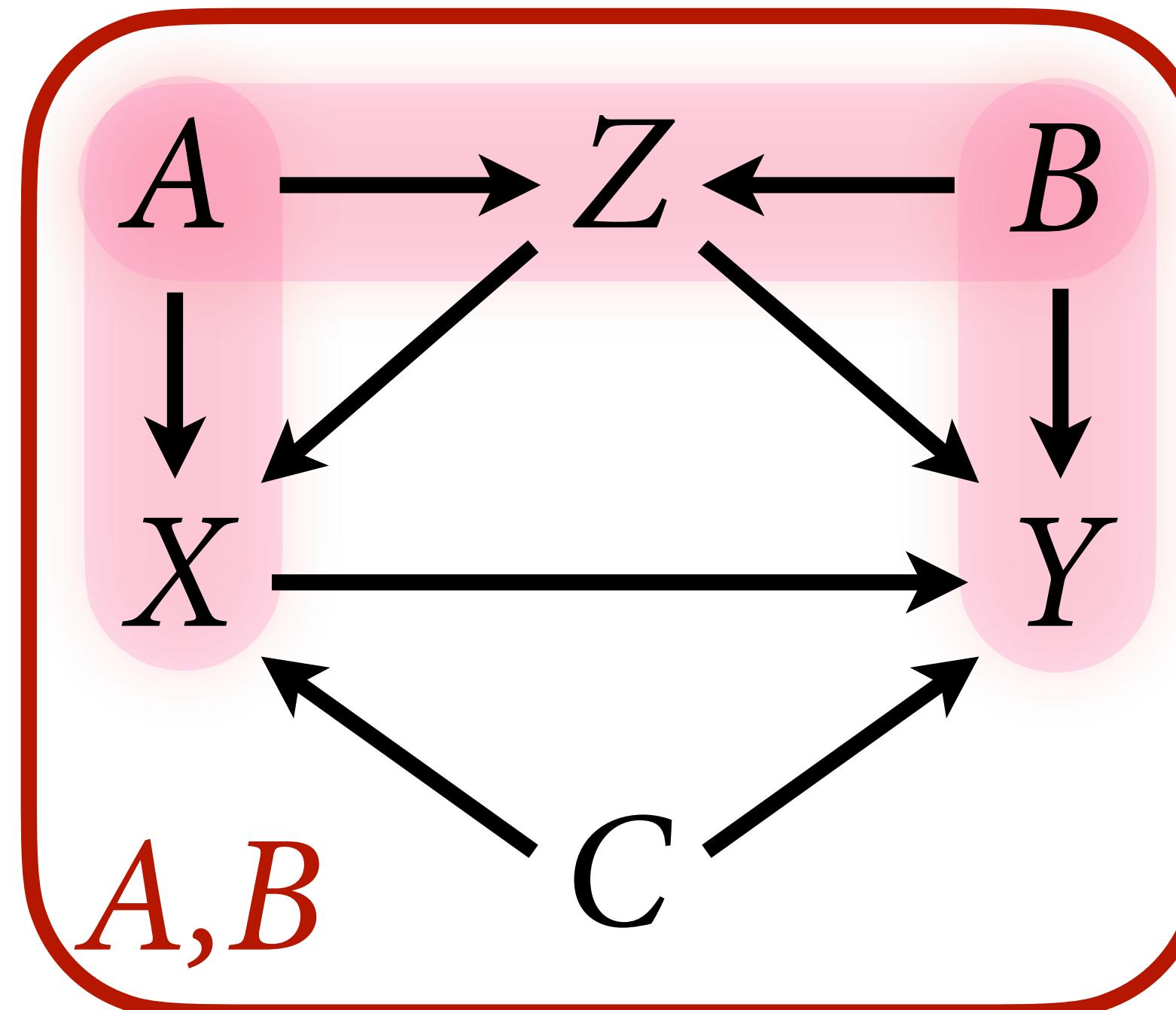
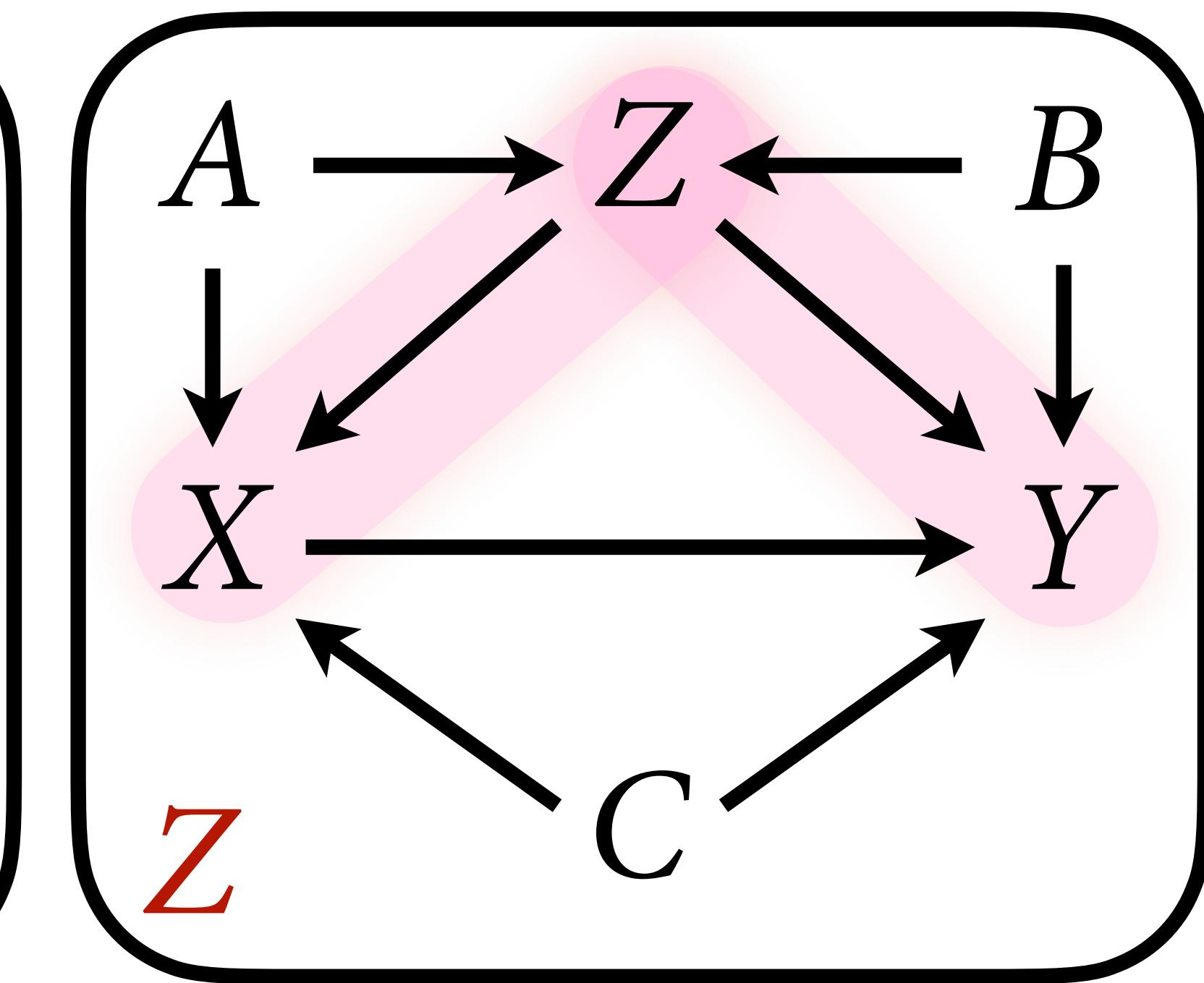
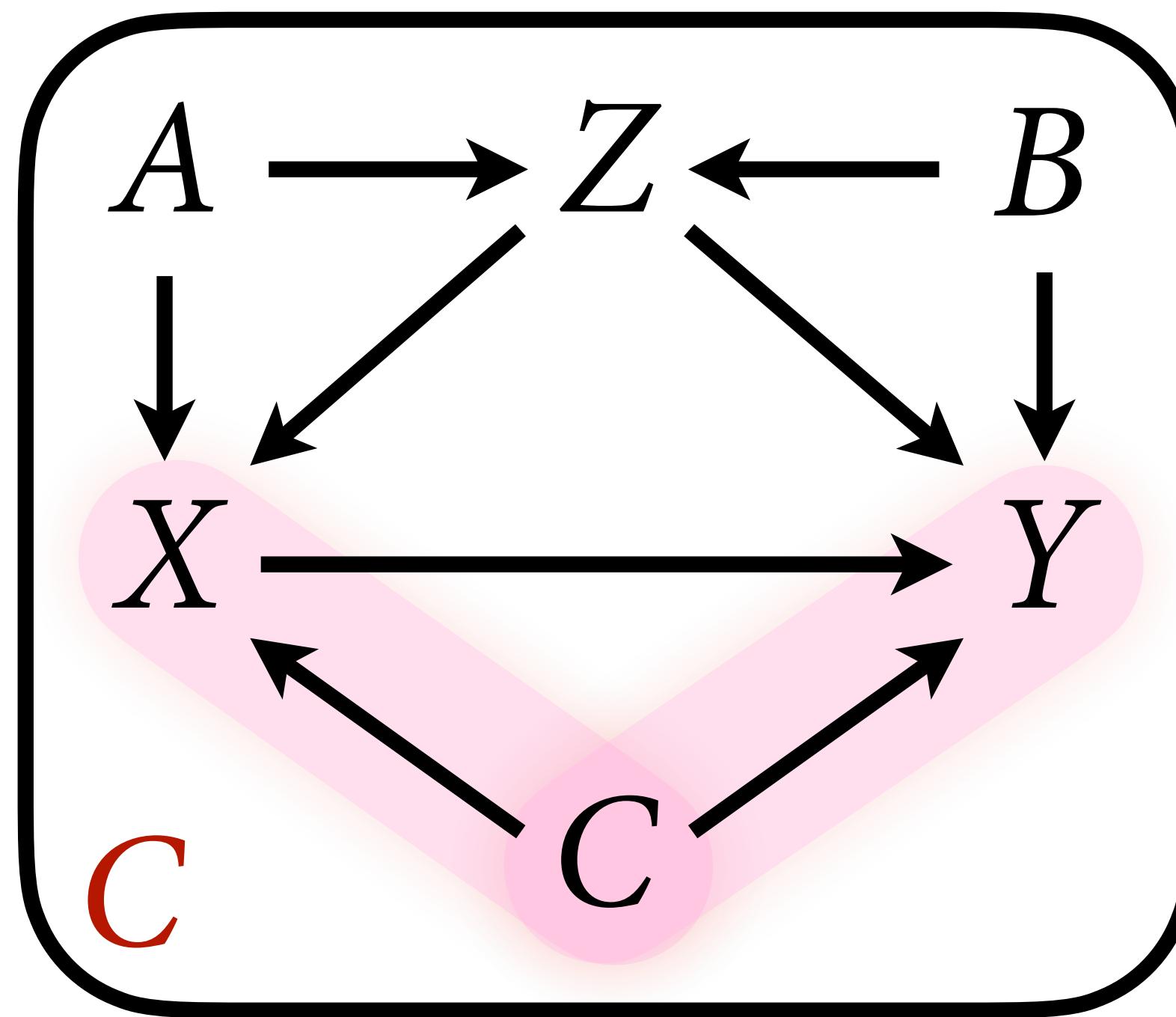
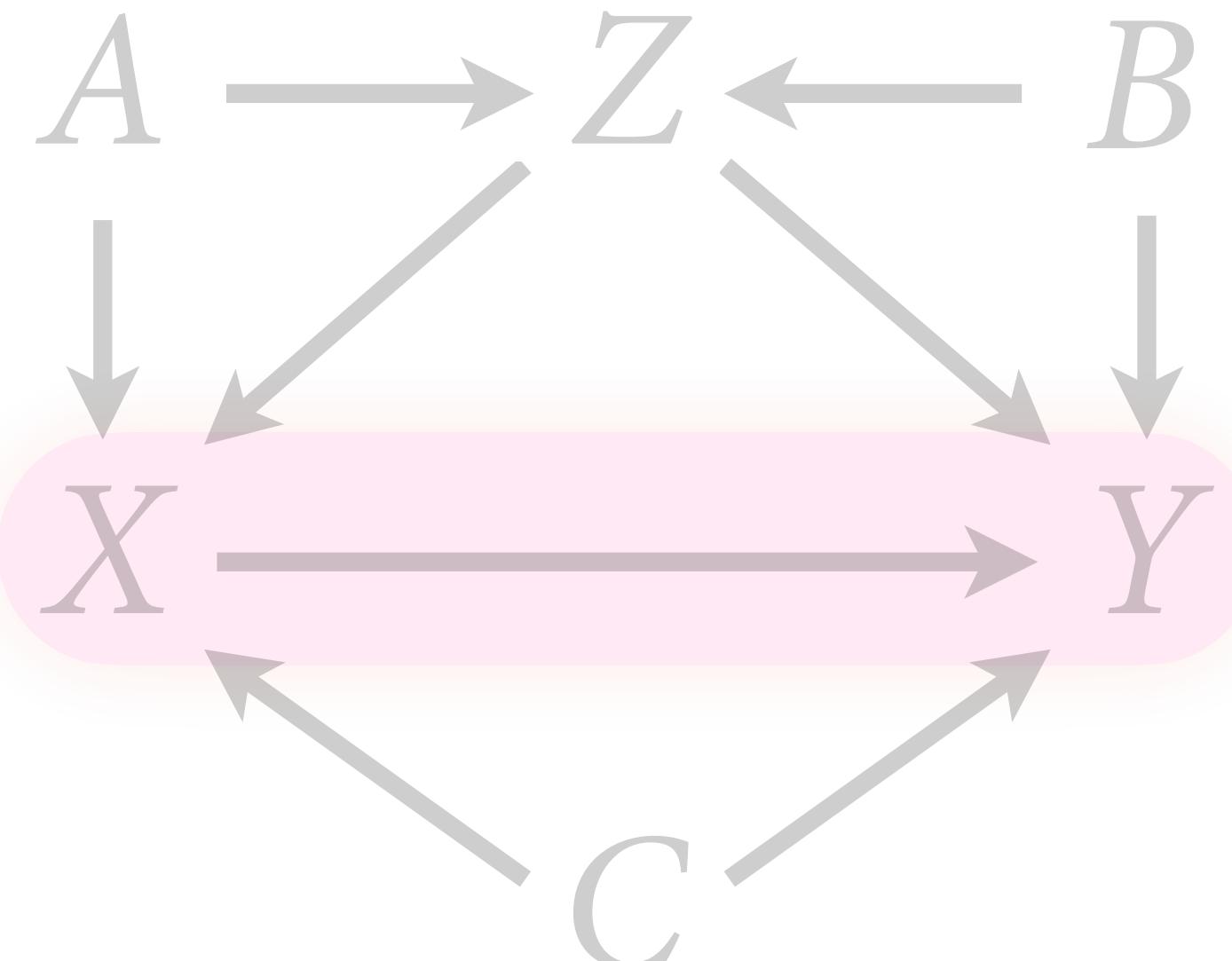
Backdoor path, open  
Close with  $Z$

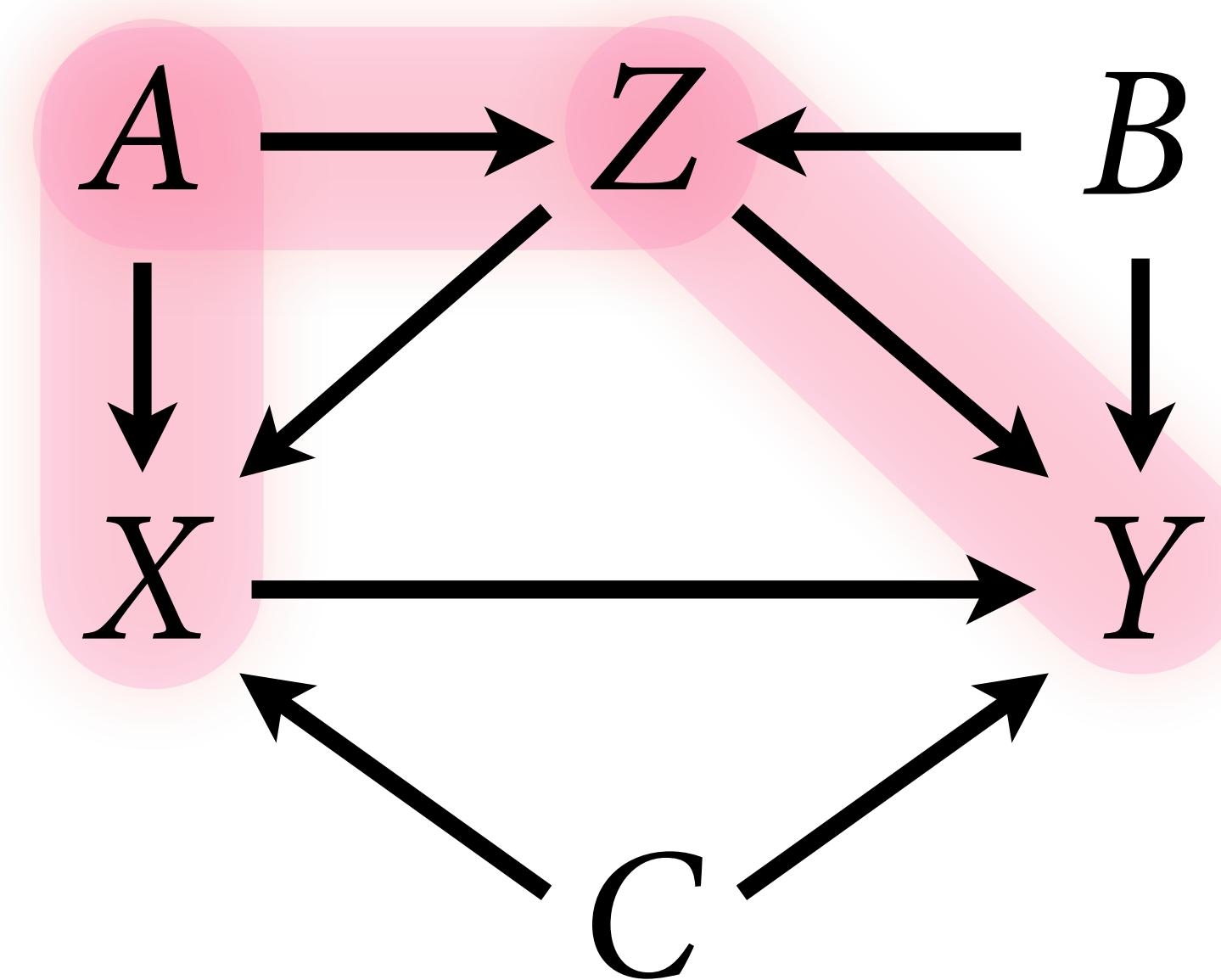




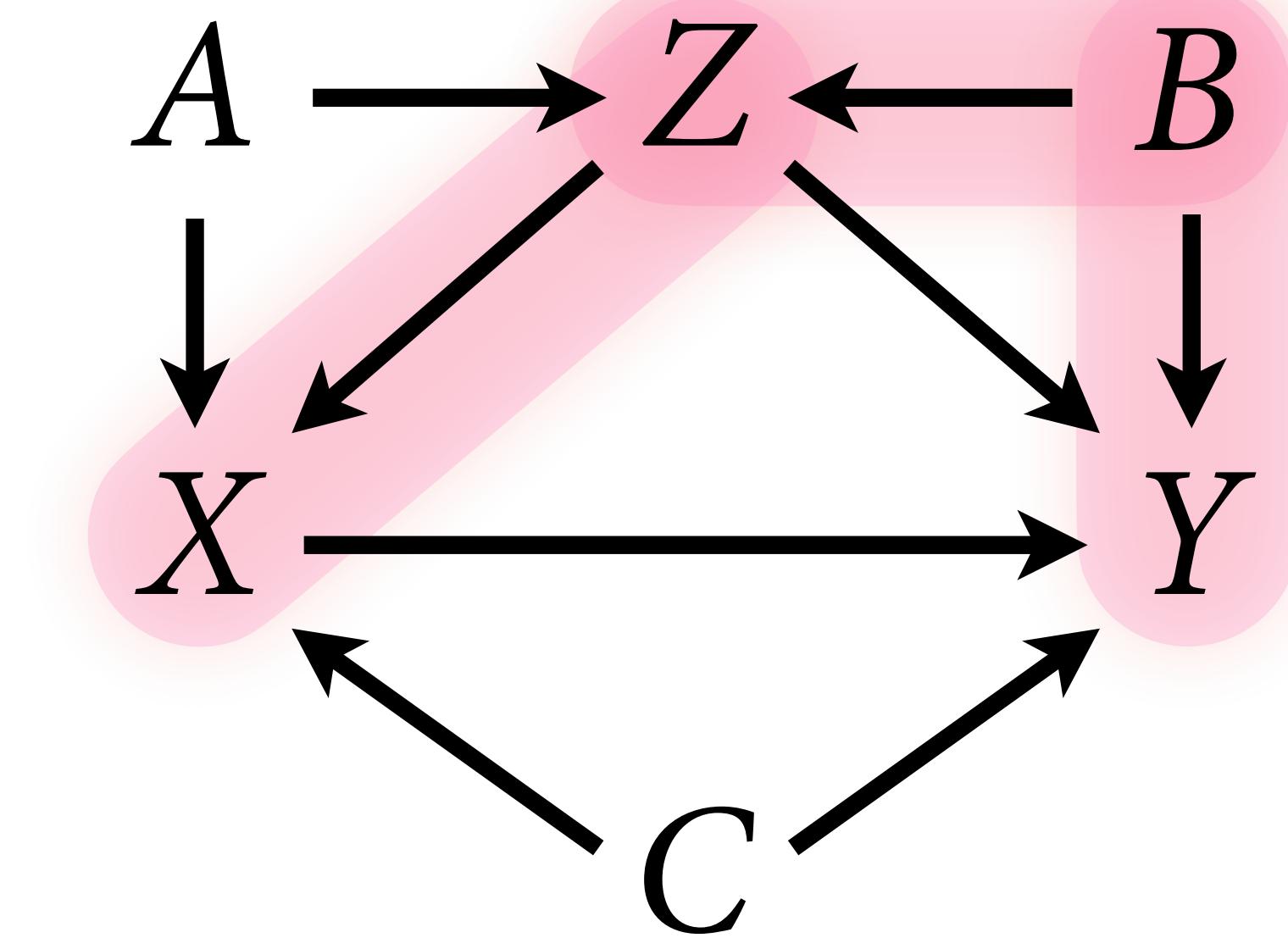
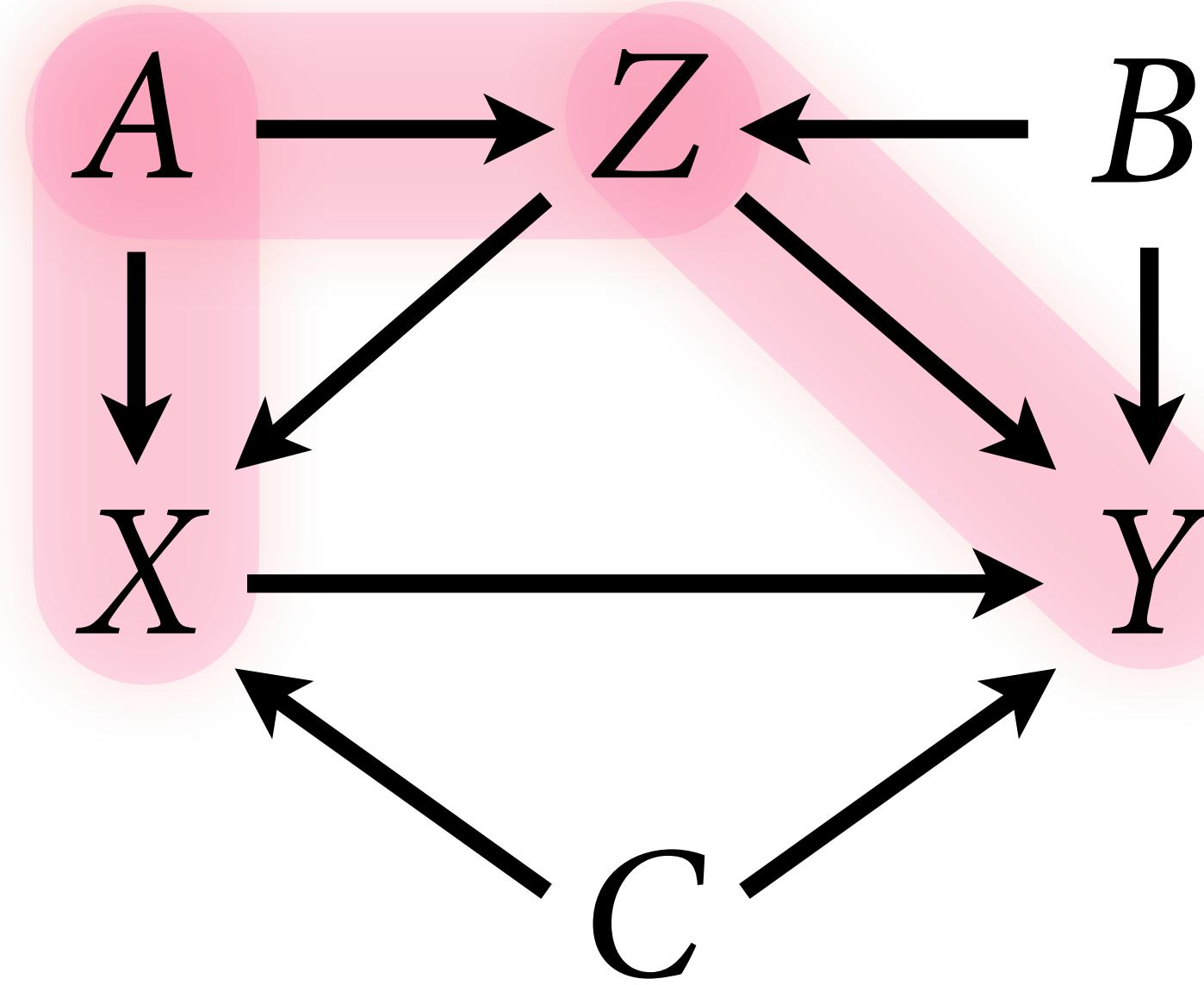
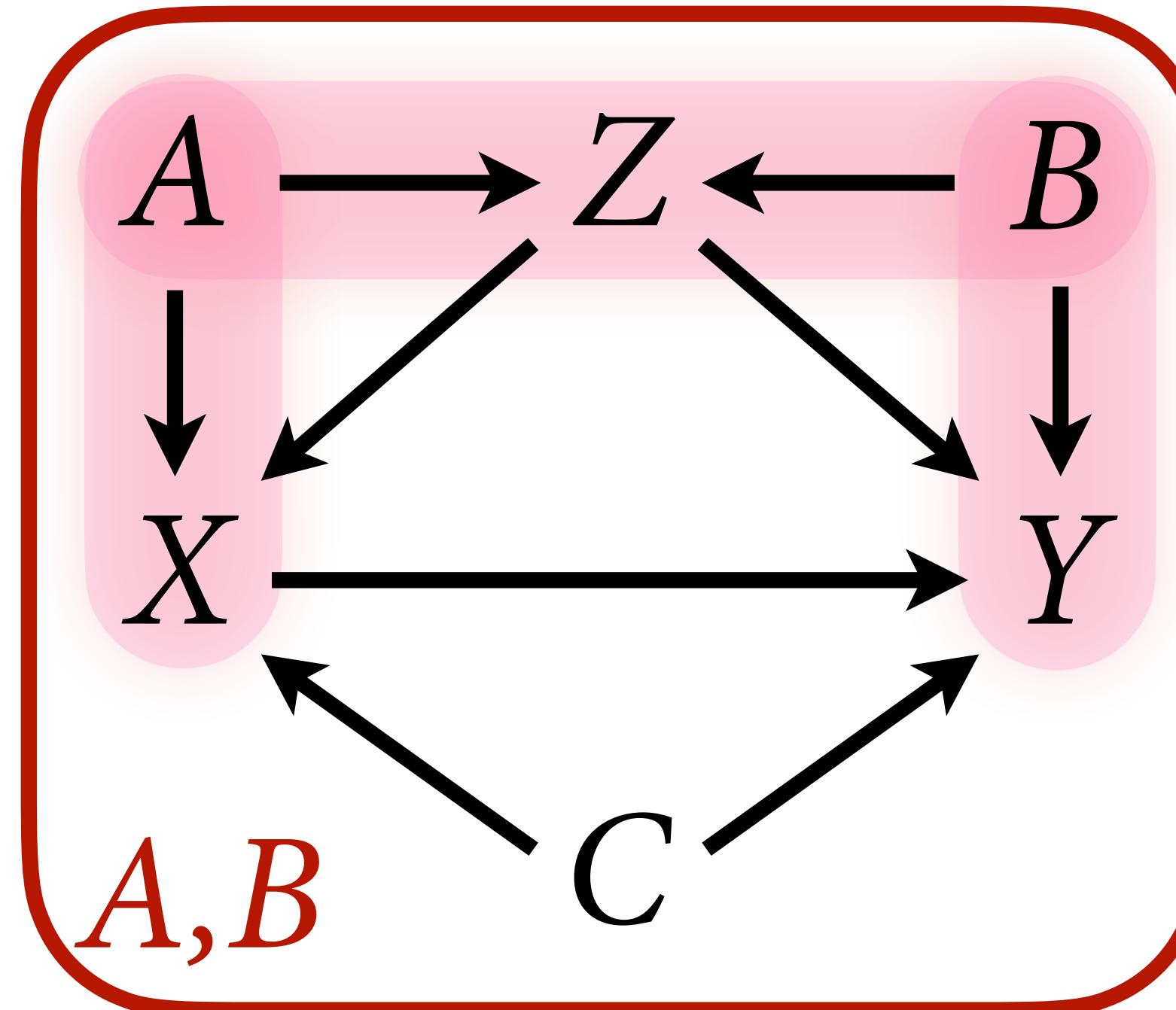
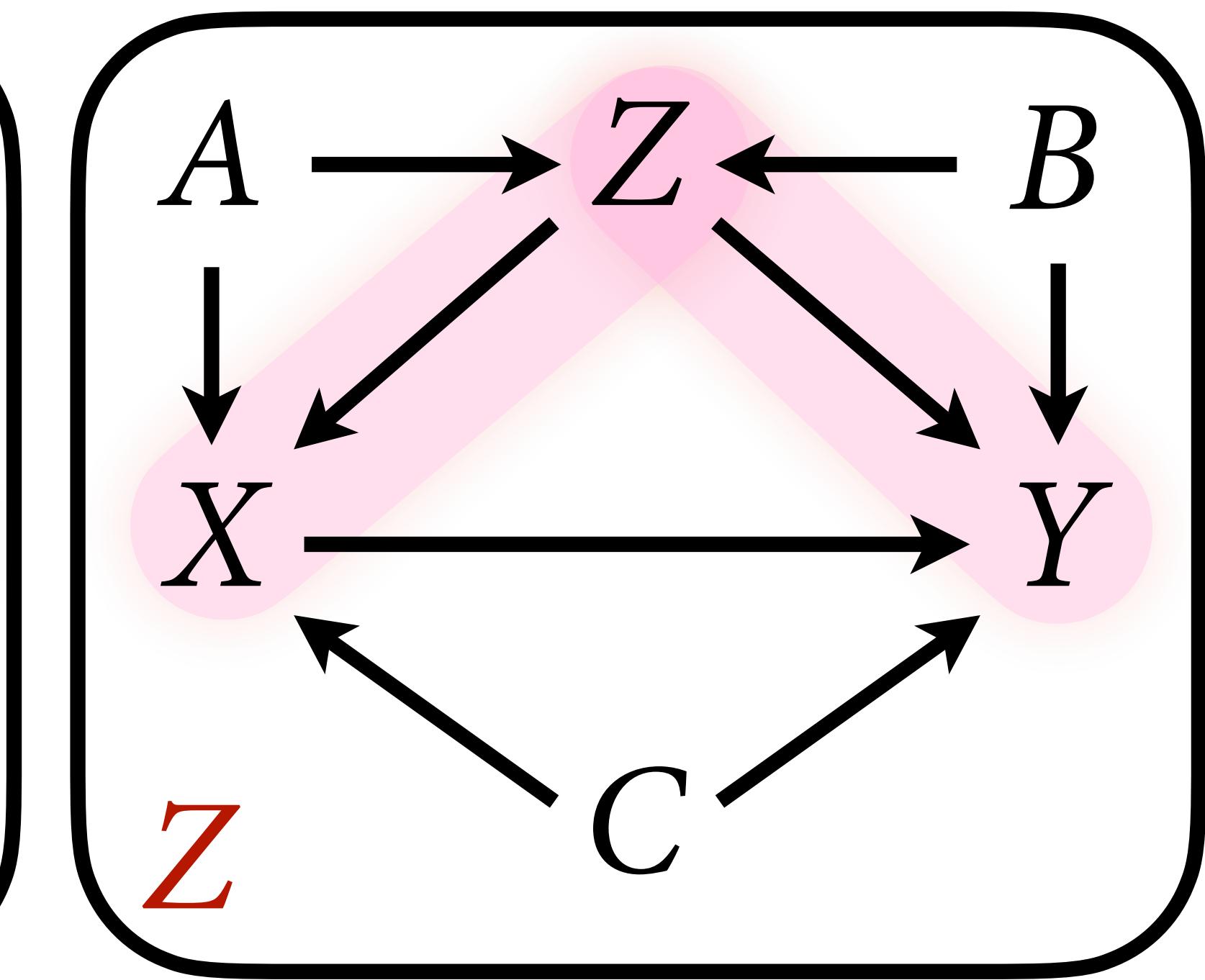
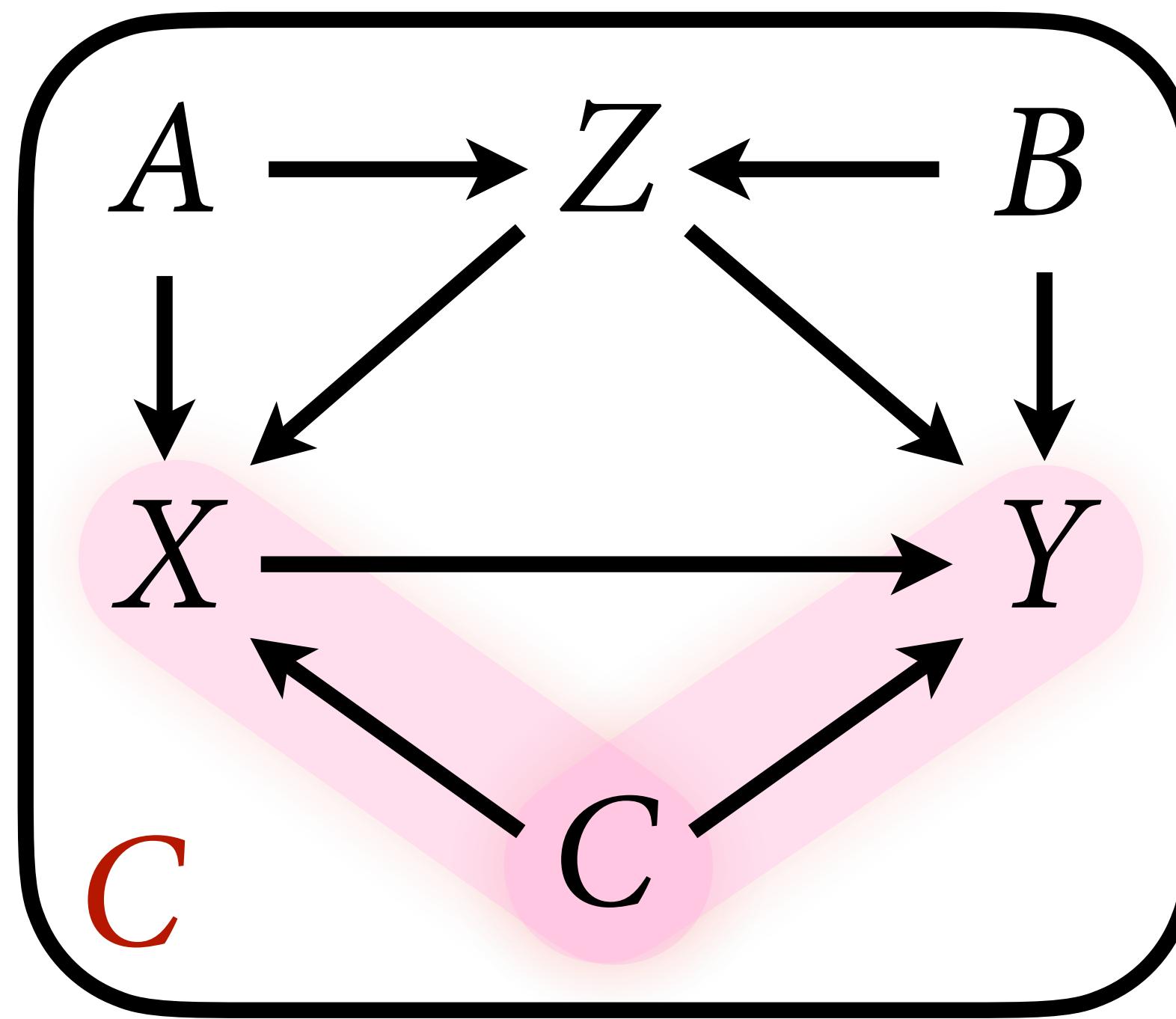
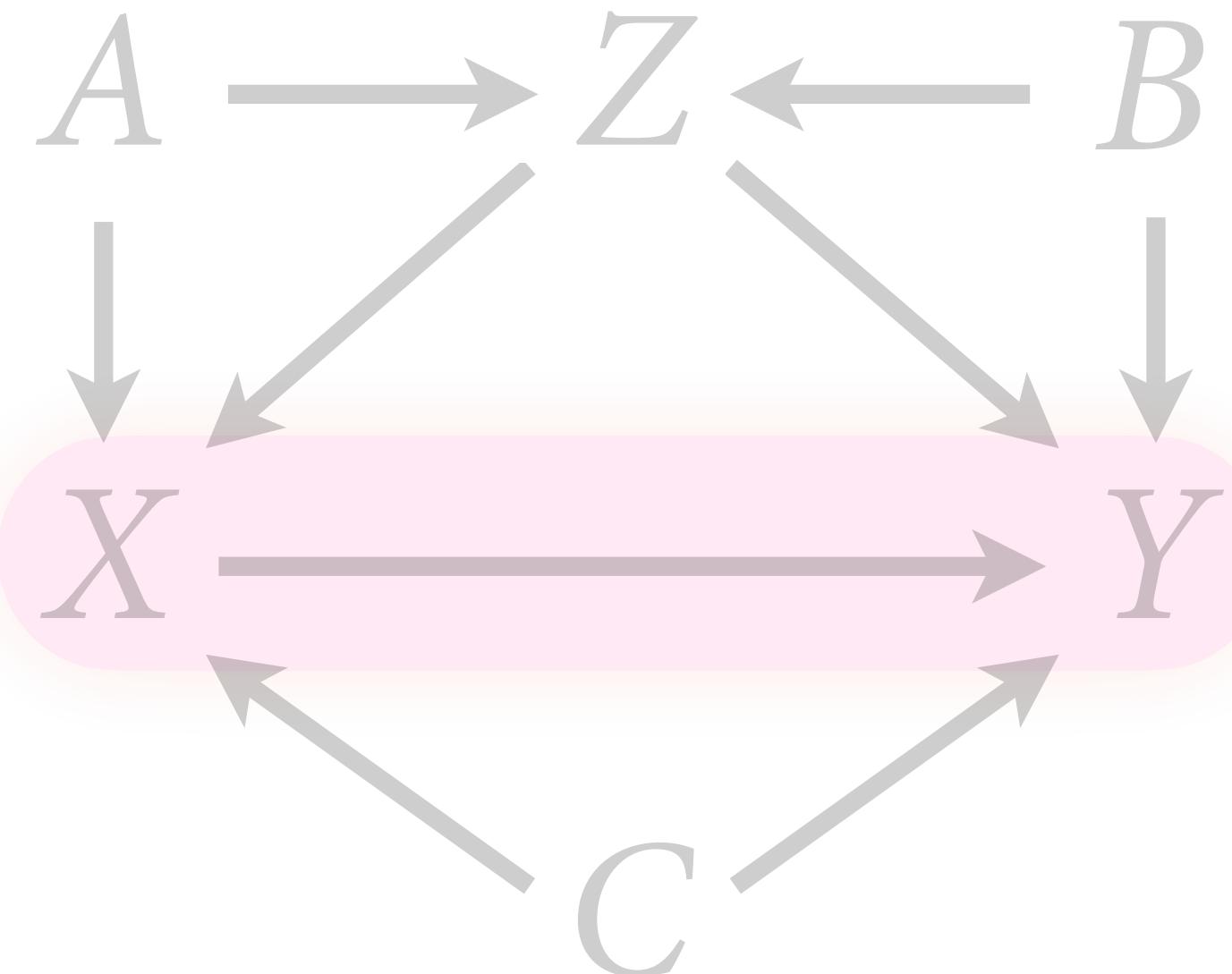
Backdoor path, opened by  $Z$   
 $A$  or  $B$  to close

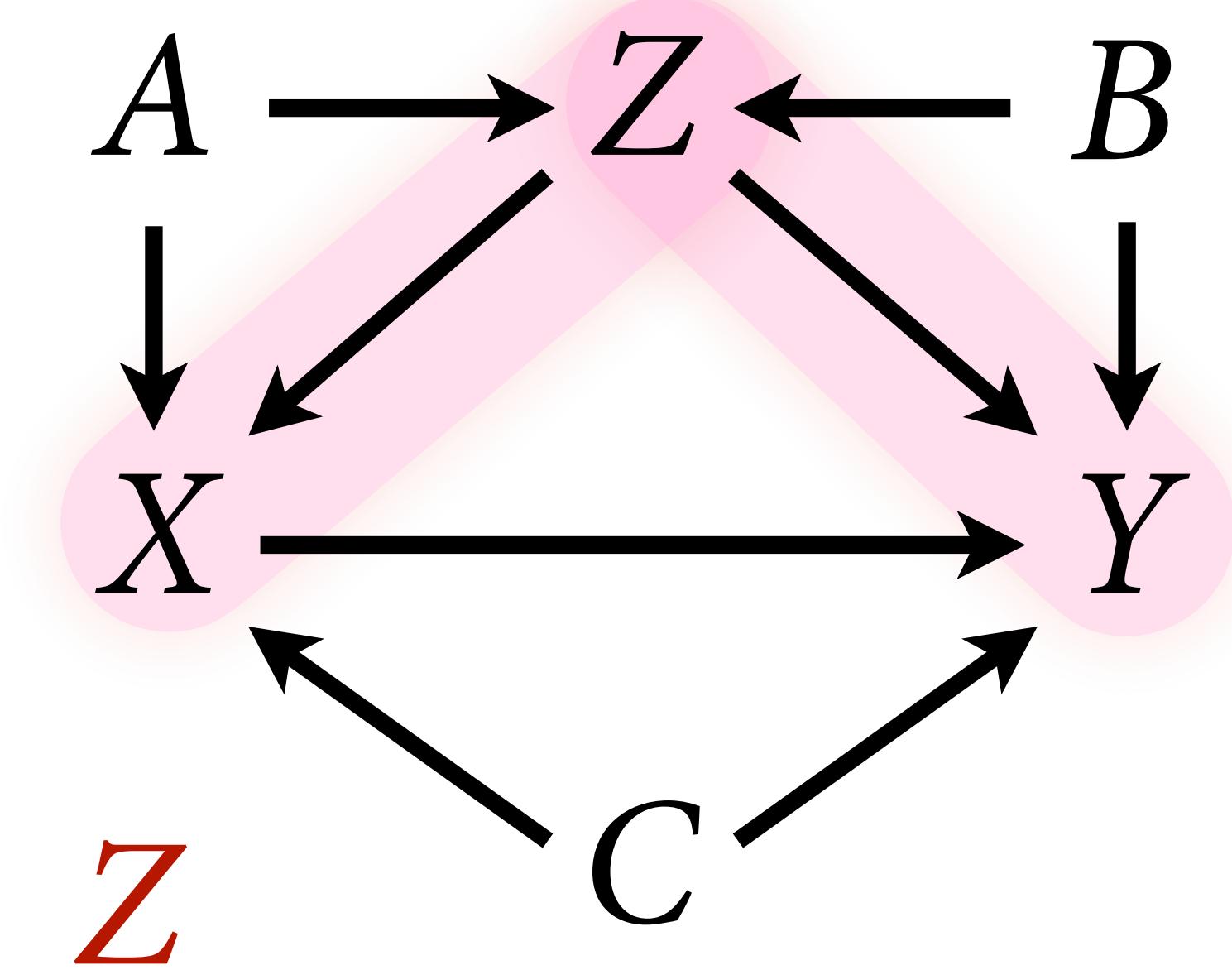
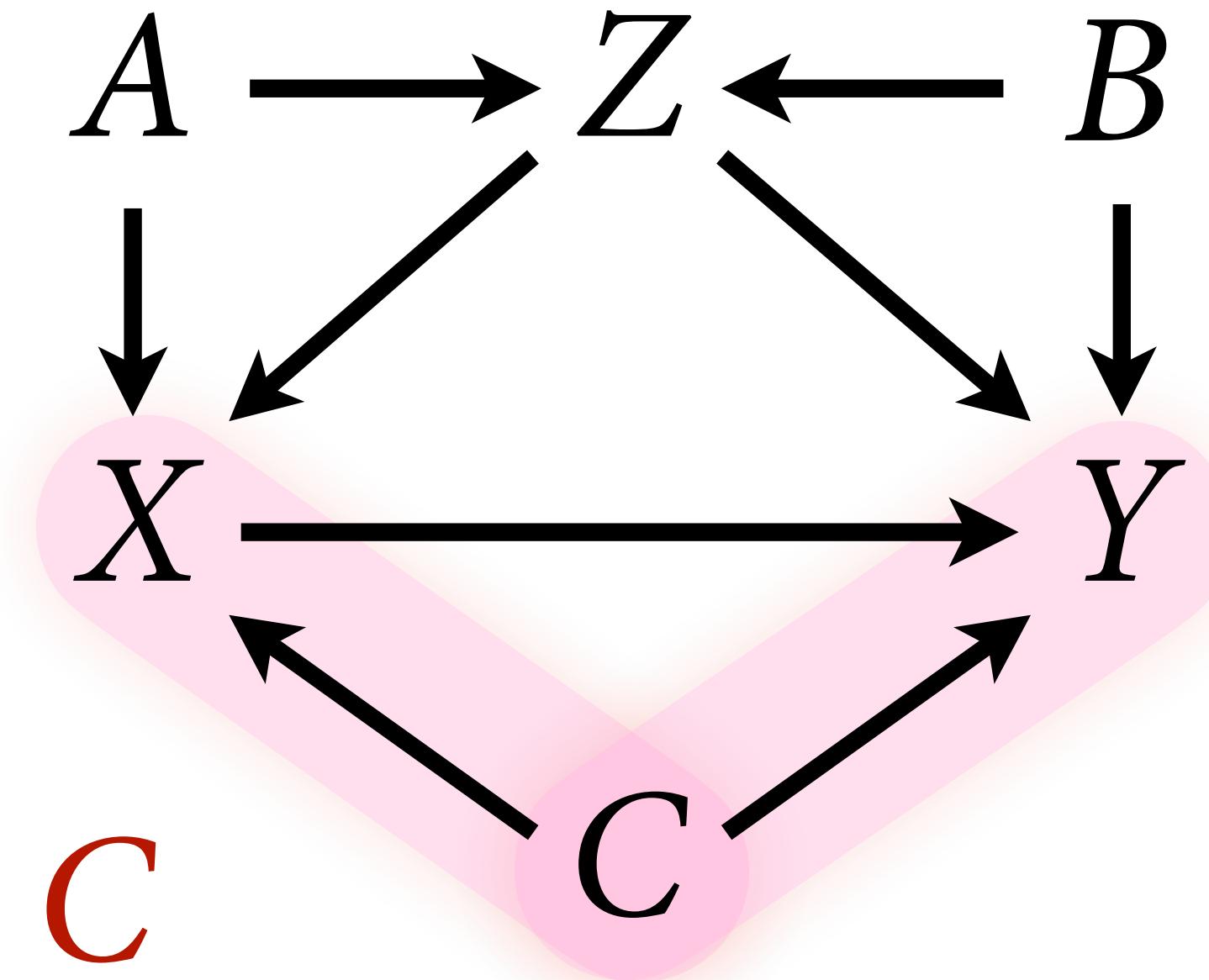
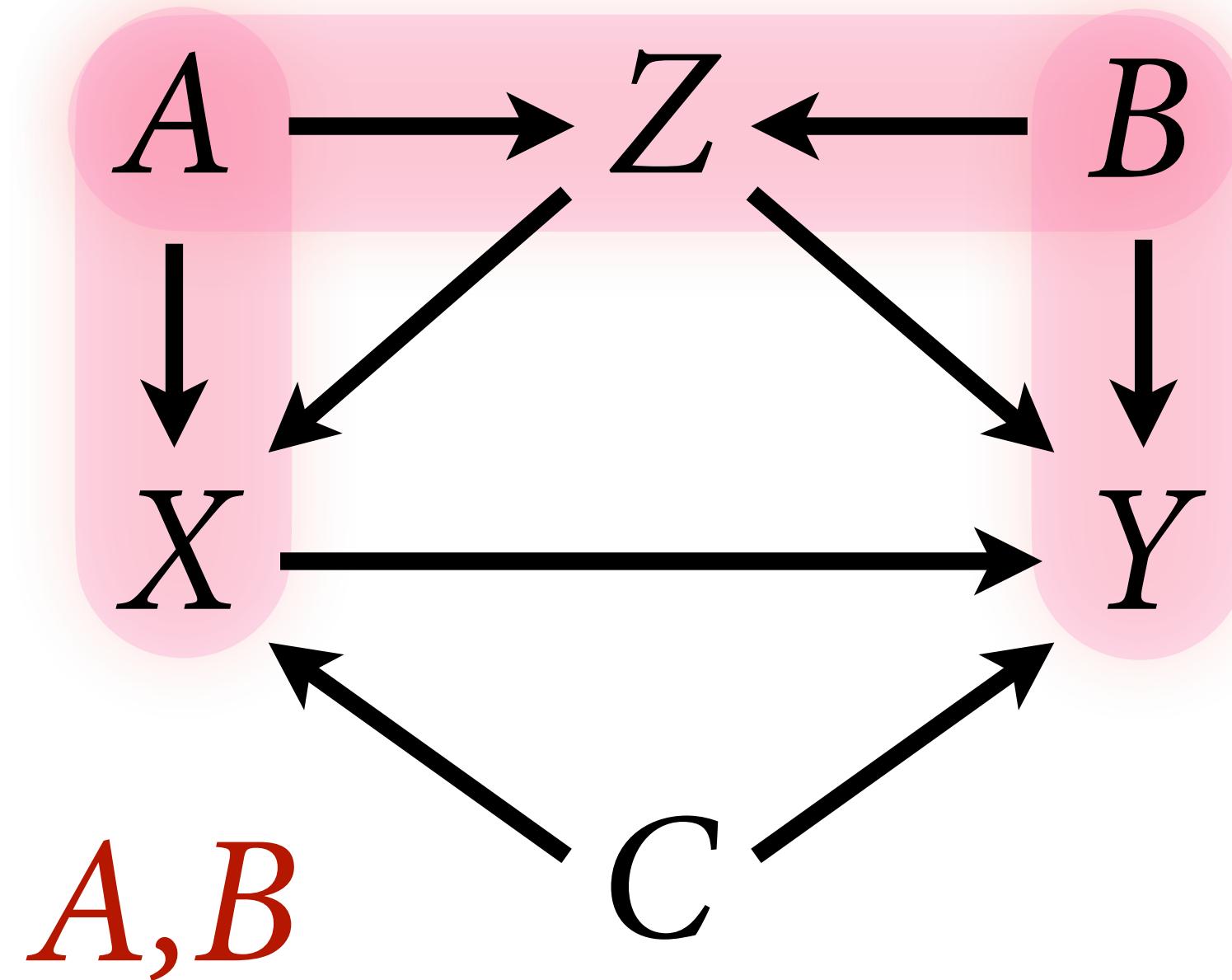




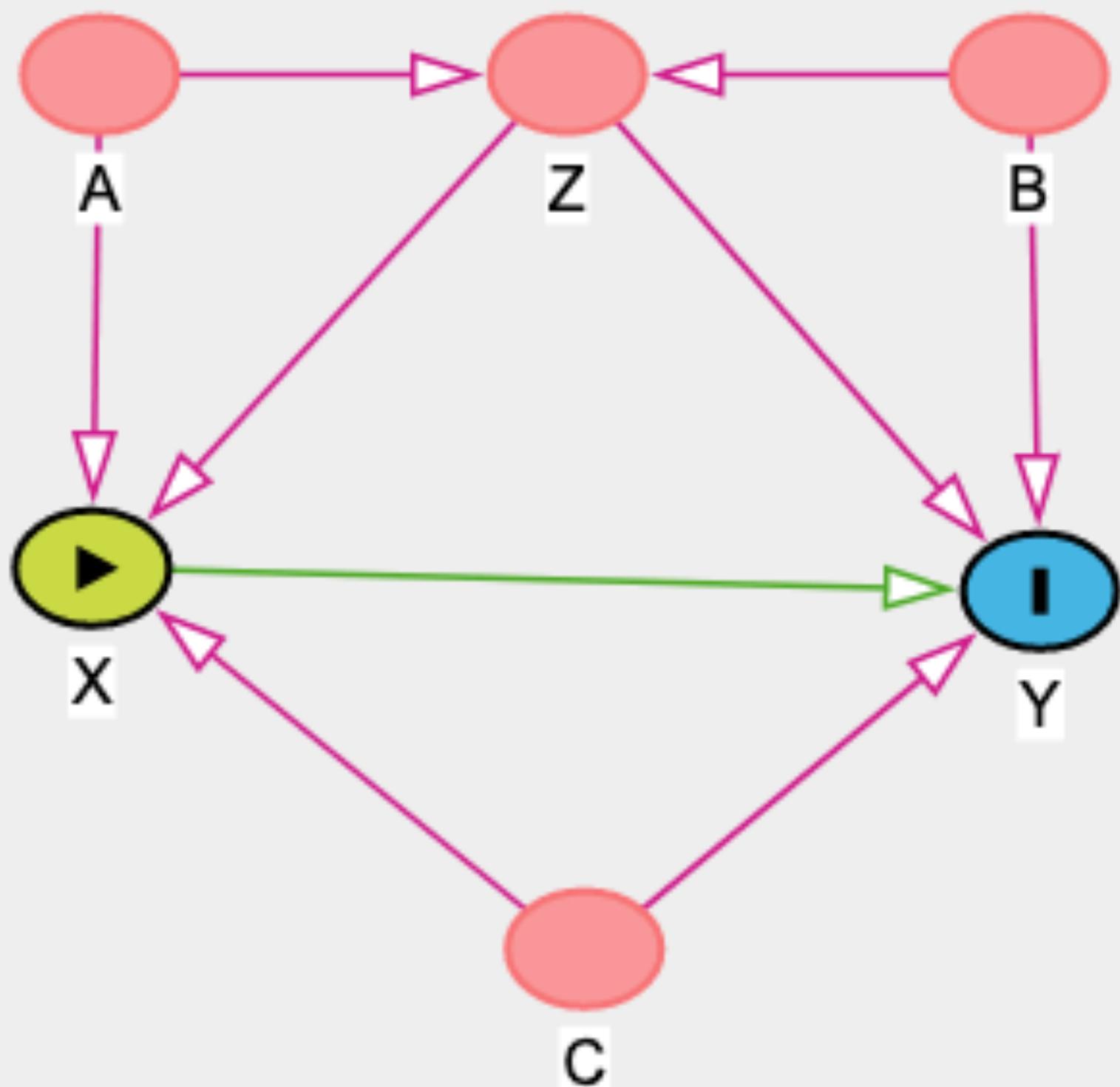


Backdoor path, open  
Close with  $A$  or  $Z$





Minimum adjustment set:  
*C, Z, and either A or B*  
 (B is better choice)



## ⊖ Causal effect identification

### Adjustment (total effect)

Minimal sufficient adjustment sets for estimating the total effect of X on Y:

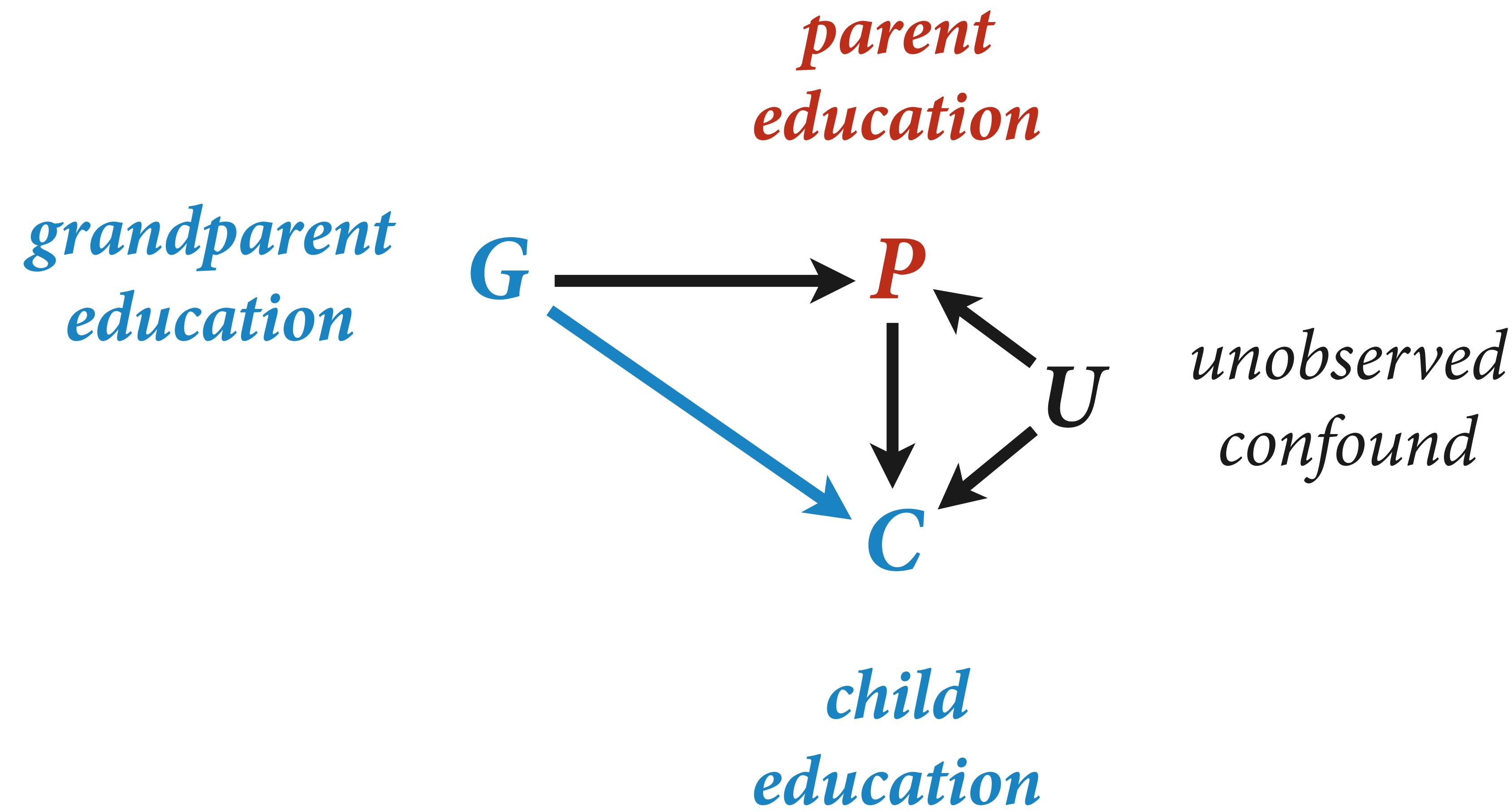
- A, C, Z
- B, C, Z

## ⊖ Testable implications

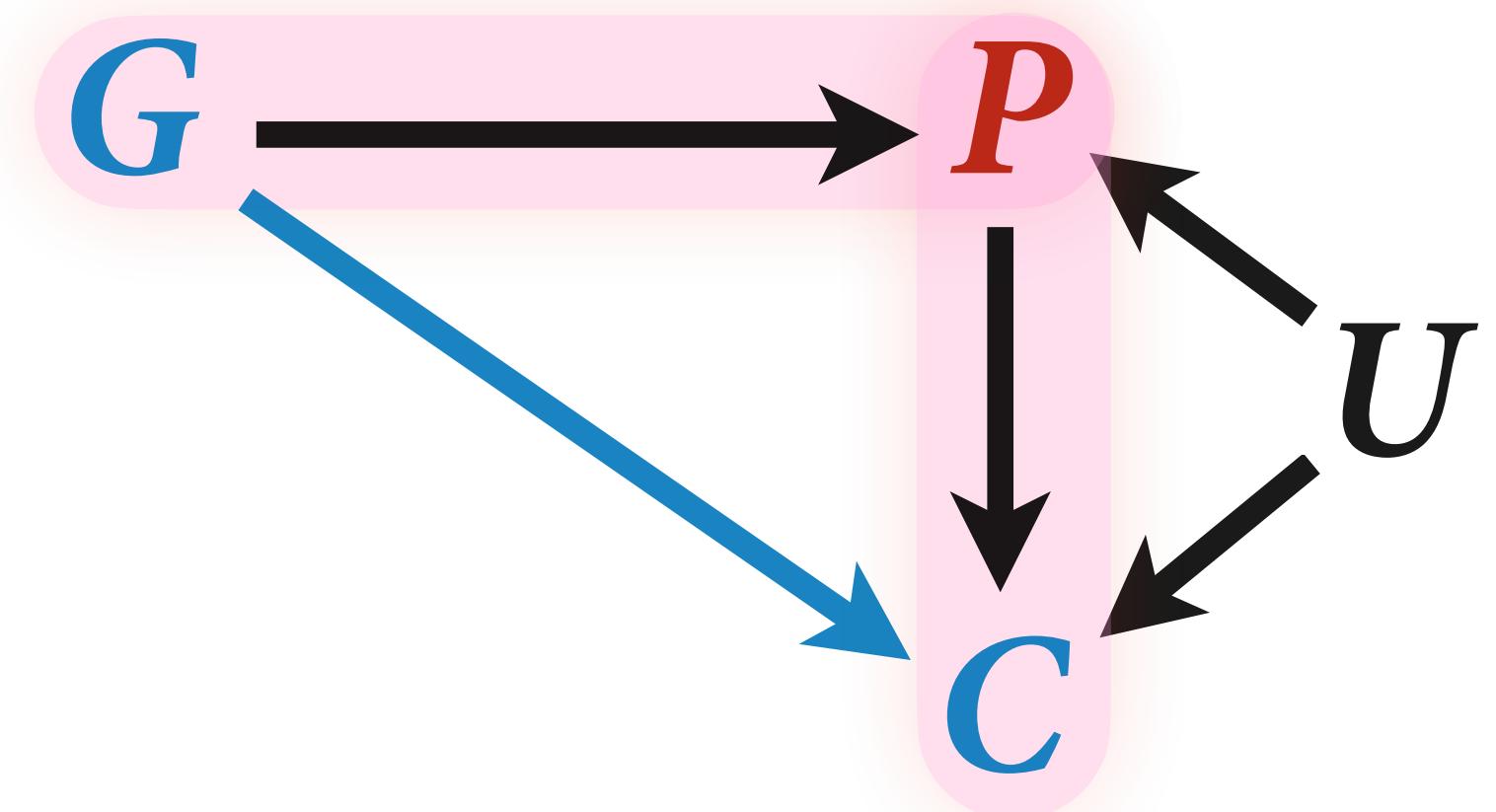
The model implies the following conditional independences:

- $X \perp\!\!\!\perp B \mid A, Z$
- $Y \perp\!\!\!\perp A \mid B, C, X, Z$
- $A \perp\!\!\!\perp B$
- $A \perp\!\!\!\perp C$
- $B \perp\!\!\!\perp C$
- $Z \perp\!\!\!\perp C$

[Export R code](#)

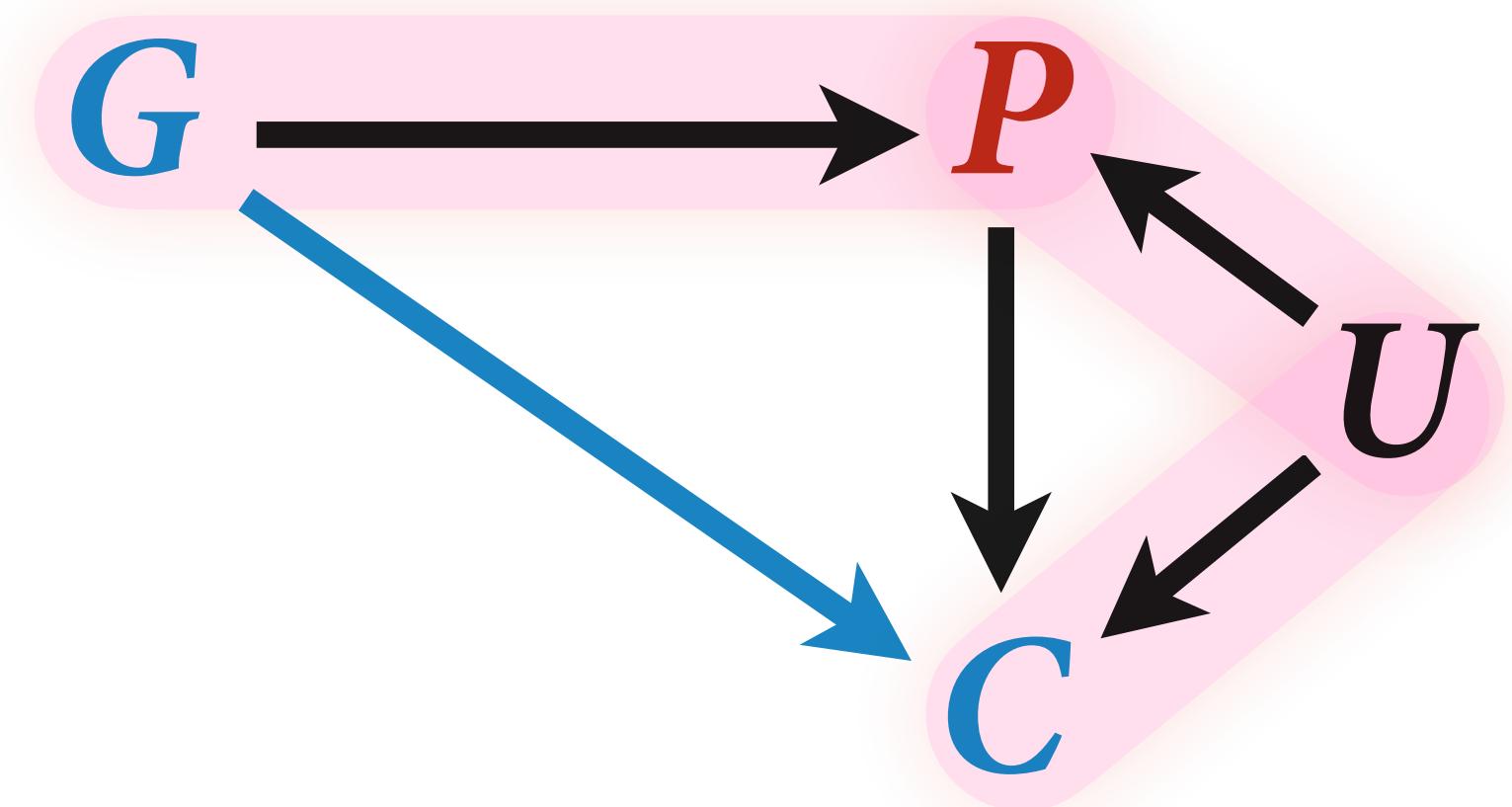


*Pipe:*  $G \rightarrow P \rightarrow C$



*P* is a mediator

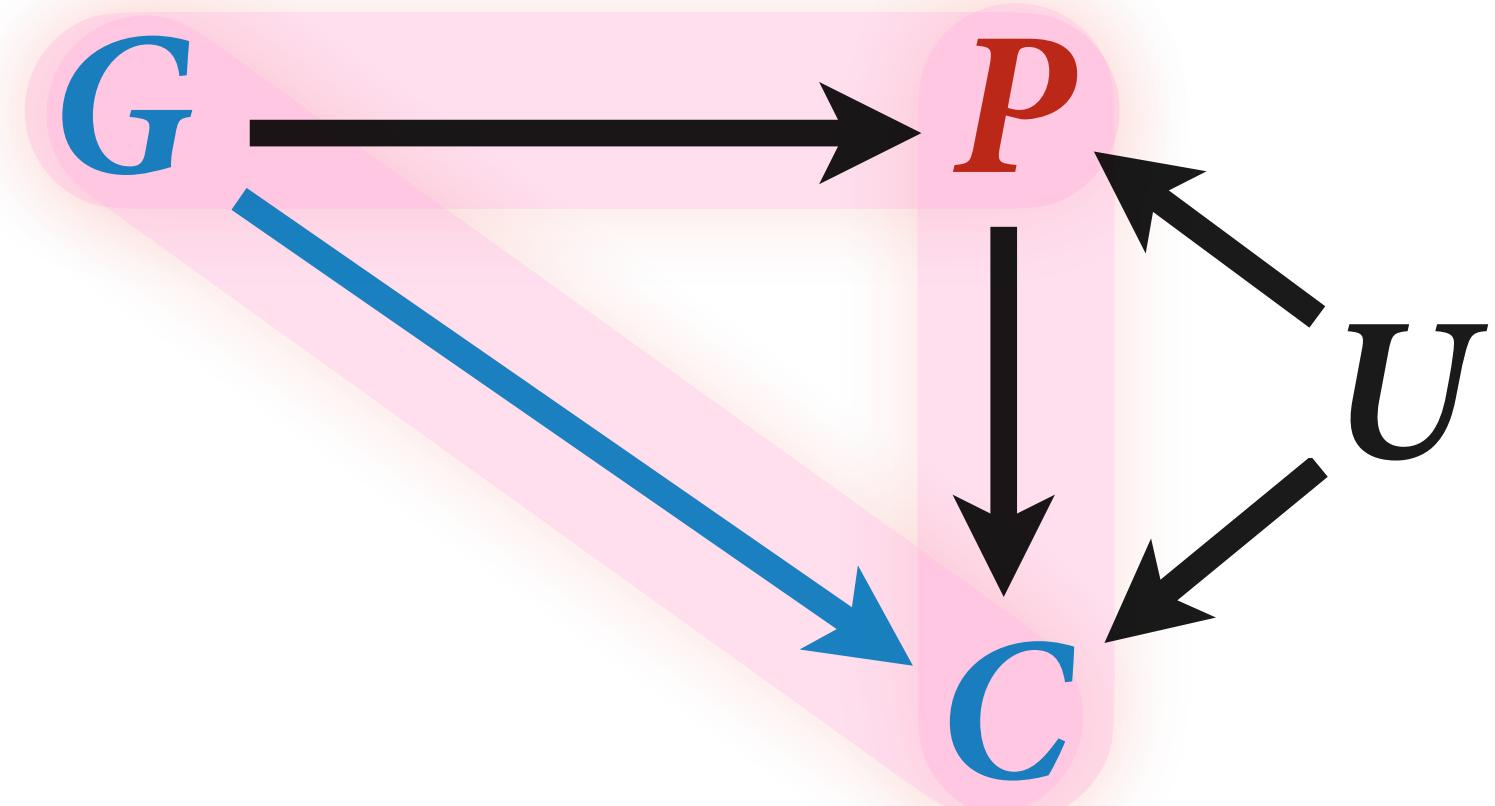
*Pipe:*  $G \rightarrow P \rightarrow C$



*P* is a collider

*Fork:*  $C \leftarrow U \rightarrow P$

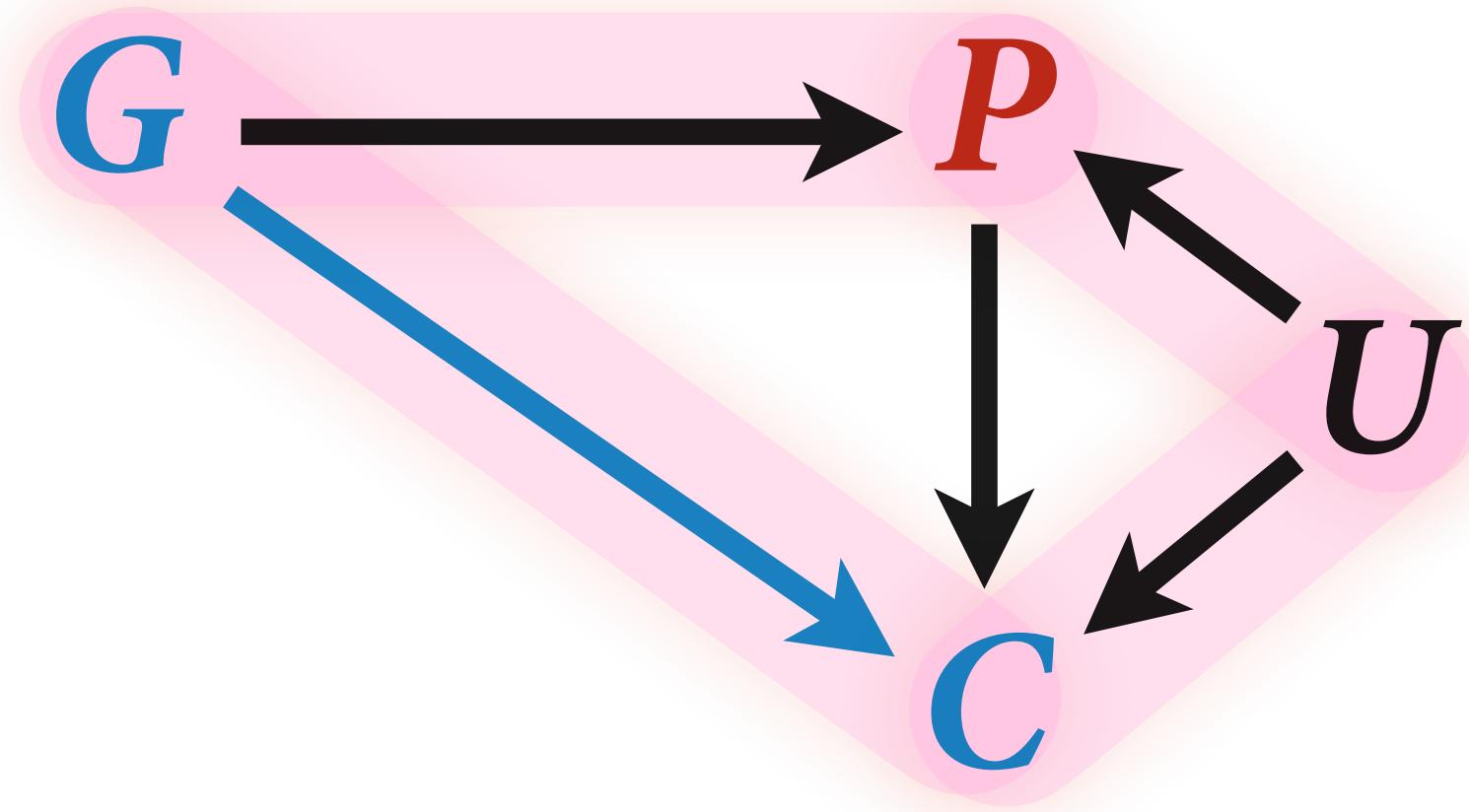
Can estimate **total**  
effect of  $G$  on  $C$



$$C_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_G G_i$$

Cannot estimate  
**direct** effect



$$C_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_G G_i + \beta_P P_i$$

# Backdoor Criterion

do-calc more than backdoors & adjustment sets

Full Luxury Bayes: use all variables, but in separate sub-models instead of single regression

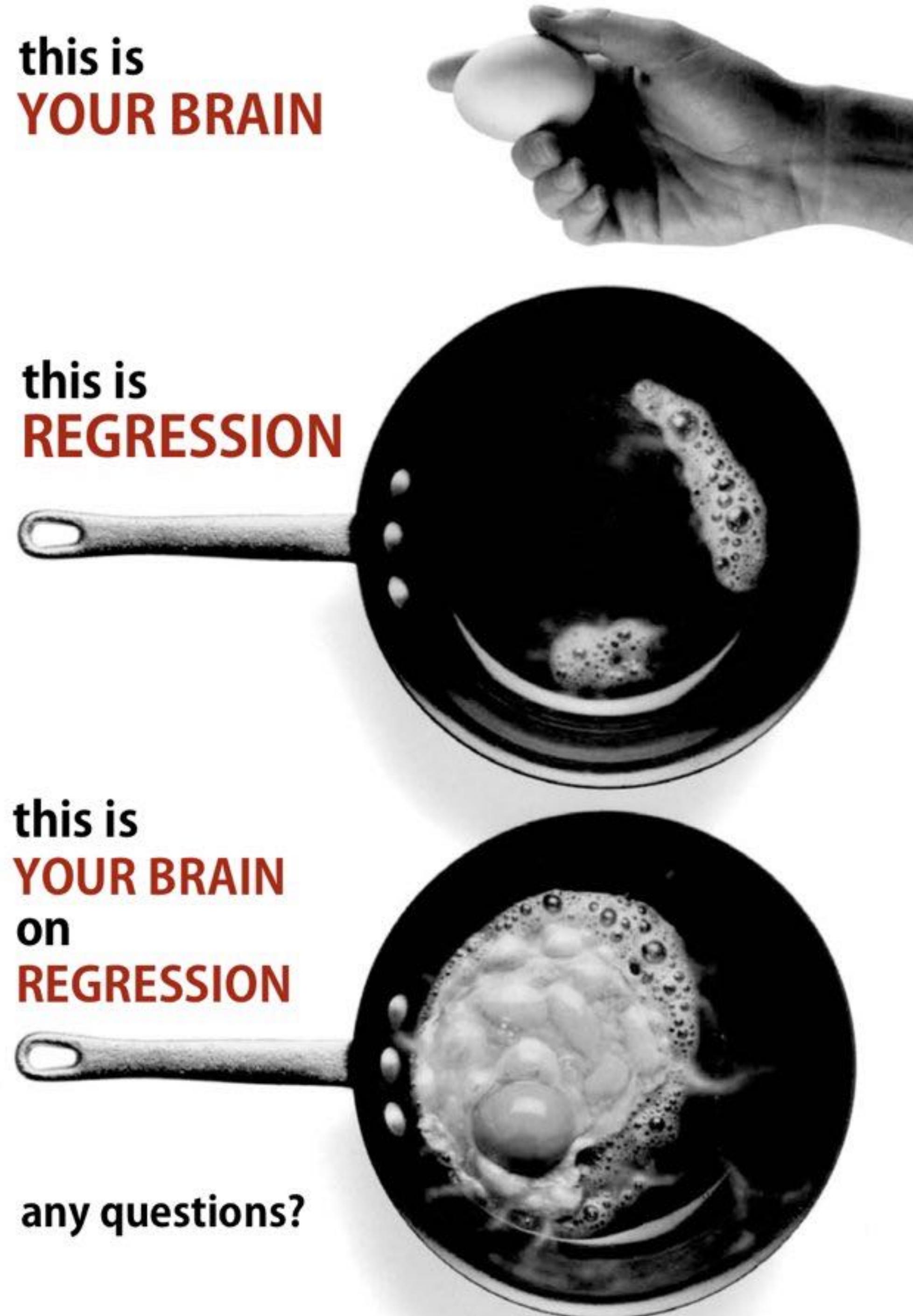
do-calc less demanding: finds relevant variables; saves us having to make some assumptions; not always a regression

this is  
**YOUR BRAIN**

this is  
**REGRESSION**

this is  
**YOUR BRAIN**  
on  
**REGRESSION**

any questions?



**PAUSE**

# Good & Bad Controls

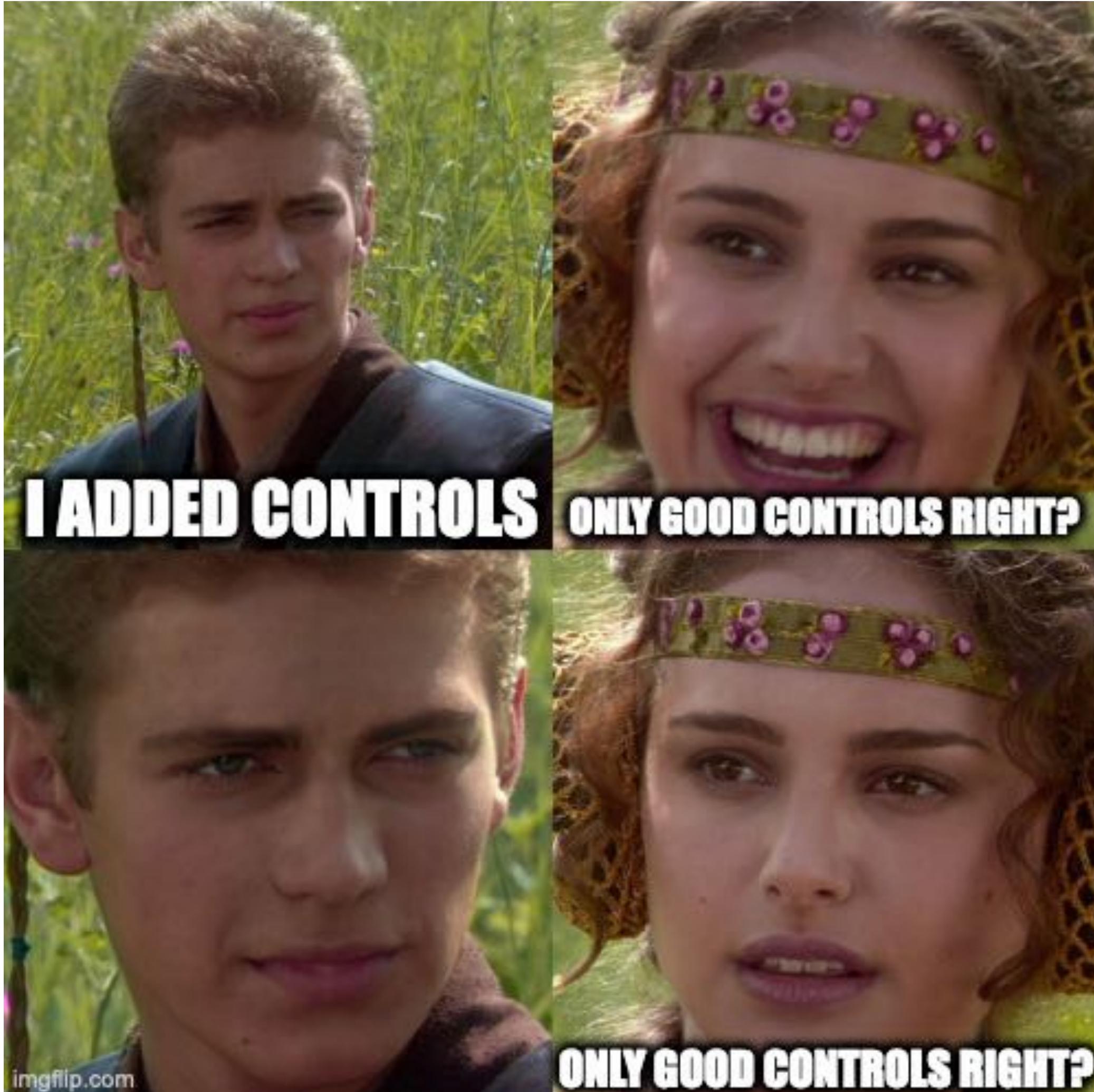
**“Control” variable:** Variable introduced to an analysis so that a causal estimate is possible

Common **wrong** heuristics for choosing control variables

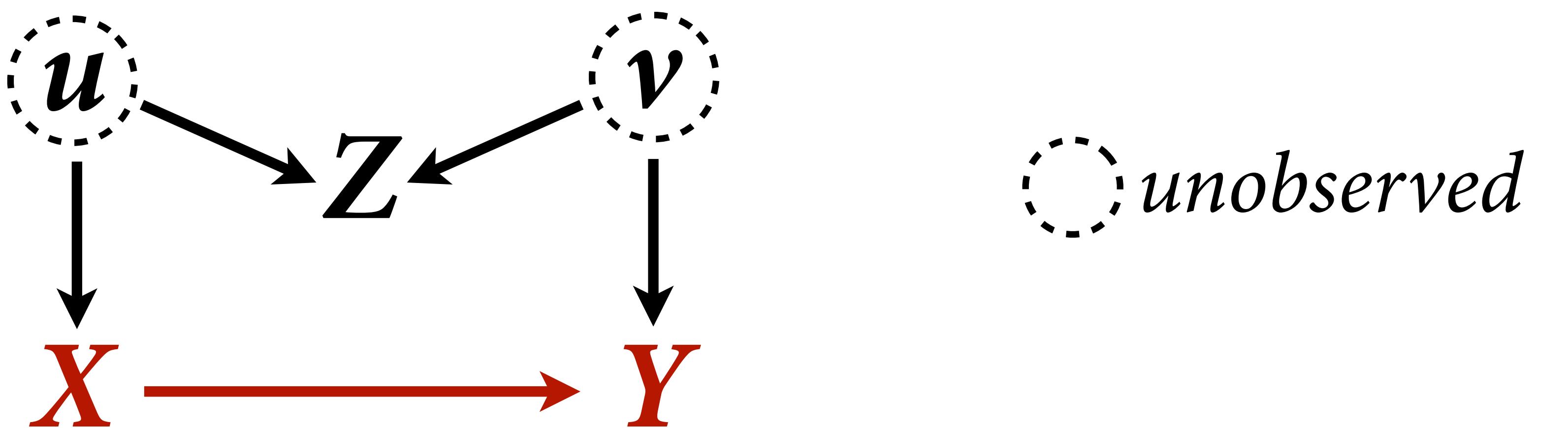
Anything in the spreadsheet **YOLO!**

Any variables not highly **collinear**

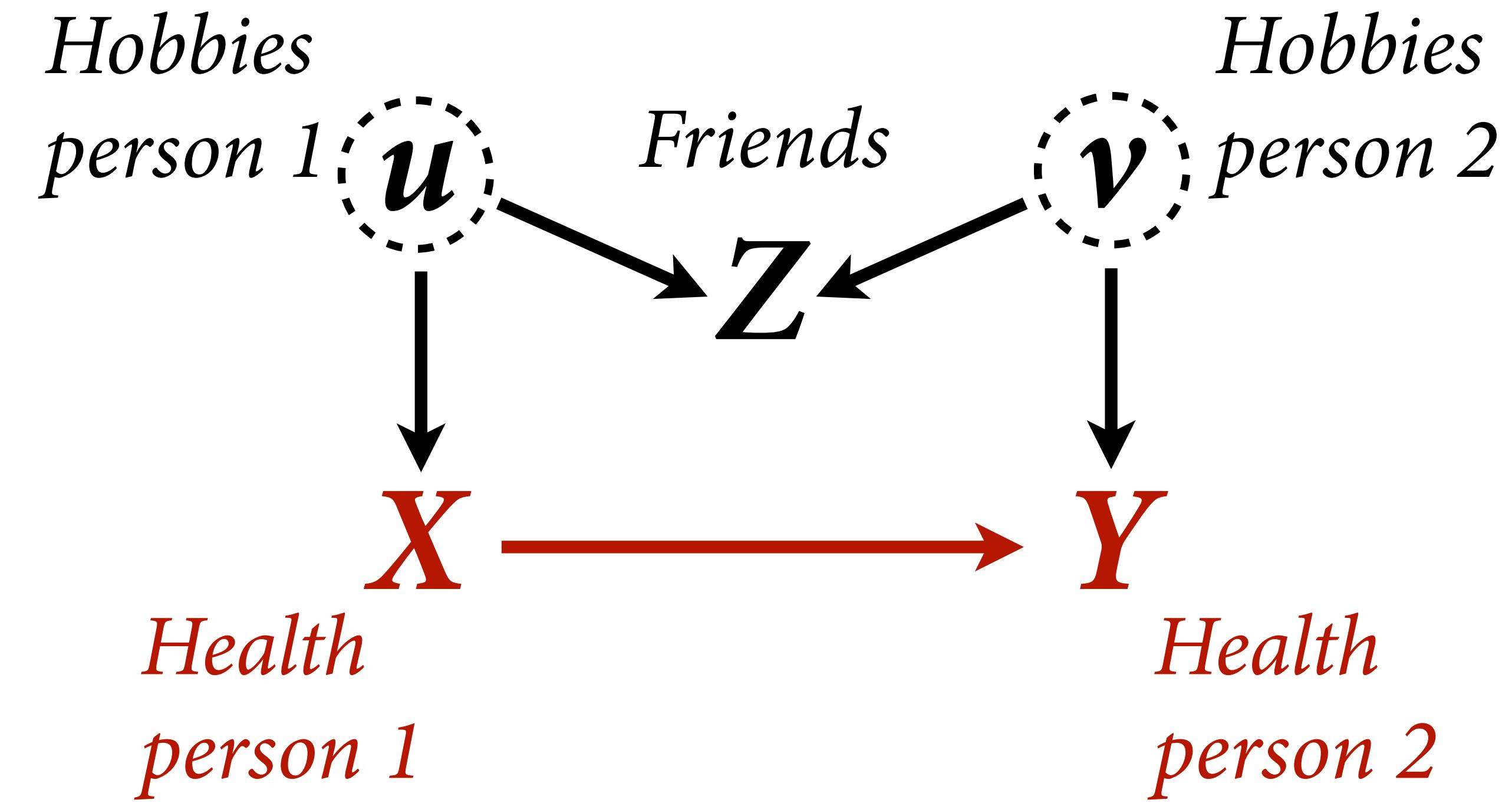
Any **pre-treatment** measurement (baseline)



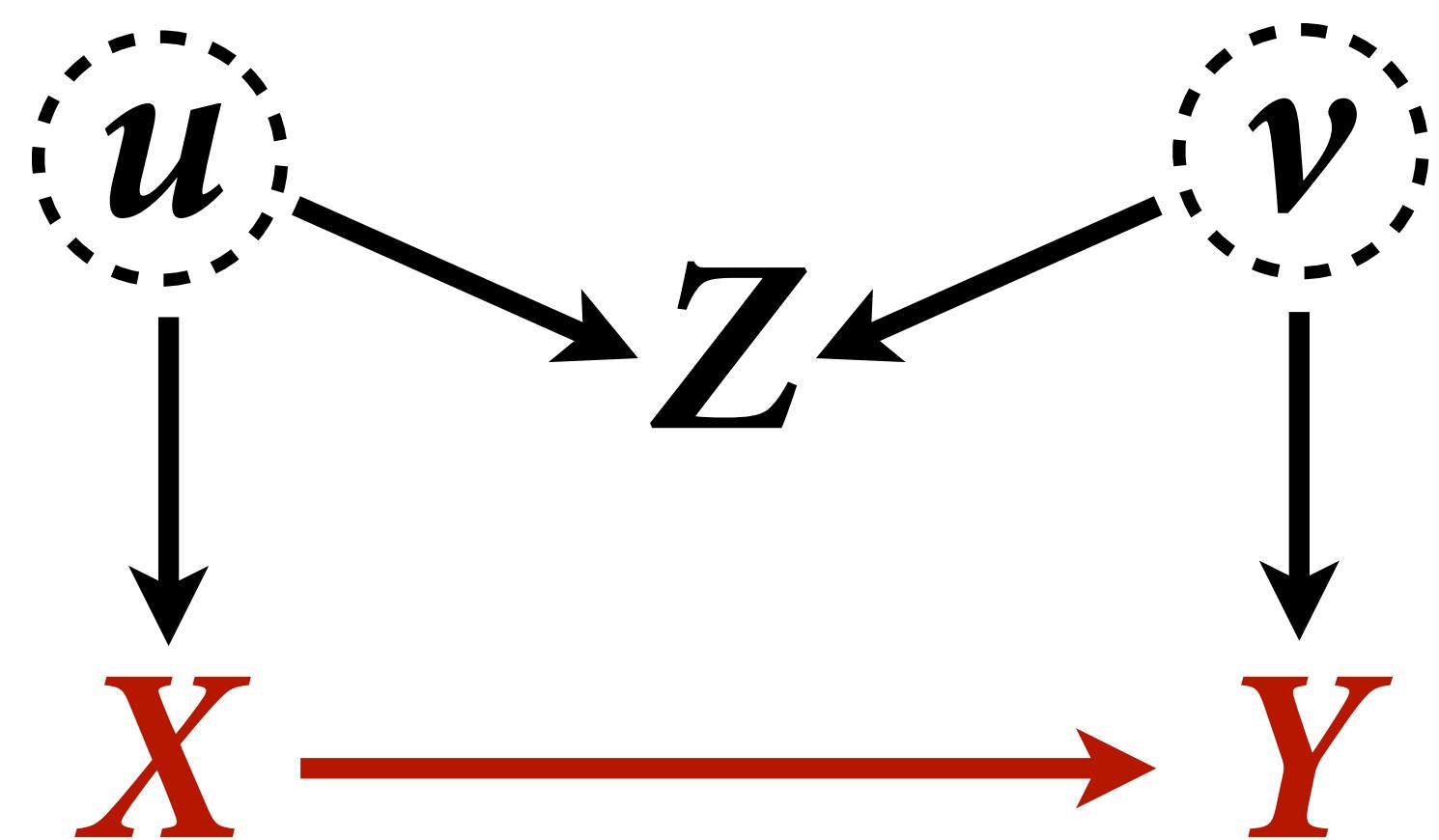
$$X \longrightarrow Y$$



*unobserved*

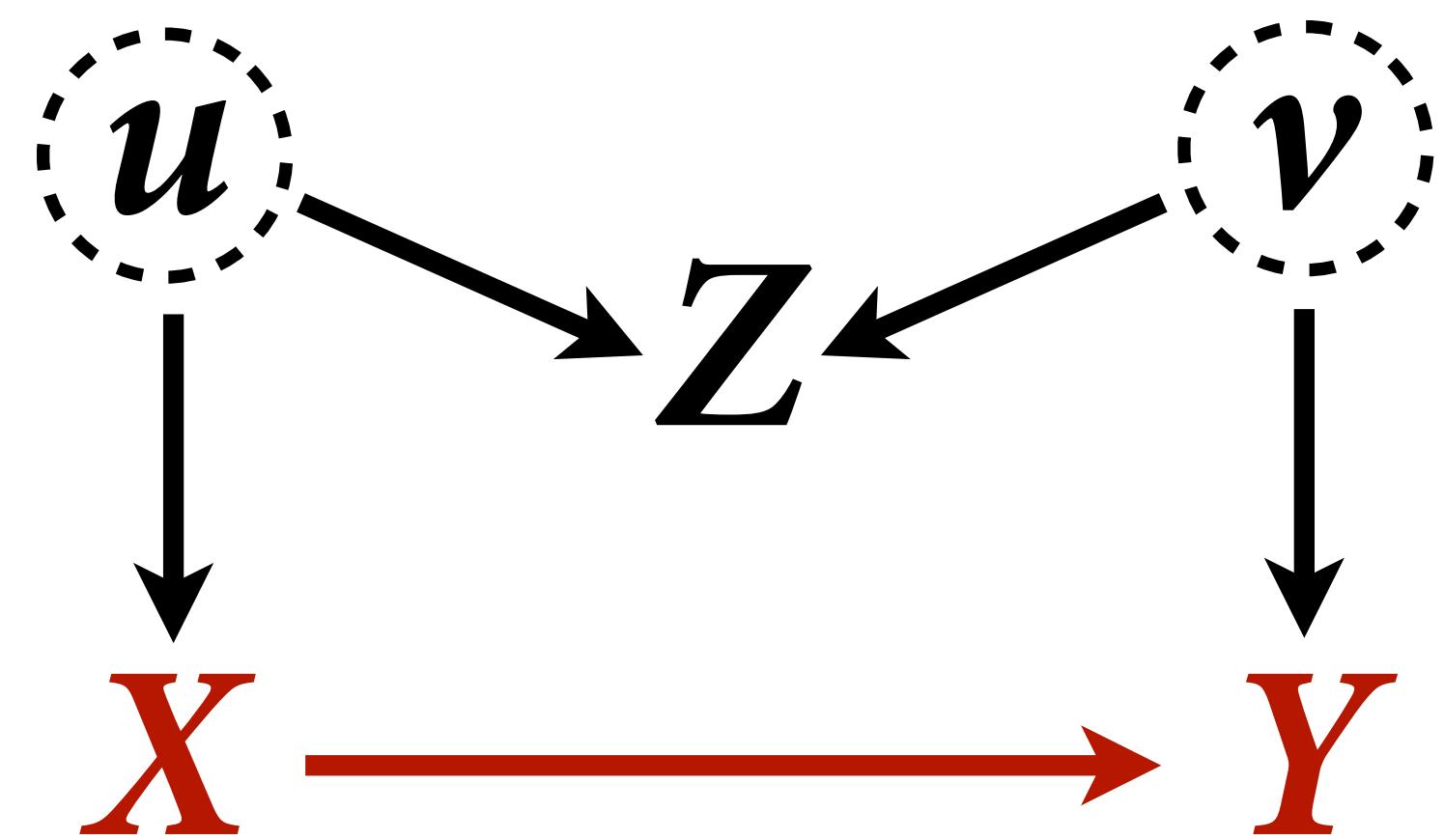


(1) List the paths



(1) List the paths

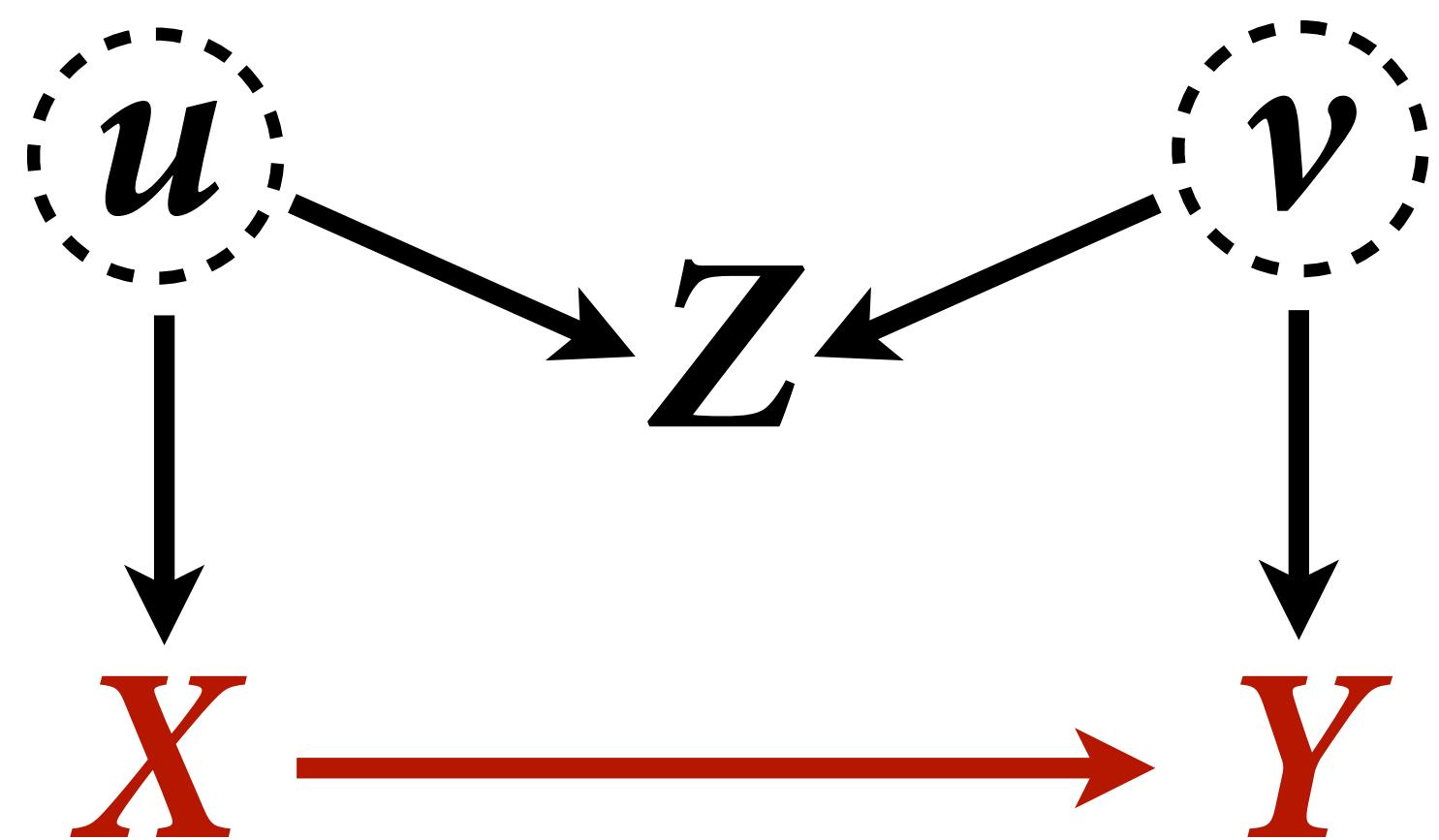
$X \rightarrow Y$



(1) List the paths

$$X \rightarrow Y$$

$$X \leftarrow u \rightarrow Z \leftarrow v \rightarrow Y$$



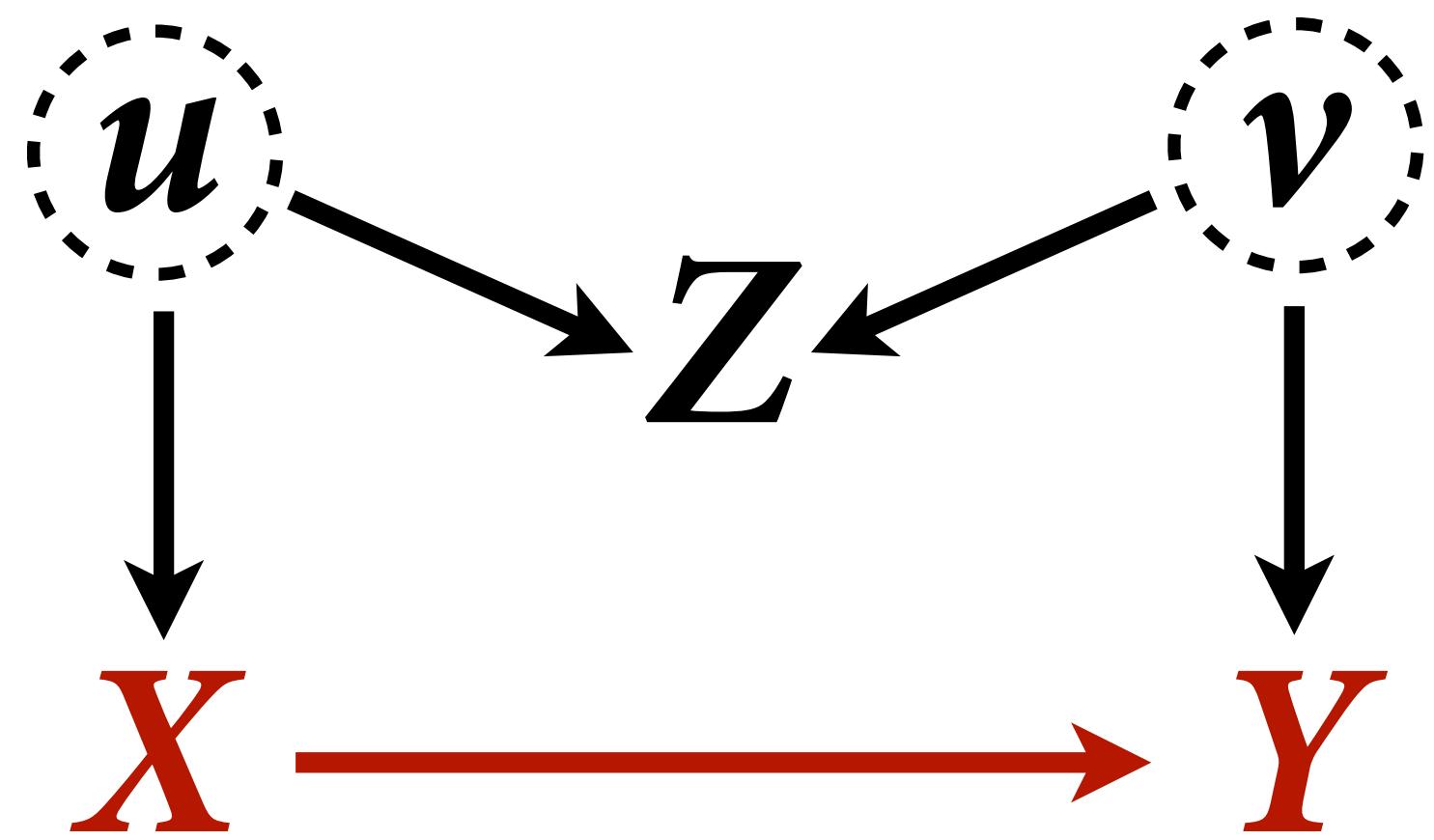
(1) List the paths    (2) Find backdoors

$$X \rightarrow Y$$

*frontdoor & open*

$$X \leftarrow u \rightarrow Z \leftarrow v \rightarrow Y$$

*backdoor & closed*



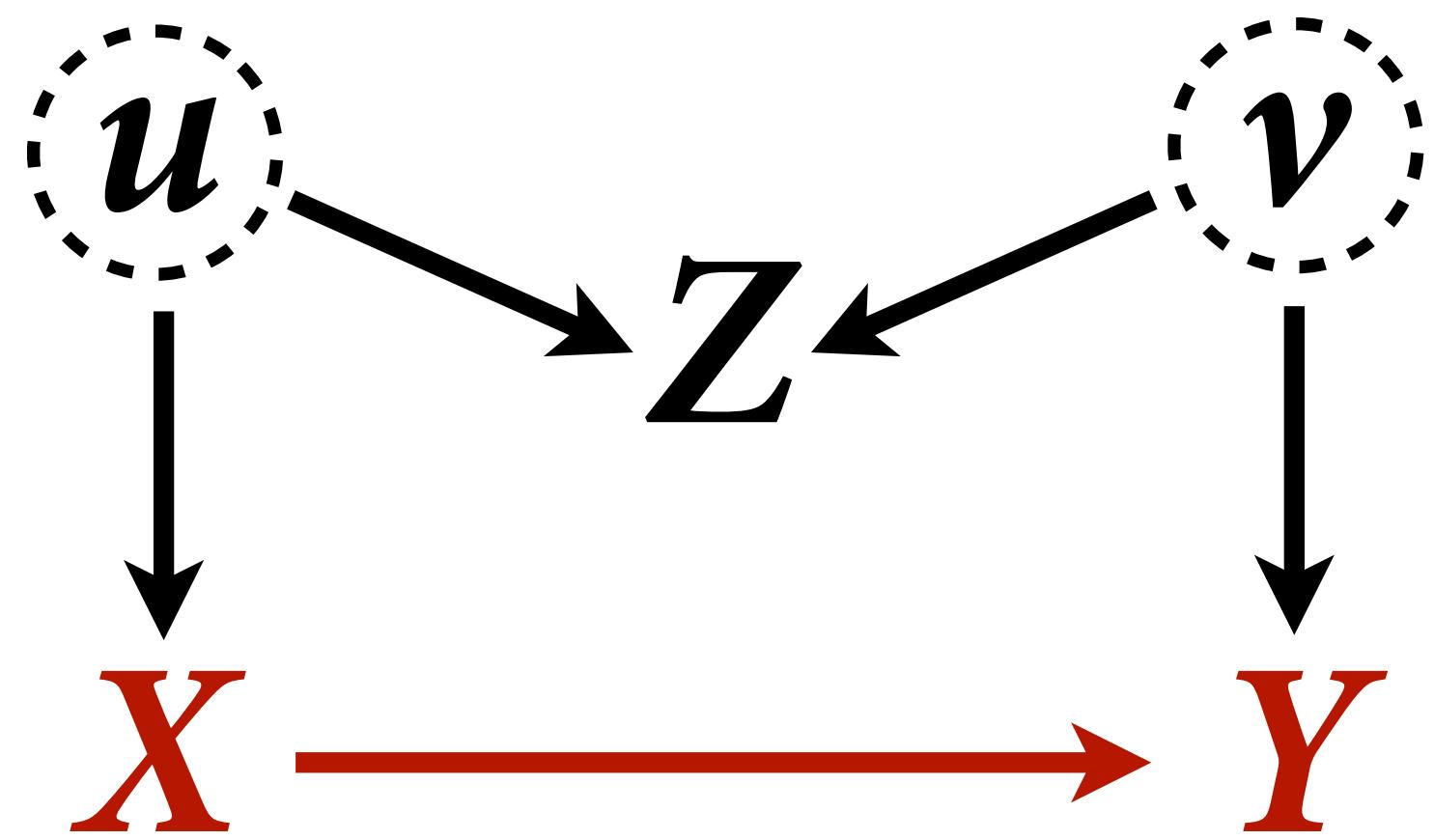
(1) List the paths    (2) Find backdoors

$$X \rightarrow Y$$

*frontdoor & open*

$$X \leftarrow u \rightarrow Z \leftarrow v \rightarrow Y$$

*backdoor & closed*



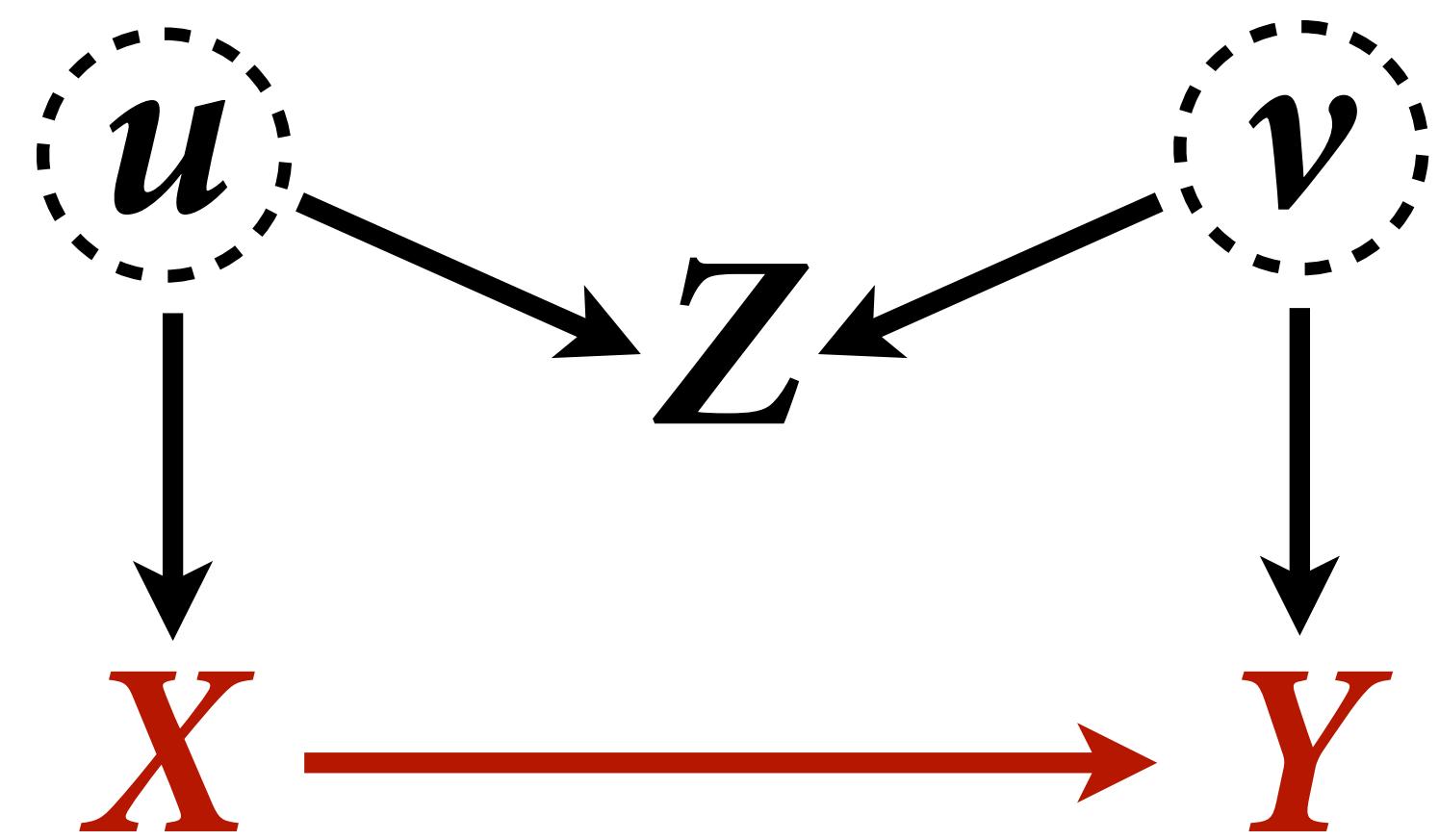
(1) List the paths    (2) Find backdoors    (3) Close backdoors

$$X \rightarrow Y$$

*frontdoor & open*

$$X \leftarrow u \rightarrow Z \leftarrow v \rightarrow Y$$

*backdoor & closed*

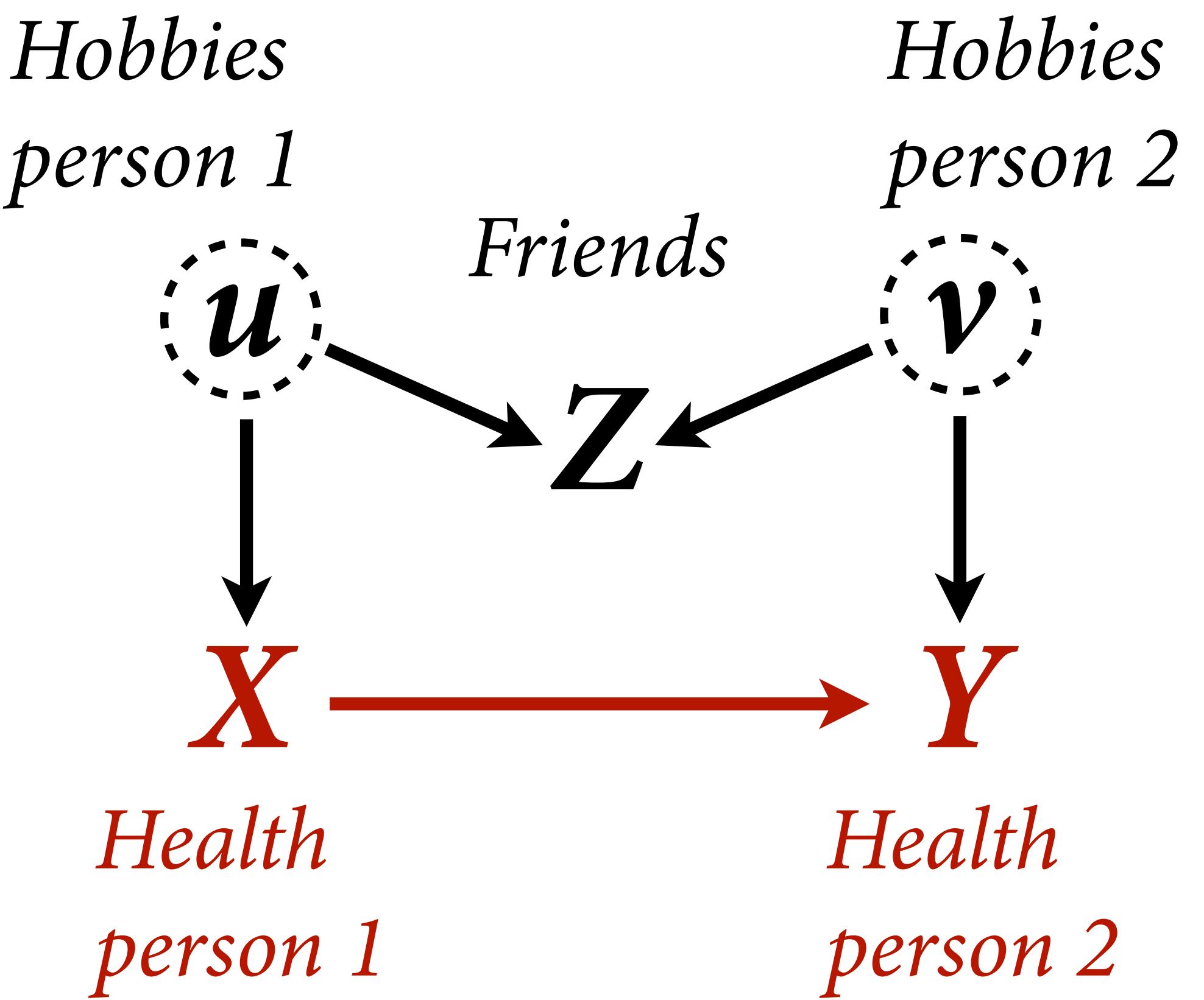


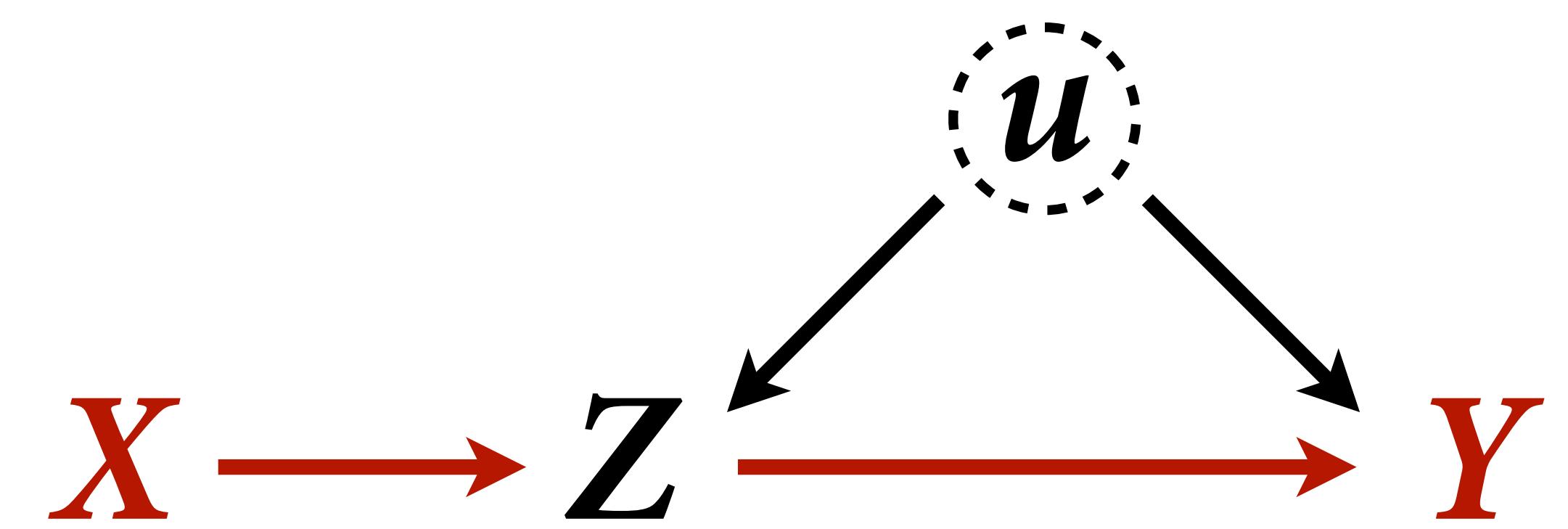
What happens if you stratify by  
 $Z$ ?

Opens the backdoor path

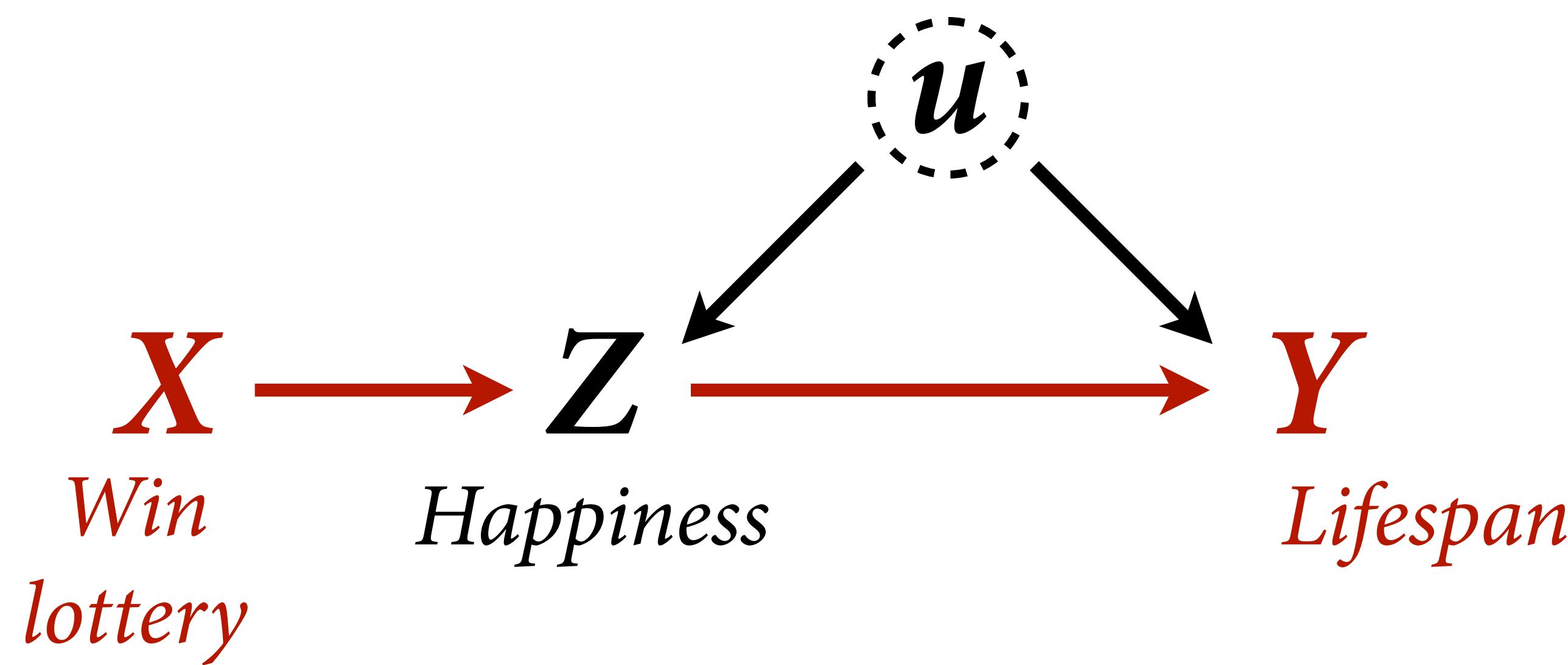
$Z$  could be a **pre-treatment**  
variable

Not safe to always control pre-  
treatment measurements





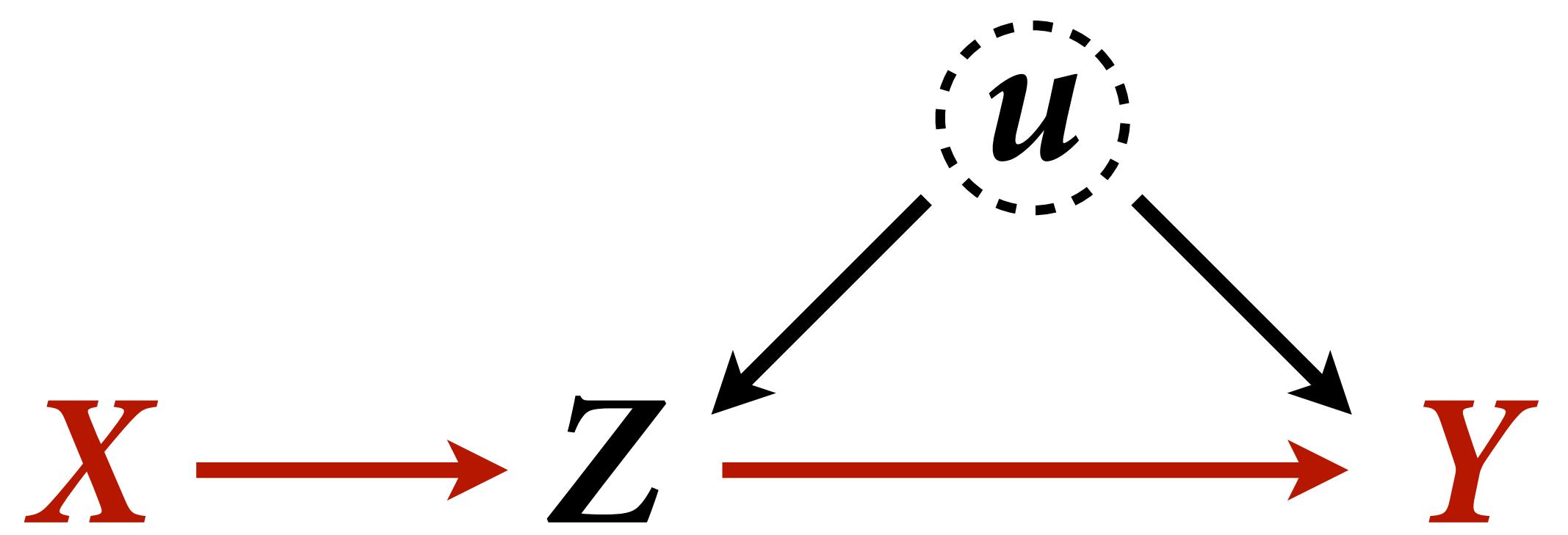
*Contextual  
confounds*



$$X \rightarrow Z \rightarrow Y$$

$$X \rightarrow Z \leftarrow u \rightarrow Y$$

No backdoor, no need  
to control for  $Z$



```

f <- function(n=100,bXZ=1,bZY=1) {
  X <- rnorm(n)
  u <- rnorm(n)
  Z <- rnorm(n, bXZ*X + u)
  Y <- rnorm(n, bZY*Z + u )
  bX <- coef( lm(Y ~ X) )['X']
  bXZ <- coef( lm(Y ~ X + Z) )['X']
  return( c(bX,bXZ) )
}

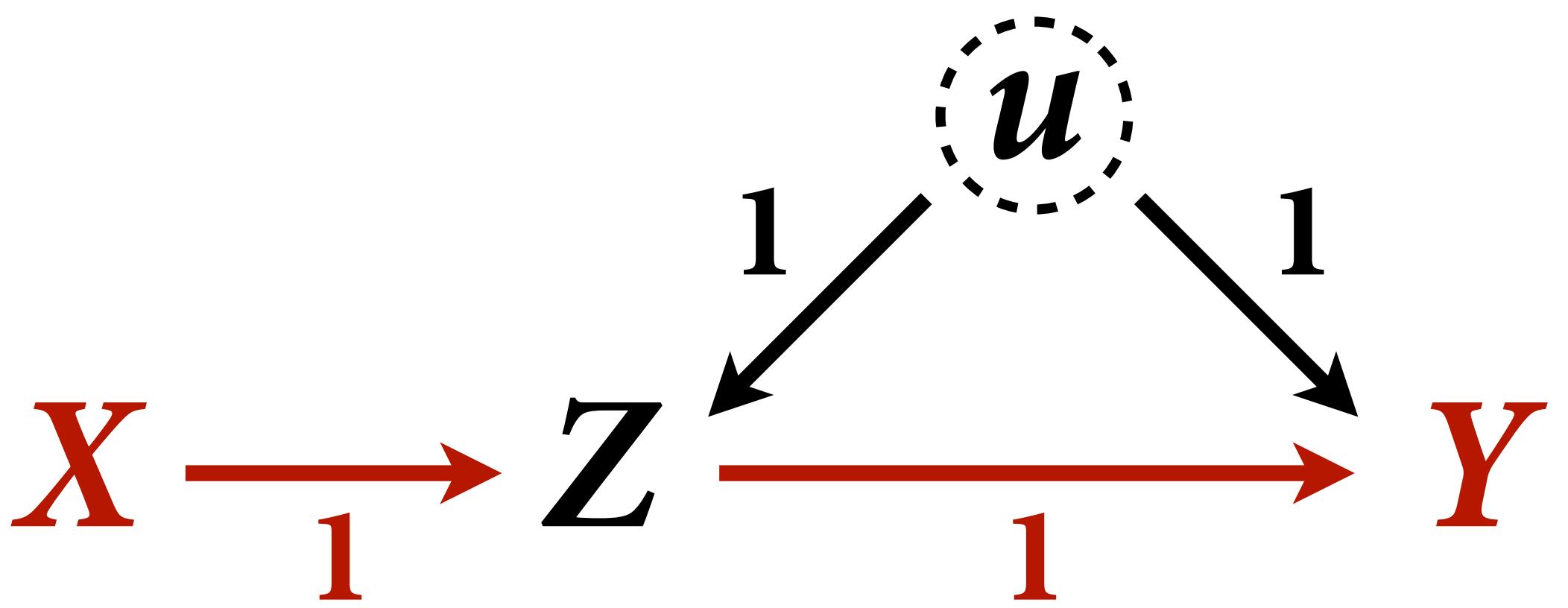
```

```

sim <- mcreplicate( 1e4 , f() , mc.cores=8 )

dens( sim[1,] , lwd=3 , xlab="posterior mean" )
dens( sim[2,] , lwd=3 , col=2 , add=TRUE )

```



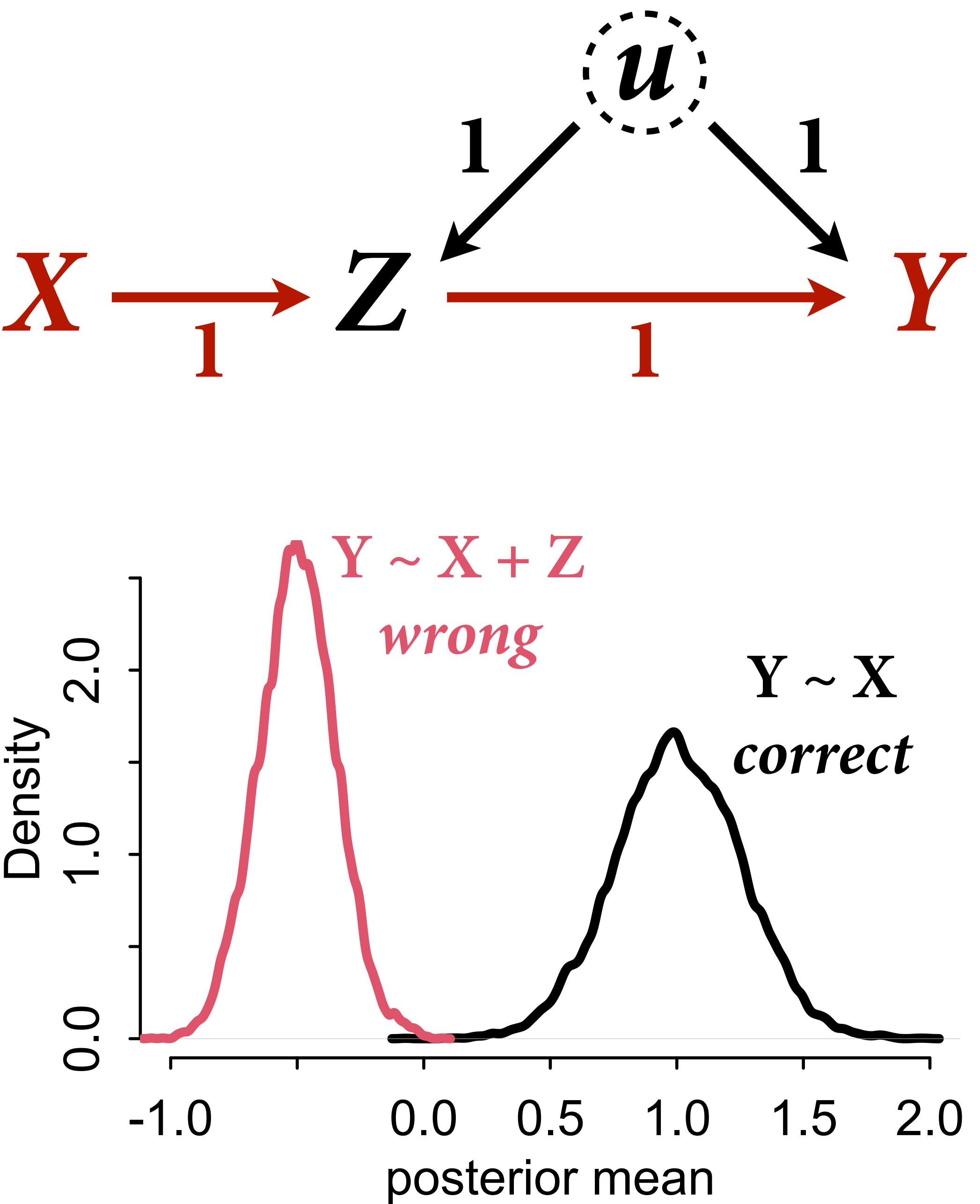
```

f <- function(n=100,bXZ=1,bZY=1) {
  X <- rnorm(n)
  u <- rnorm(n)
  Z <- rnorm(n, bXZ*X + u)
  Y <- rnorm(n, bZY*Z + u )
  bX <- coef( lm(Y ~ X) )['X']
  bXZ <- coef( lm(Y ~ X + Z) )['X']
  return( c(bX,bXZ) )
}

sim <- mcreplicate( 1e4 , f() , mc.cores=8 )

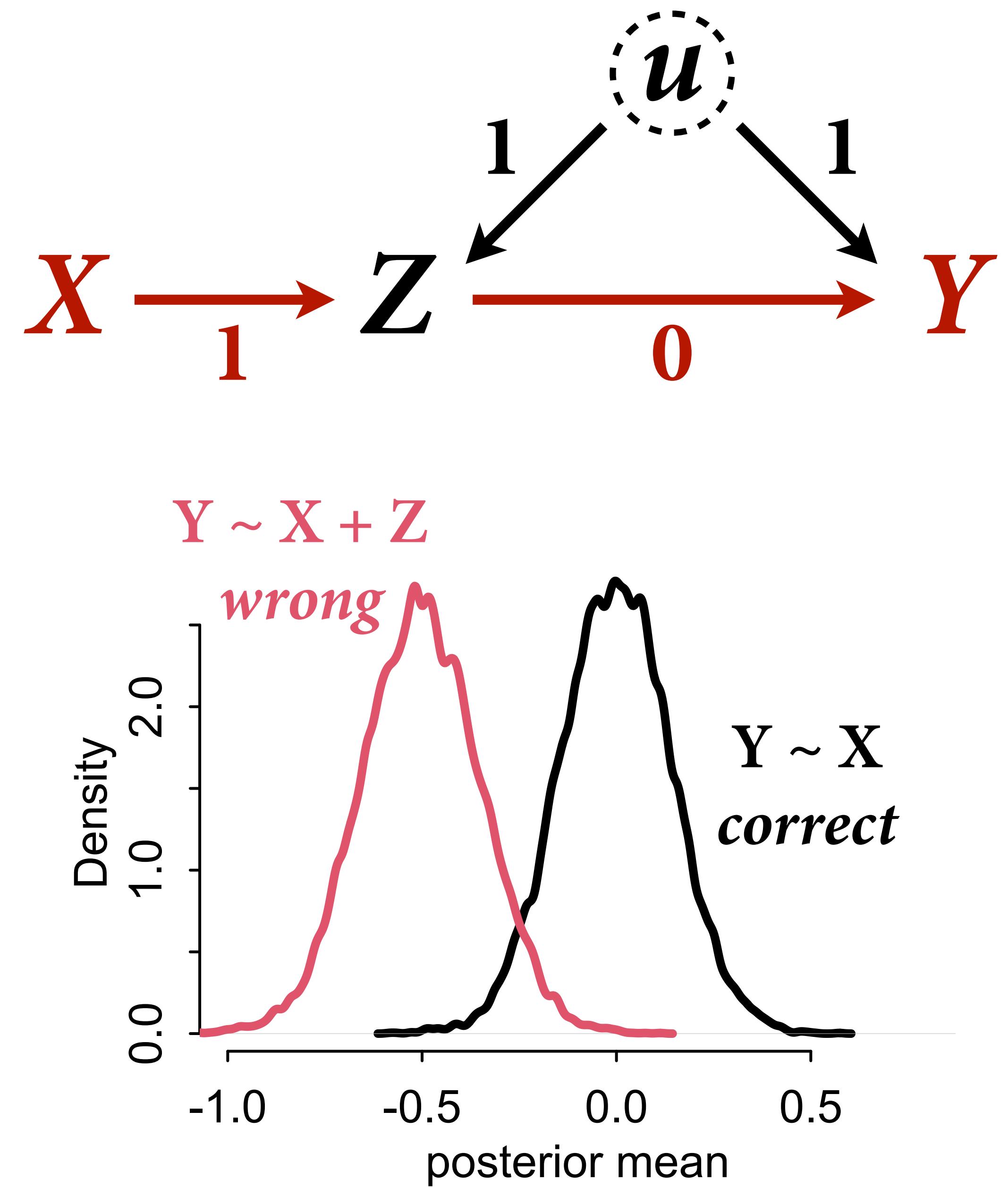
dens( sim[1,] , lwd=3 , xlab="posterior mean" )
dens( sim[2,] , lwd=3 , col=2 , add=TRUE )

```



# Change bZY to zero

```
f <- function(n=100,bXZ=1,bZY=1) {  
  X <- rnorm(n)  
  u <- rnorm(n)  
  Z <- rnorm(n, bXZ*X + u)  
  Y <- rnorm(n, bZY*Z + u )  
  bX <- coef( lm(Y ~ X) )['X']  
  bXZ <- coef( lm(Y ~ X + Z) )['X']  
  return( c(bX,bXZ) )  
}  
  
sim <- mcreplicate( 1e4 , f(bZY=0) , mc.cores=8 )  
  
dens( sim[1,] , lwd=3 , xlab="posterior mean" )  
dens( sim[2,] , lwd=3 , col=2 , add=TRUE )
```



$$X \rightarrow Z \rightarrow Y$$

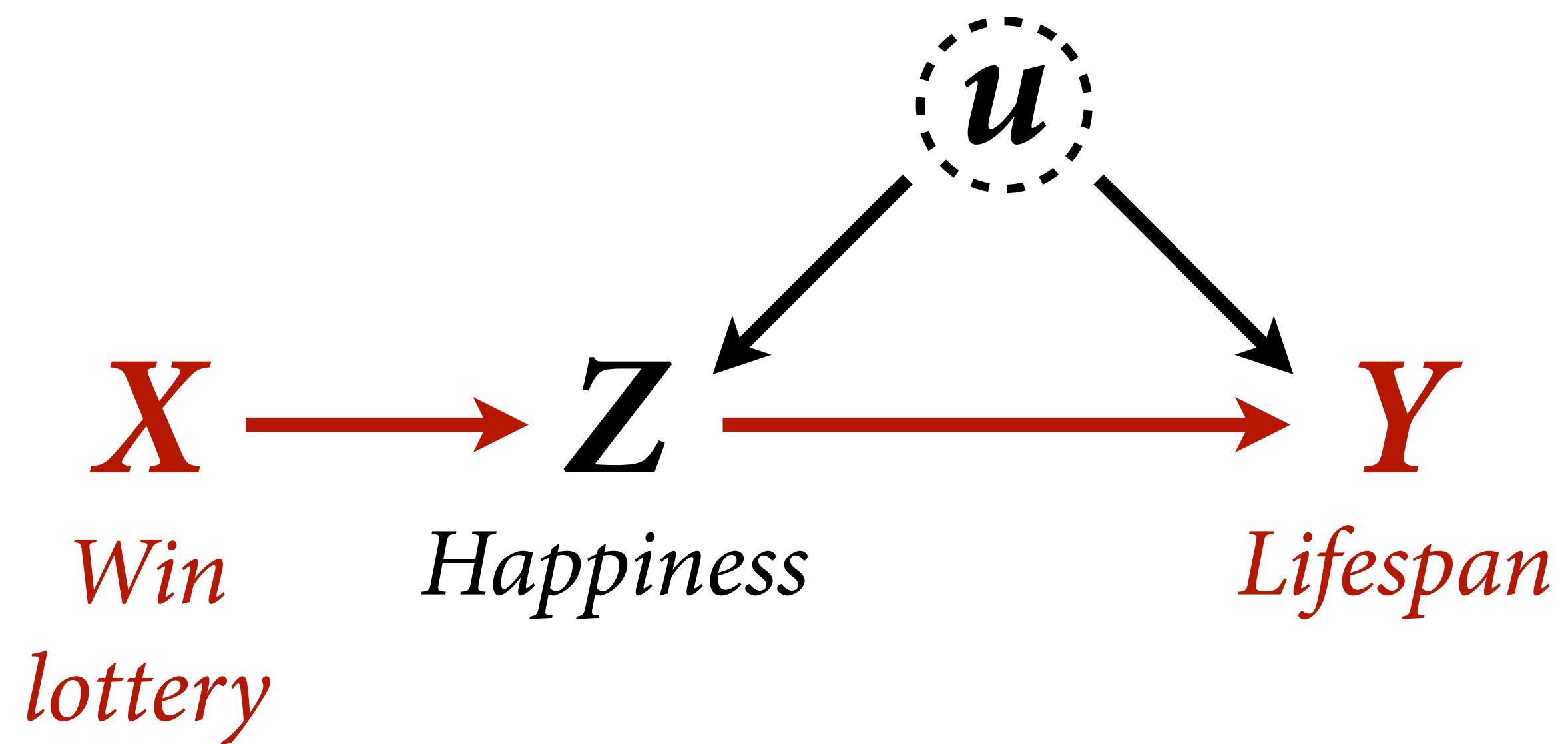
$$X \rightarrow Z \leftarrow u \rightarrow Y$$

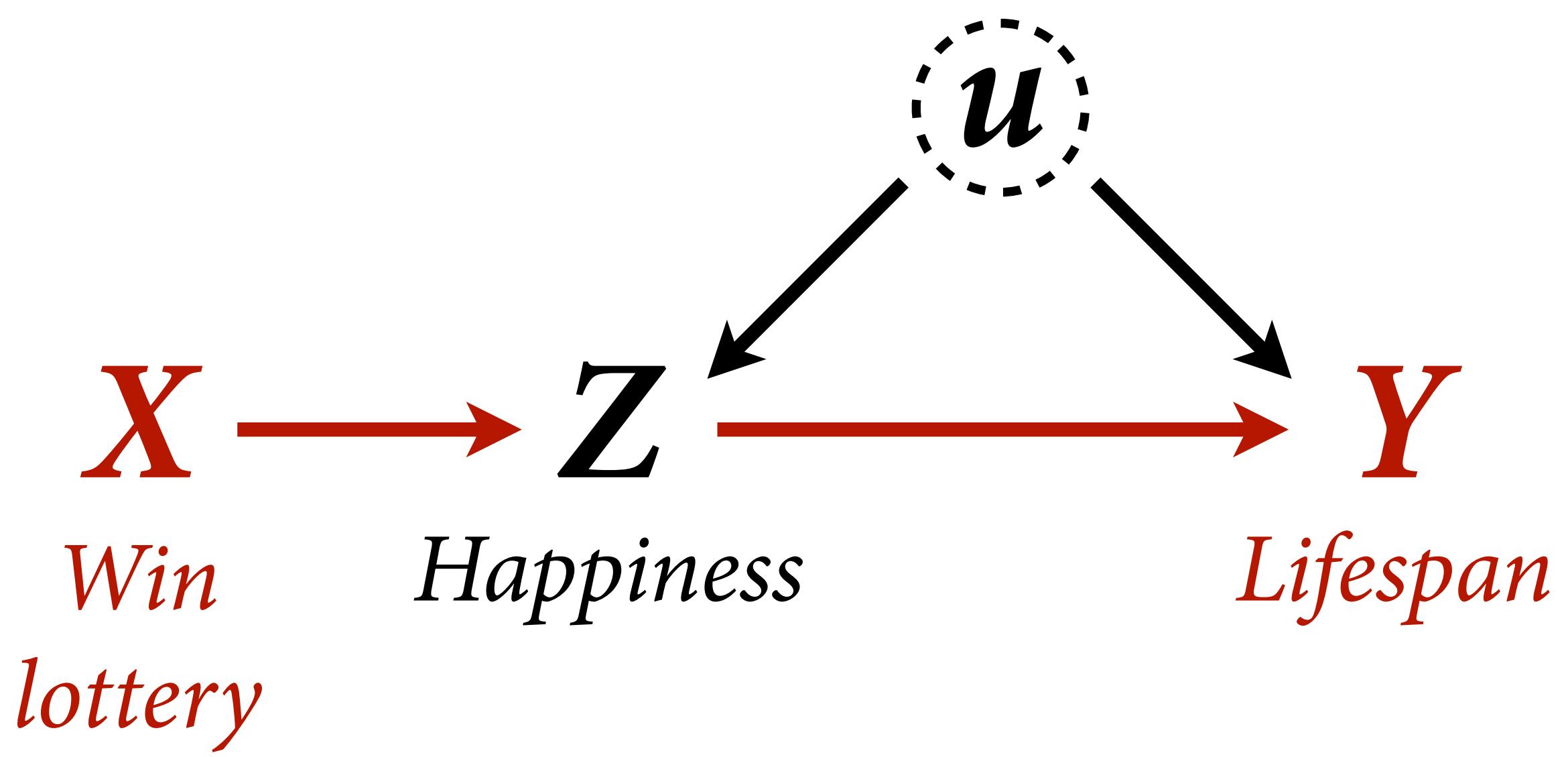
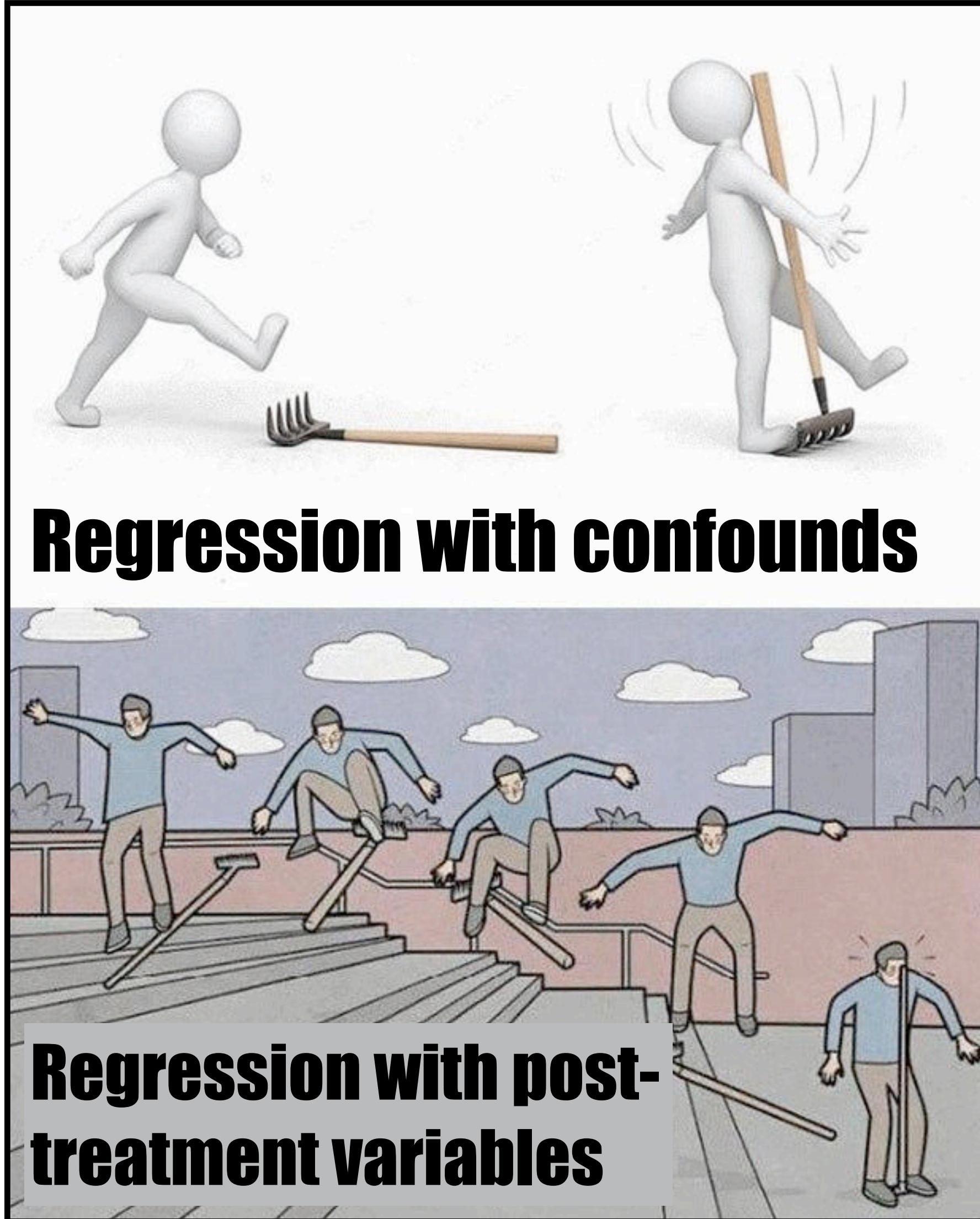
Controlling for  $Z$  biases  
treatment estimate  $X$

Controlling for  $Z$  opens biasing  
path through  $u$

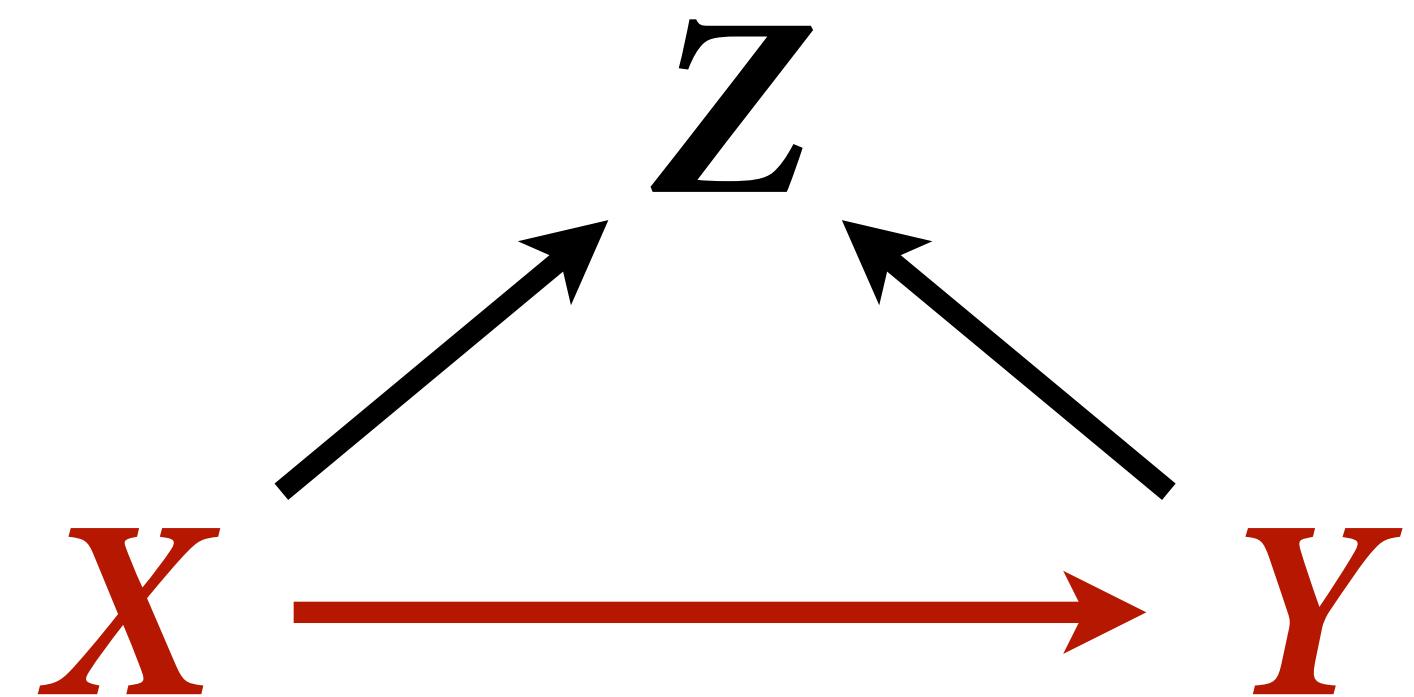
Can estimate effect of  $X$ ; Cannot  
estimate mediation effect  $Z$

No backdoor, no need  
to control for  $Z$

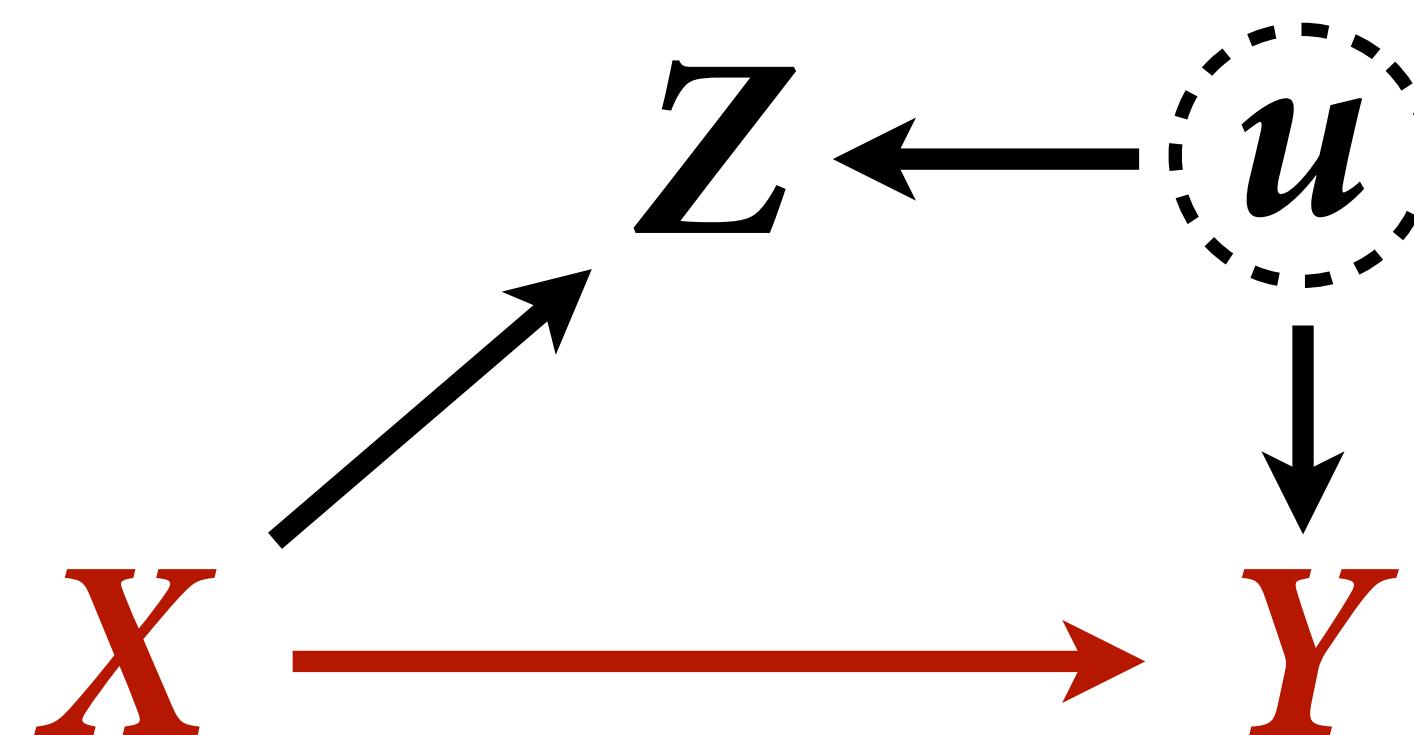


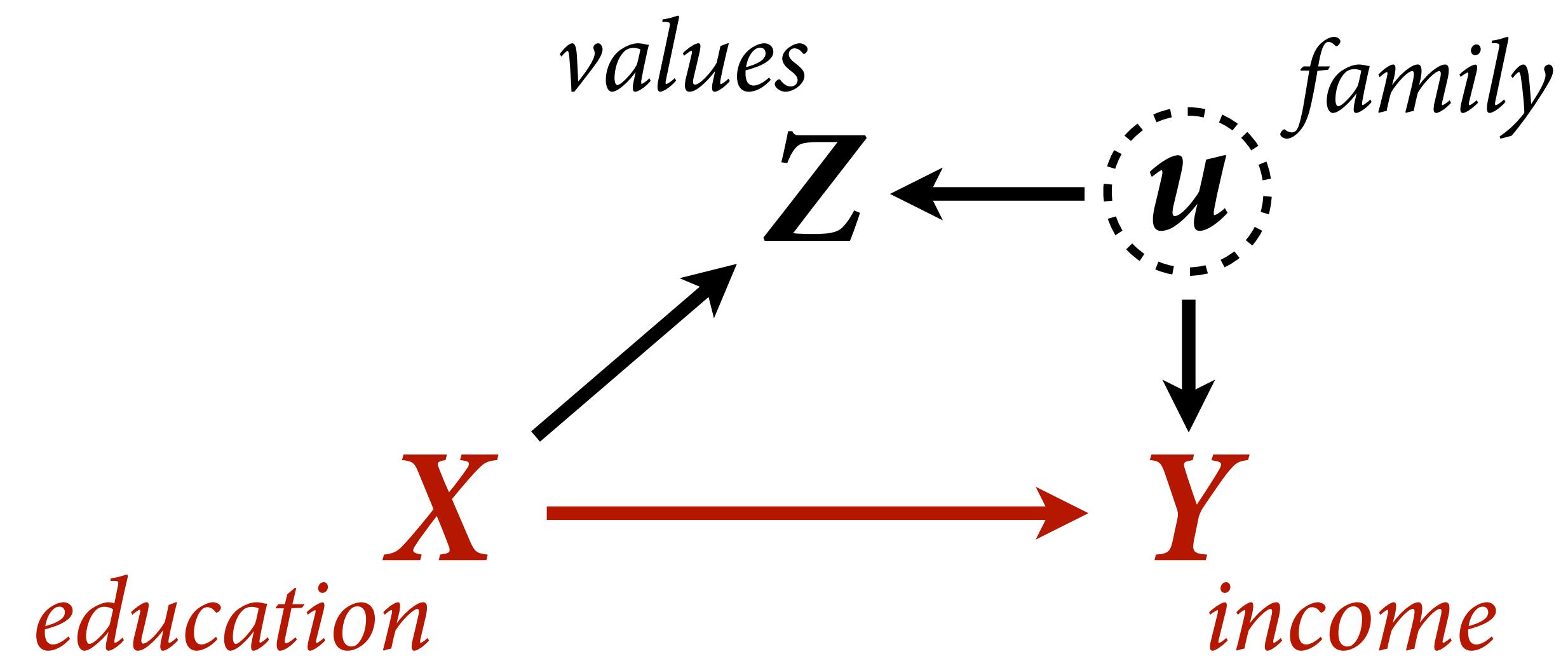


Do not touch the collider!

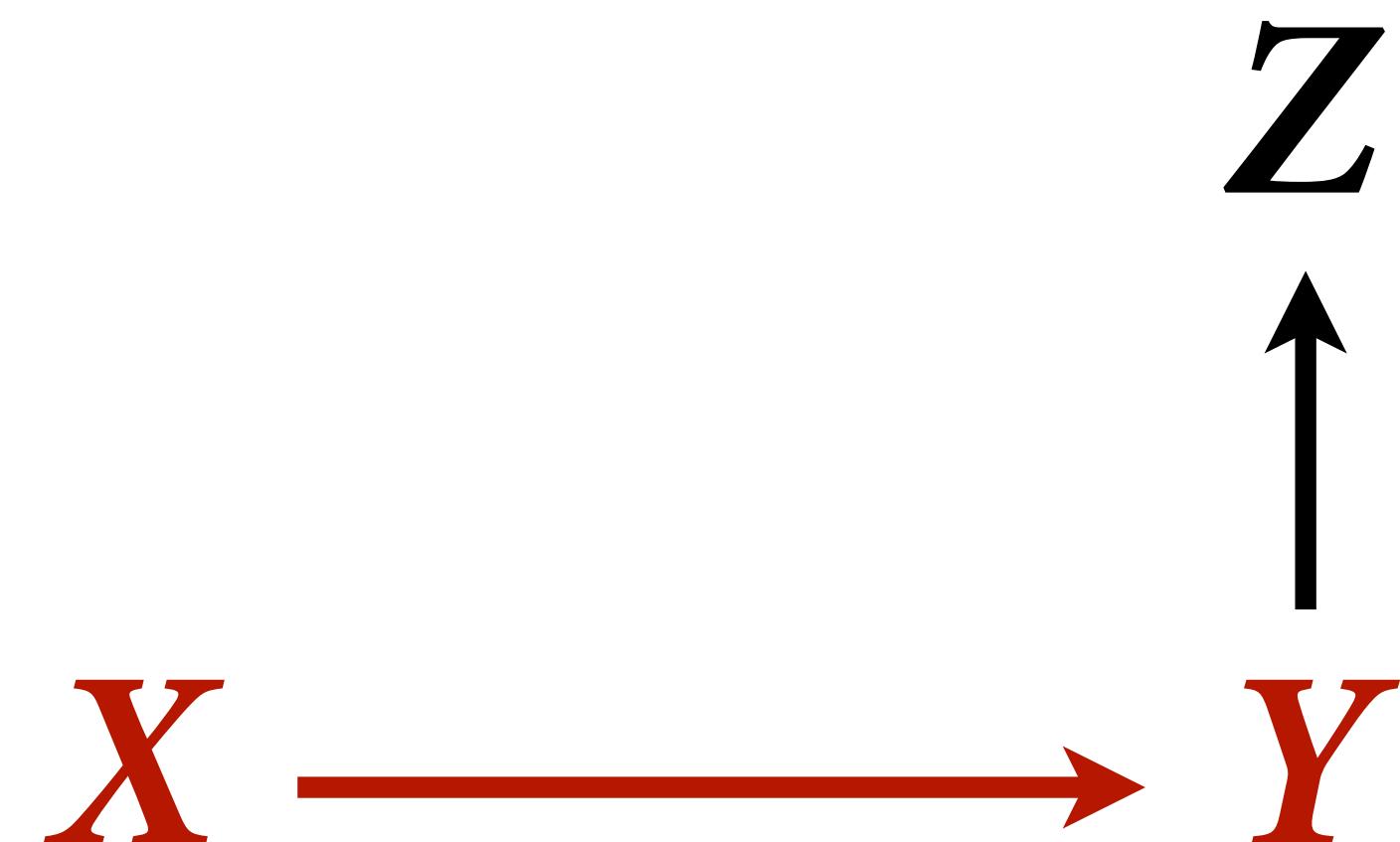


Colliders not always so obvious

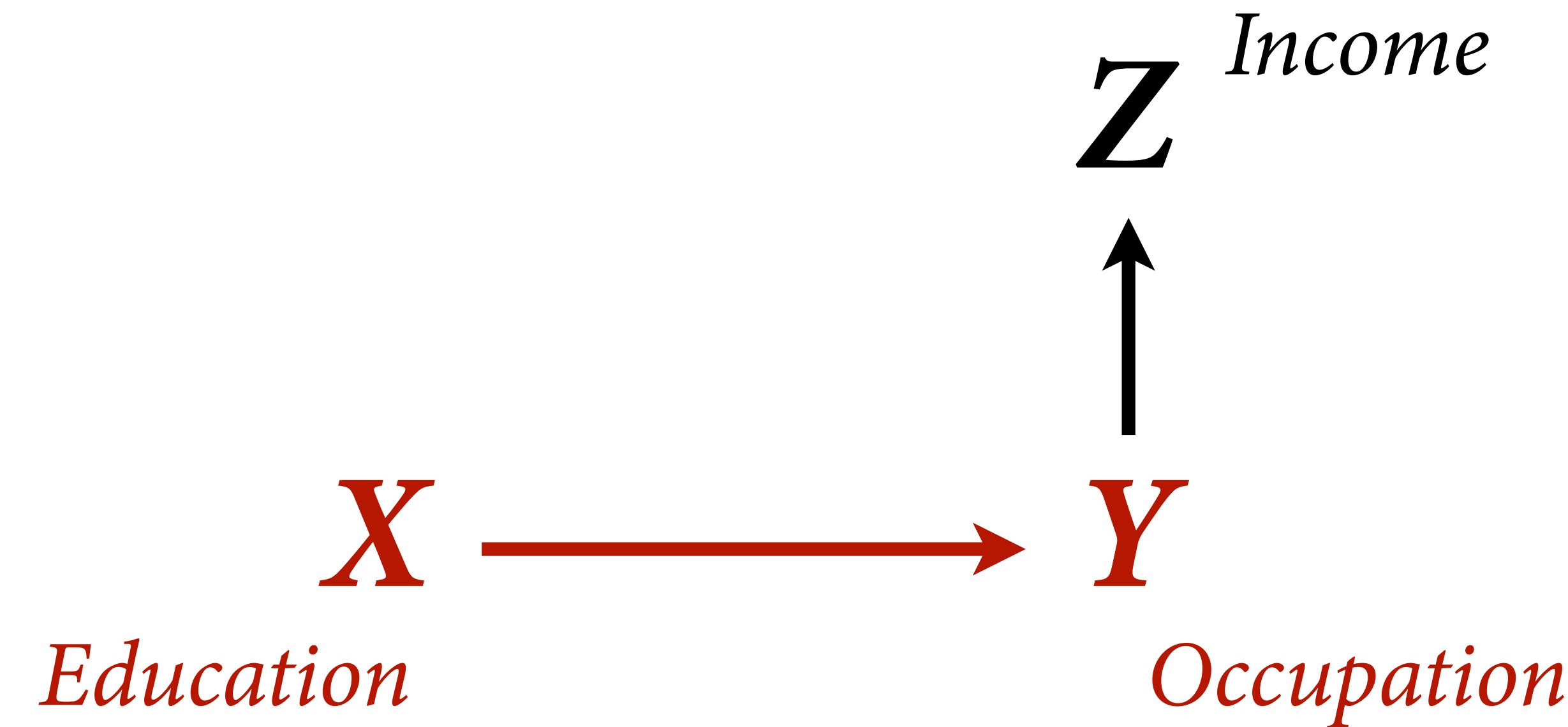




# Case-control bias (selection on outcome)

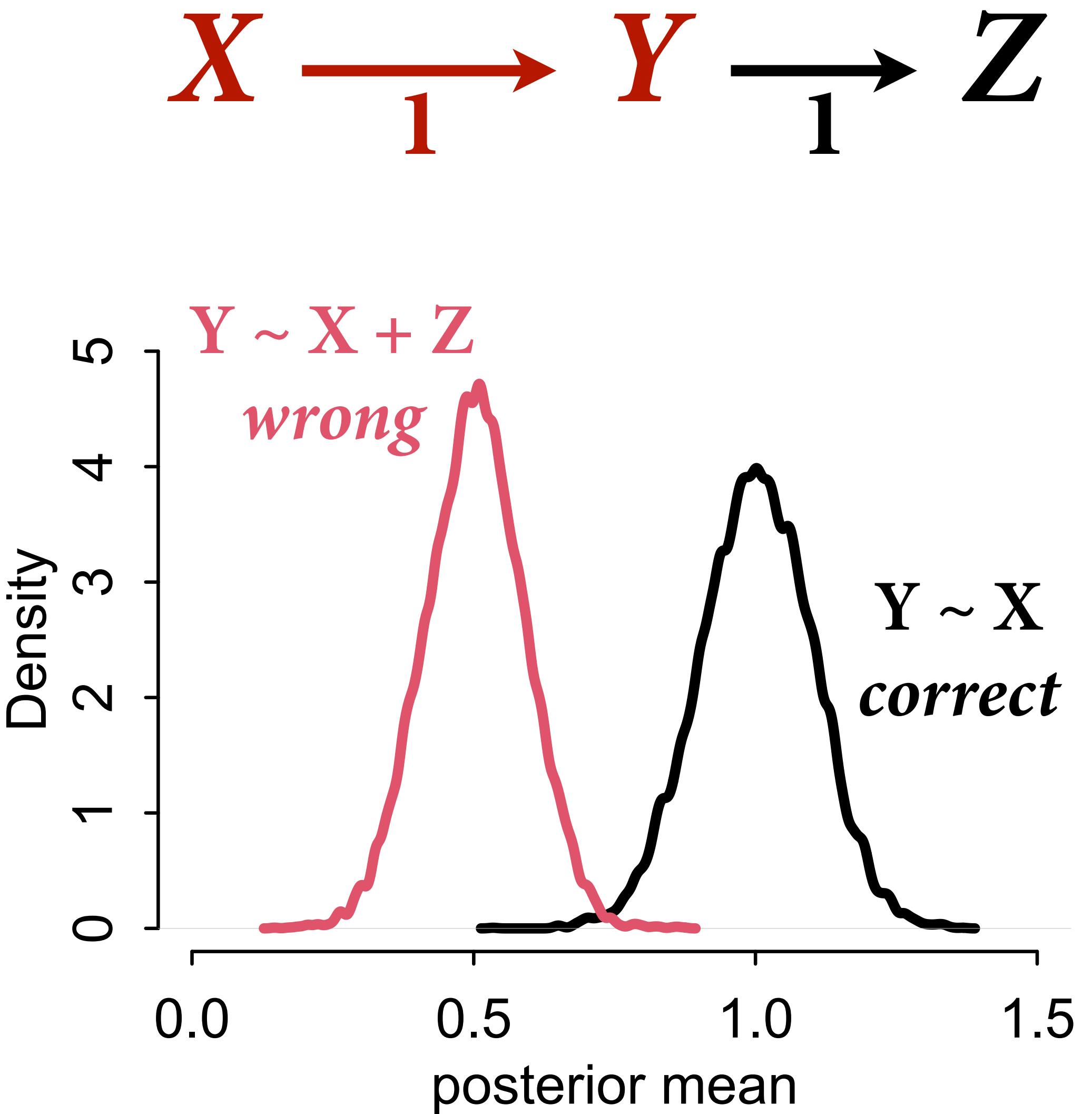


# Case-control bias (selection on outcome)



# Case-control bias (selection on outcome)

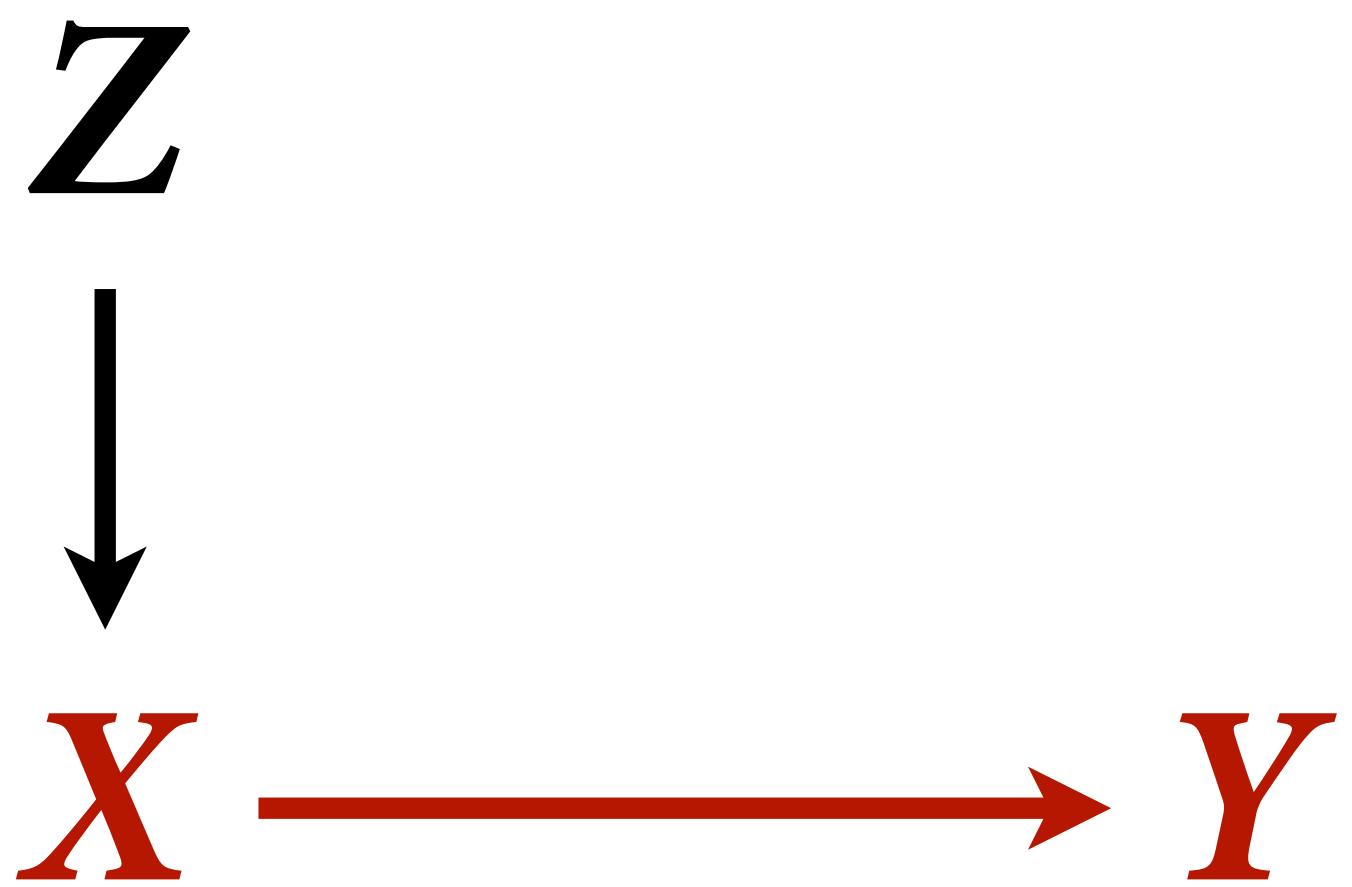
```
f <- function(n=100,bXY=1,bYZ=1) {  
  X <- rnorm(n)  
  Y <- rnorm(n, bXY*X )  
  Z <- rnorm(n, bYZ*Y )  
  bX <- coef( lm(Y ~ X) )['X']  
  bXZ <- coef( lm(Y ~ X + Z) )['X']  
  return( c(bX,bXZ) )  
}  
  
sim <- mcreplicate( 1e4 , f() , mc.cores=8 )  
  
dens( sim[1,] , lwd=3 , xlab="posterior mean" )  
dens( sim[2,] , lwd=3 , col=2 , add=TRUE )
```



“Precision parasite”

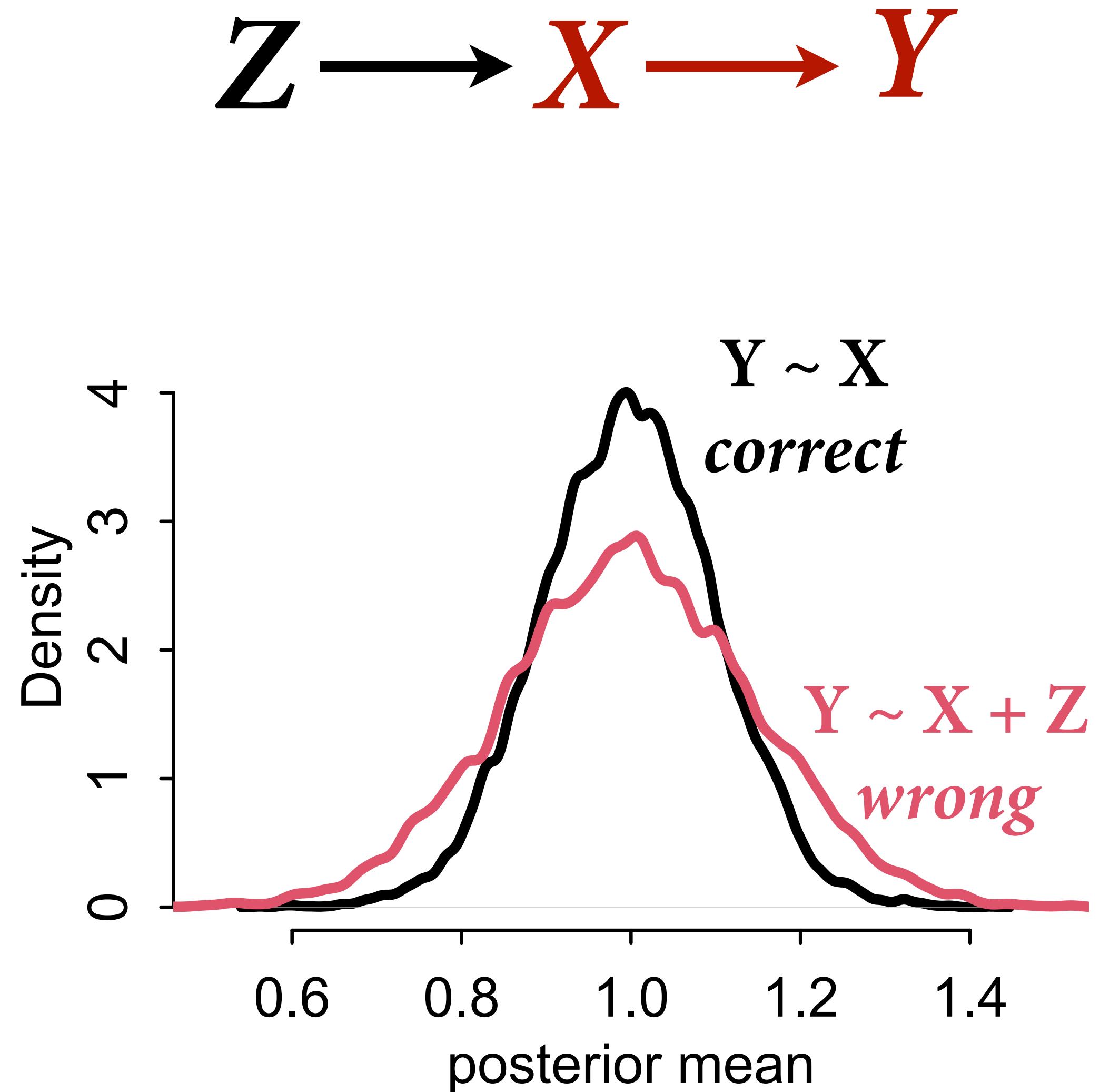
No backdoors

But still not good to  
condition on  $Z$



# “Precision parasite”

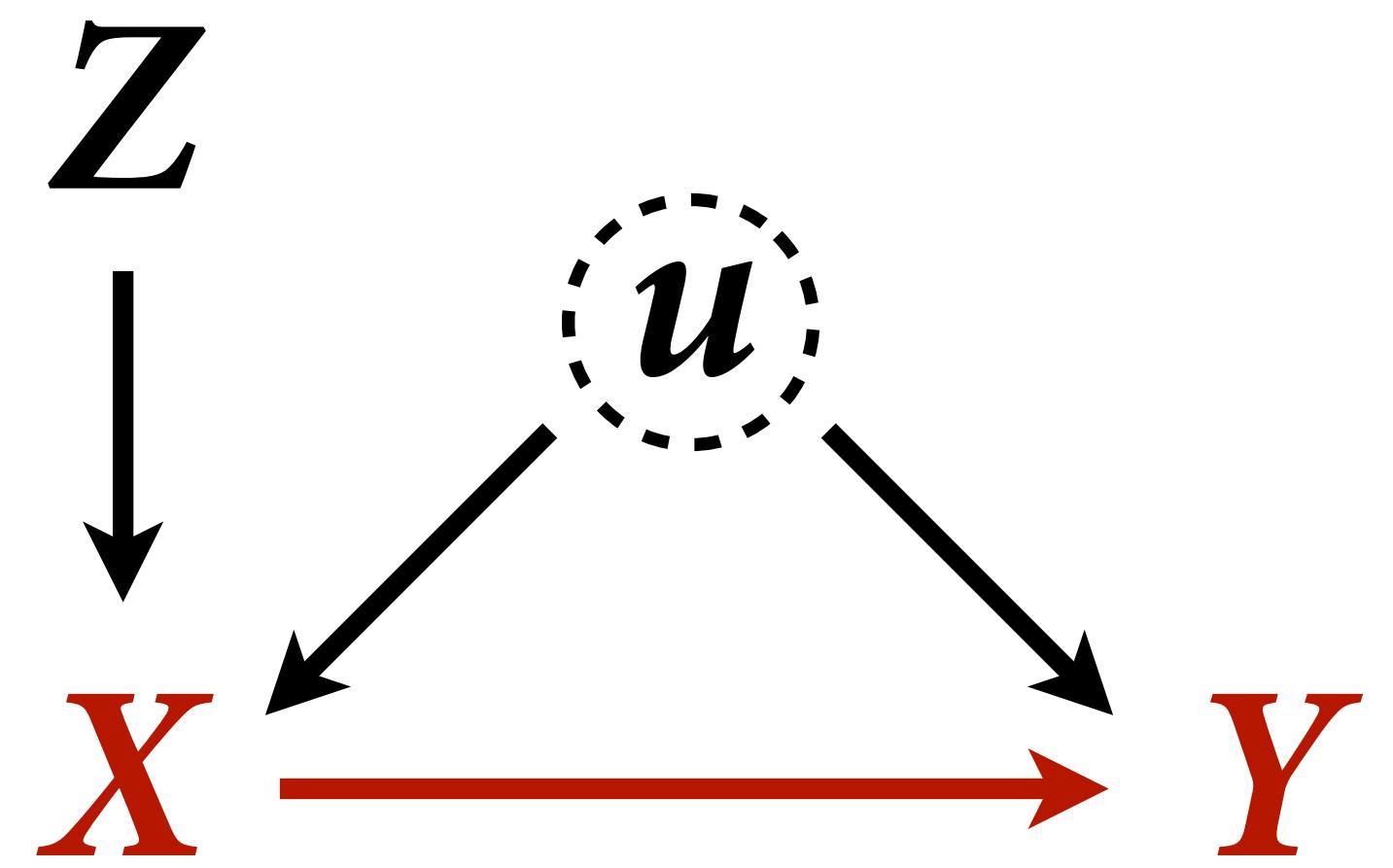
```
f <- function(n=100,bZX=1,bXY=1) {  
  Z <- rnorm(n)  
  X <- rnorm(n, bZX*Z )  
  Y <- rnorm(n, bXY*X )  
  bX <- coef( lm(Y ~ X) )['X']  
  bXZ <- coef( lm(Y ~ X + Z) )['X']  
  return( c(bX,bXZ) )  
}  
  
sim <- mcreplicate( 1e4 , f(n=50) , mc.cores=8 )  
  
dens( sim[1,] , lwd=3 , xlab="posterior mean" )  
dens( sim[2,] , lwd=3 , col=2 , add=TRUE )
```



“Bias amplification”

$X$  and  $Y$  confounded by  $u$

Something **truly awful** happens  
when we add  $Z$

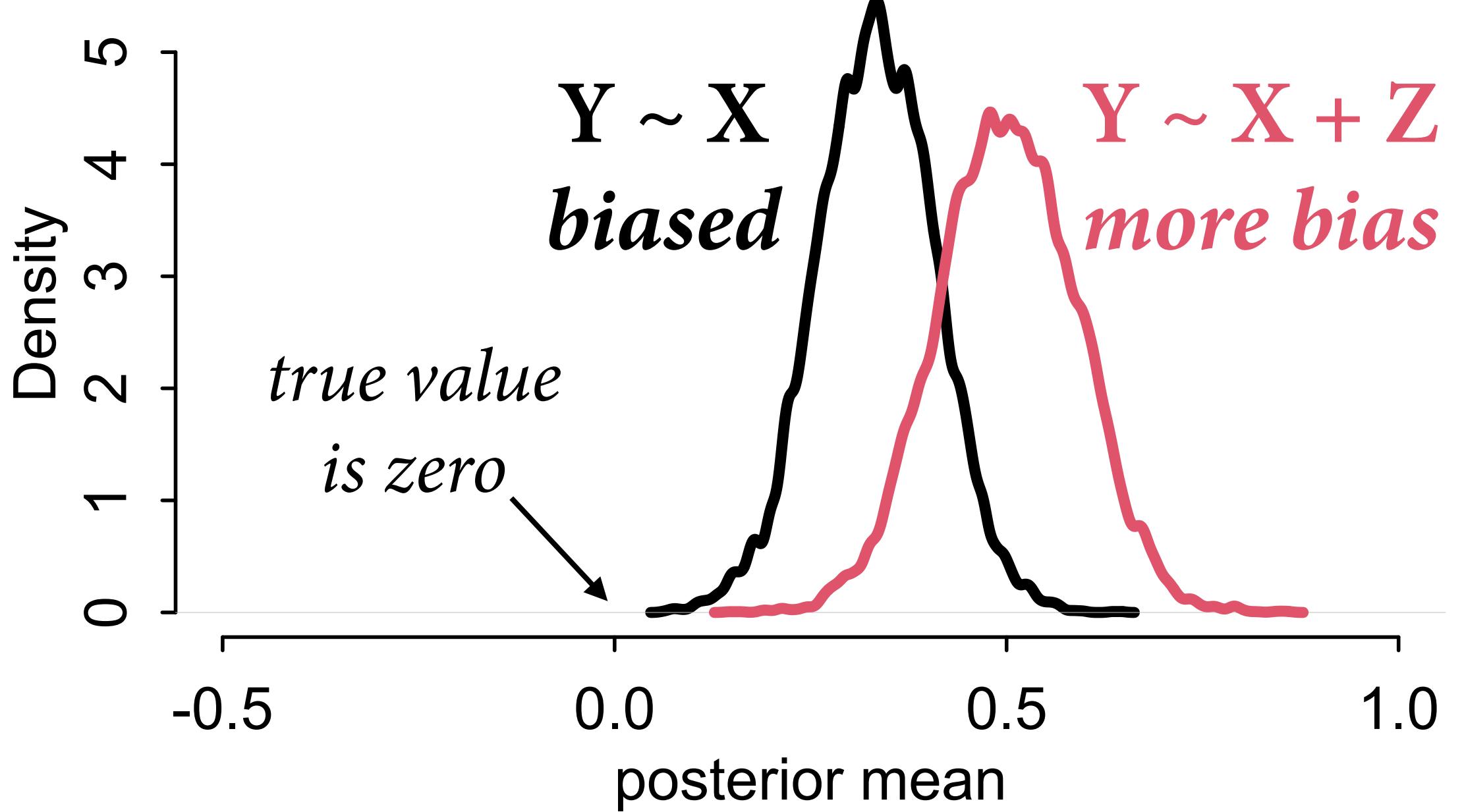
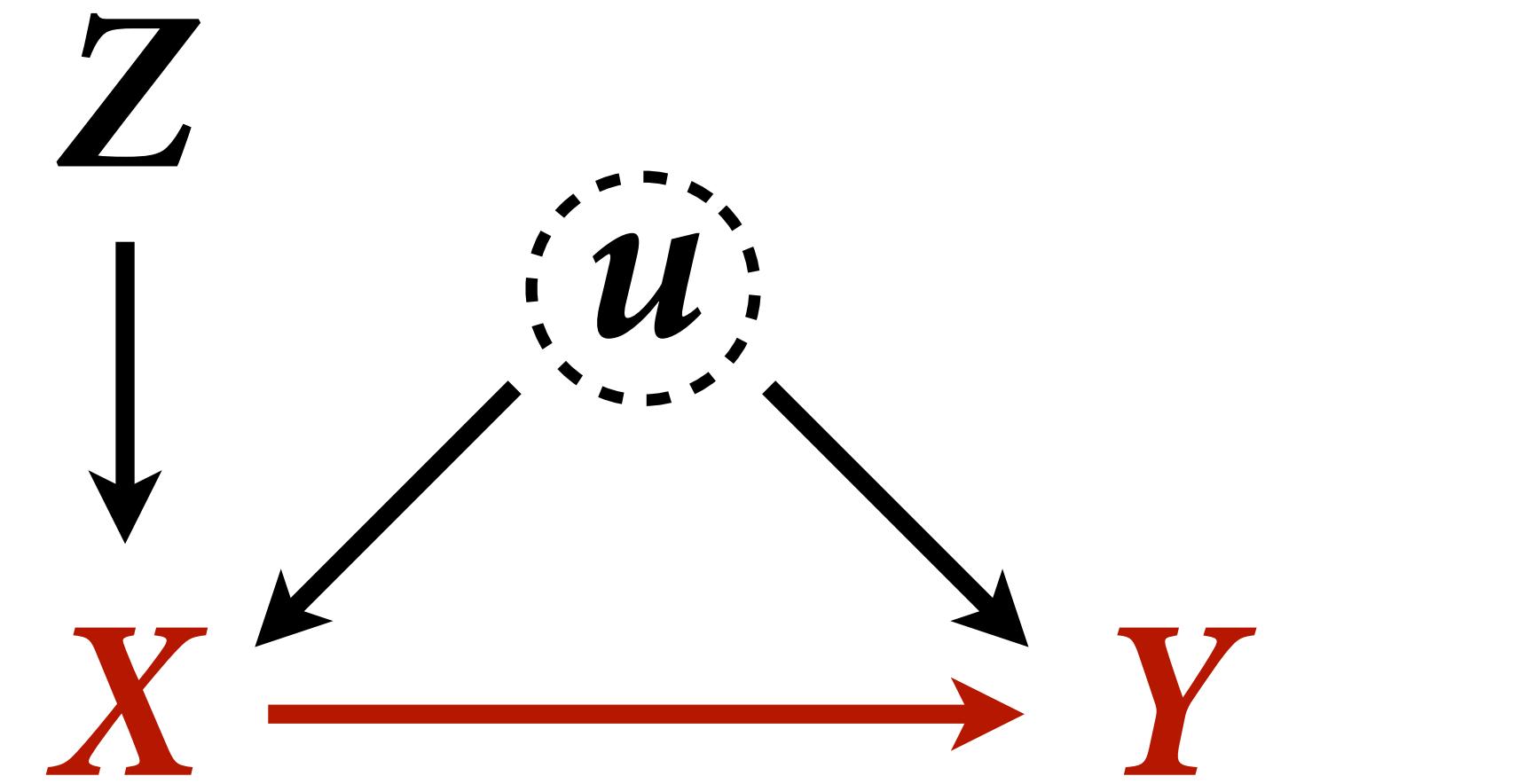


```

f <- function(n=100,bZX=1,bXY=1) {
  Z <- rnorm(n)
  u <- rnorm(n)
  X <- rnorm(n, bZX*Z + u )
  Y <- rnorm(n, bXY*X + u )
  bX <- coef( lm(Y ~ X) )['X']
  bXZ <- coef( lm(Y ~ X + Z) )['X']
  return( c(bX,bXZ) )
}
sim <- mcreplicate( 1e4 , f(bXY=0) , mc.cores=8 )

dens( sim[1,] , lwd=3 , xlab="posterior mean" )
dens( sim[2,] , lwd=3 , col=2 , add=TRUE )

```

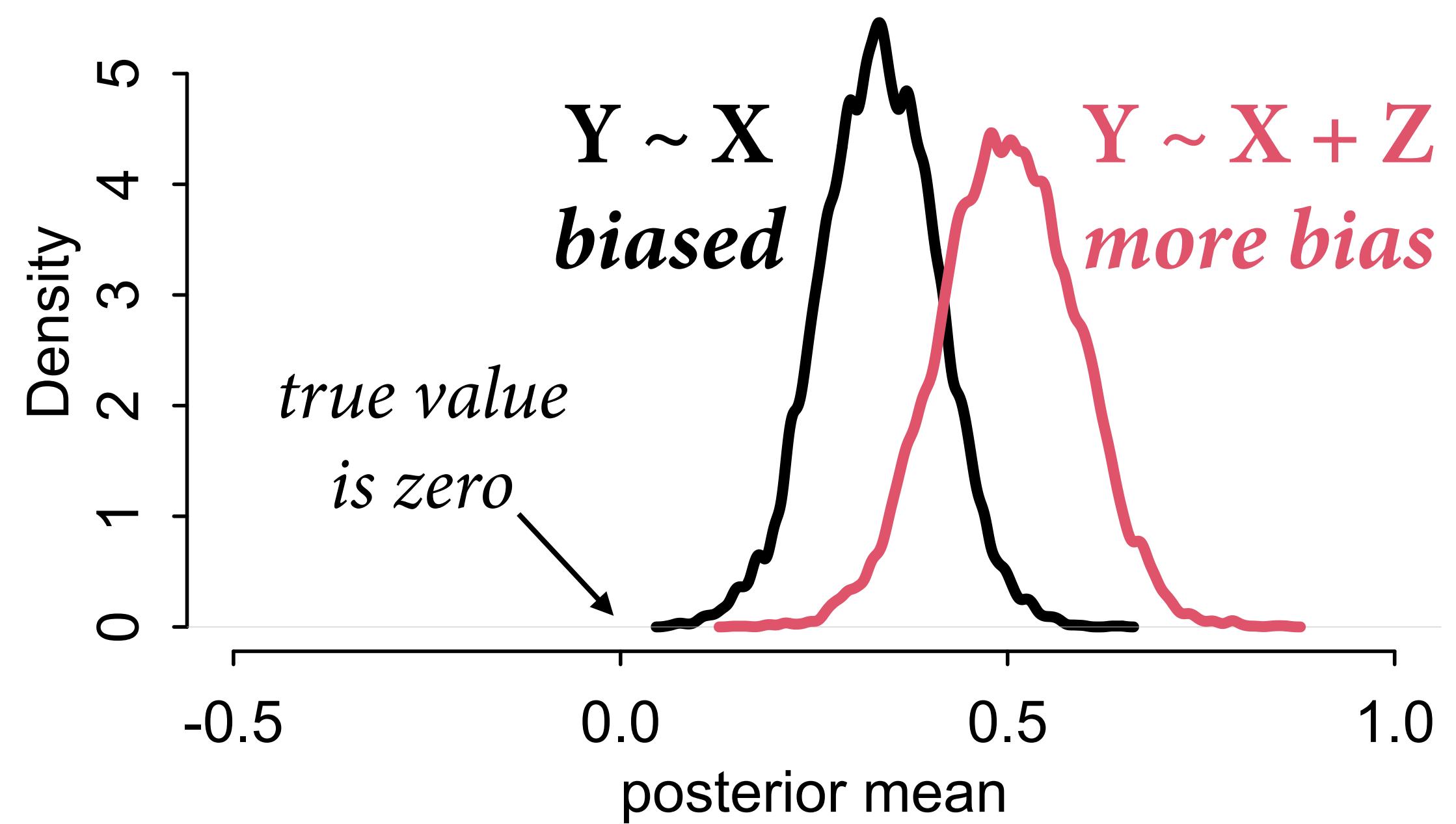
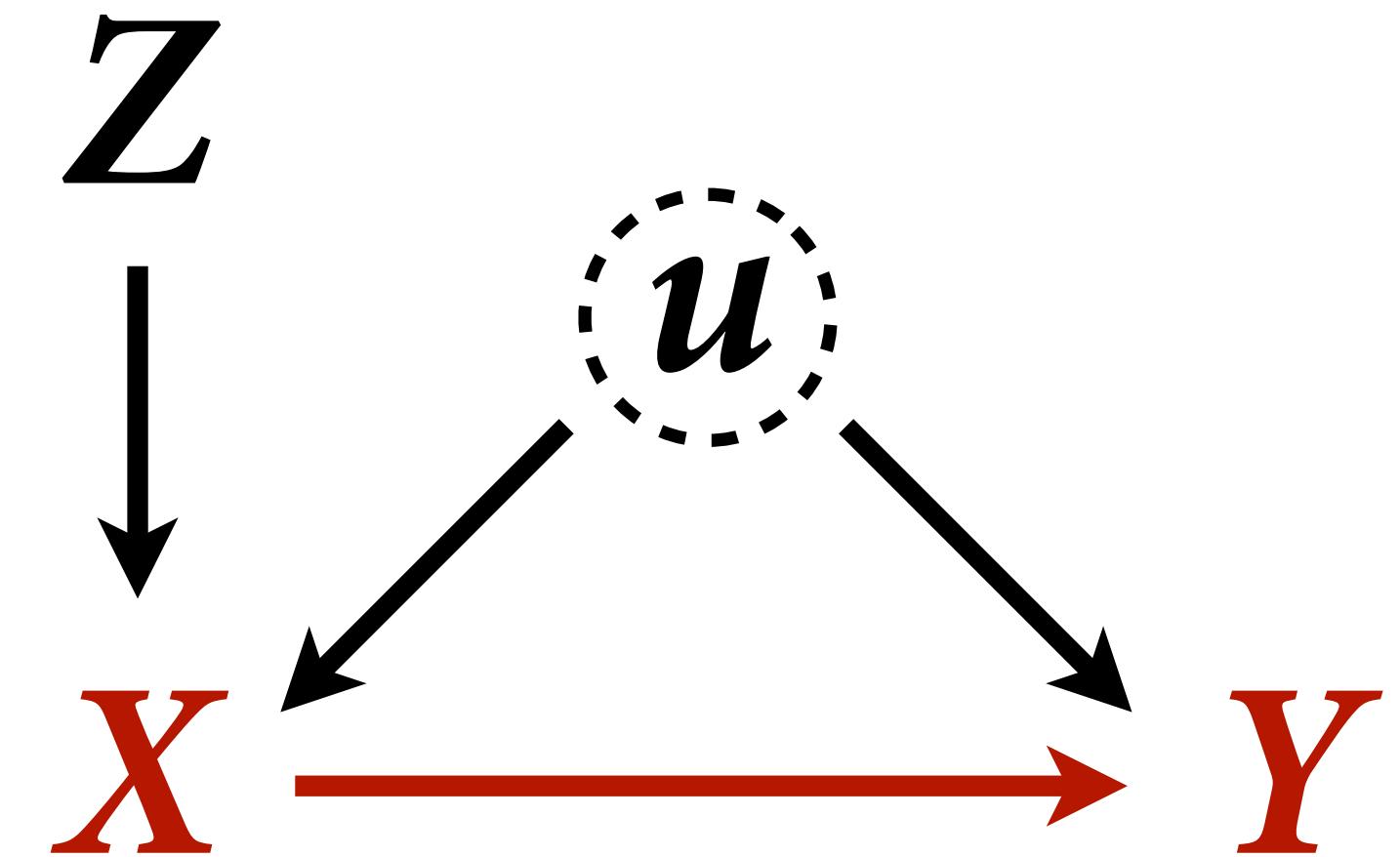


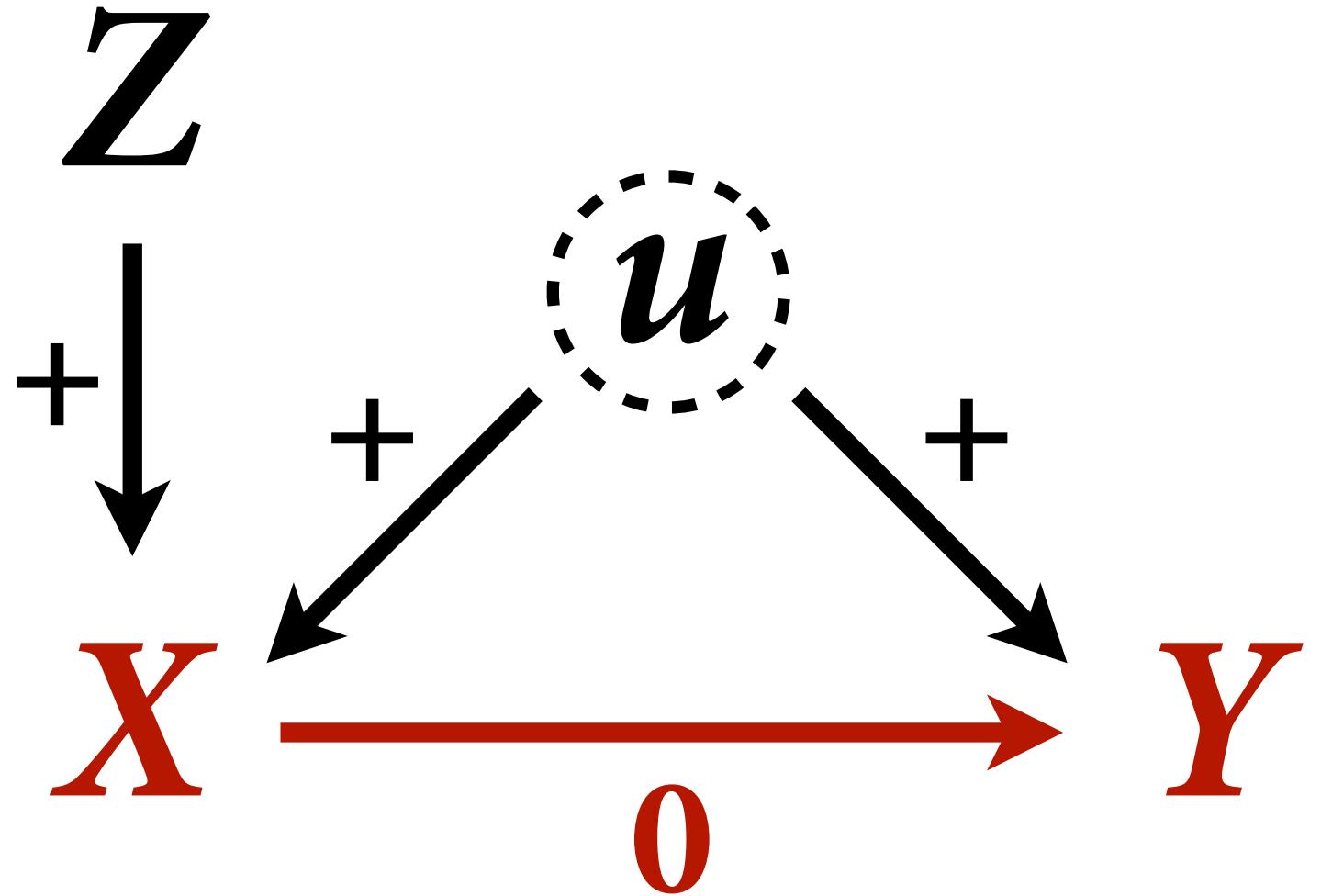
*WHY?*

Covariation  $X$  &  $Y$  requires variation in their causes

Within each level of  $Z$ , less variation in  $X$

Confound  $u$  relatively more important within each  $Z$

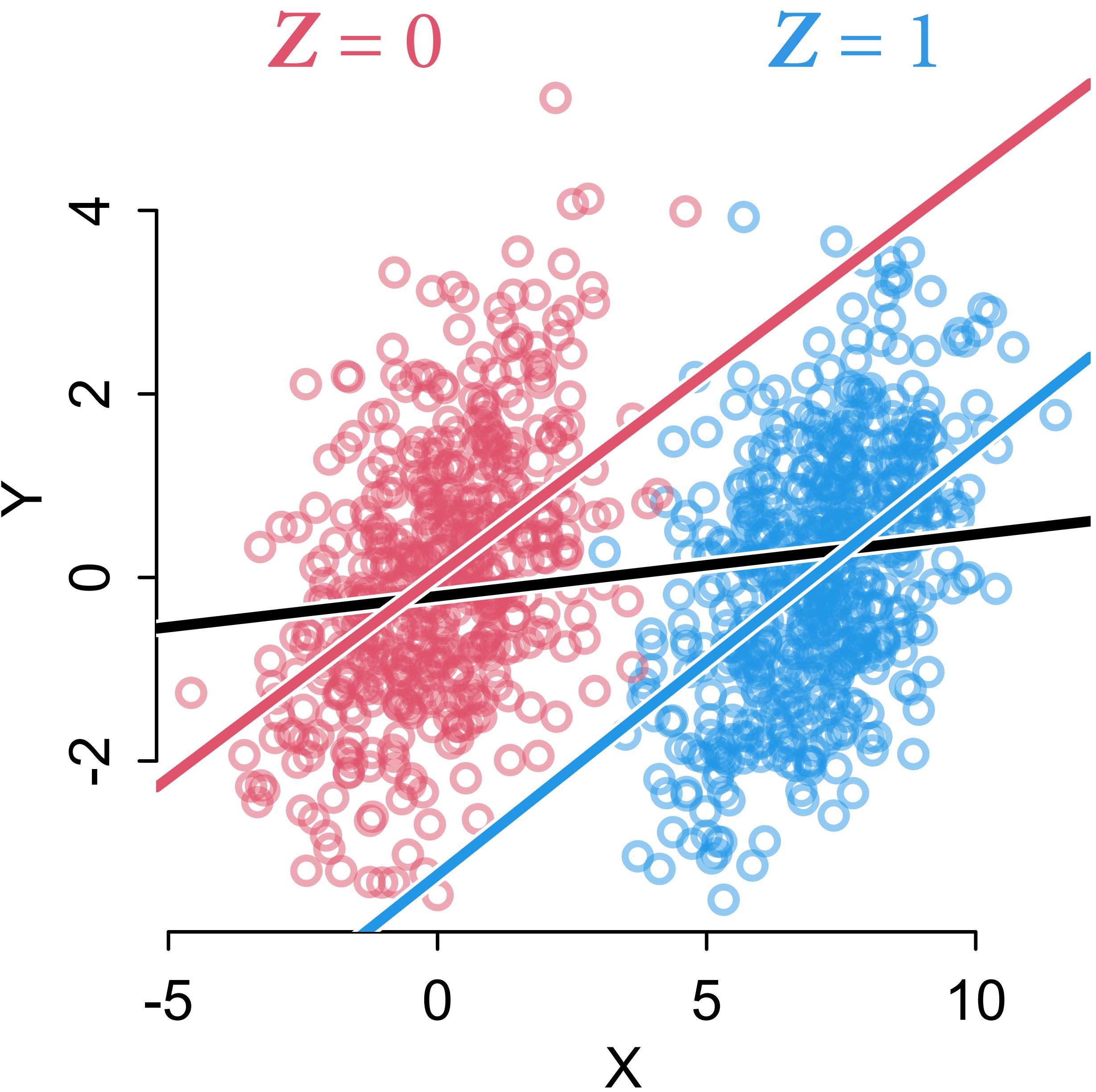


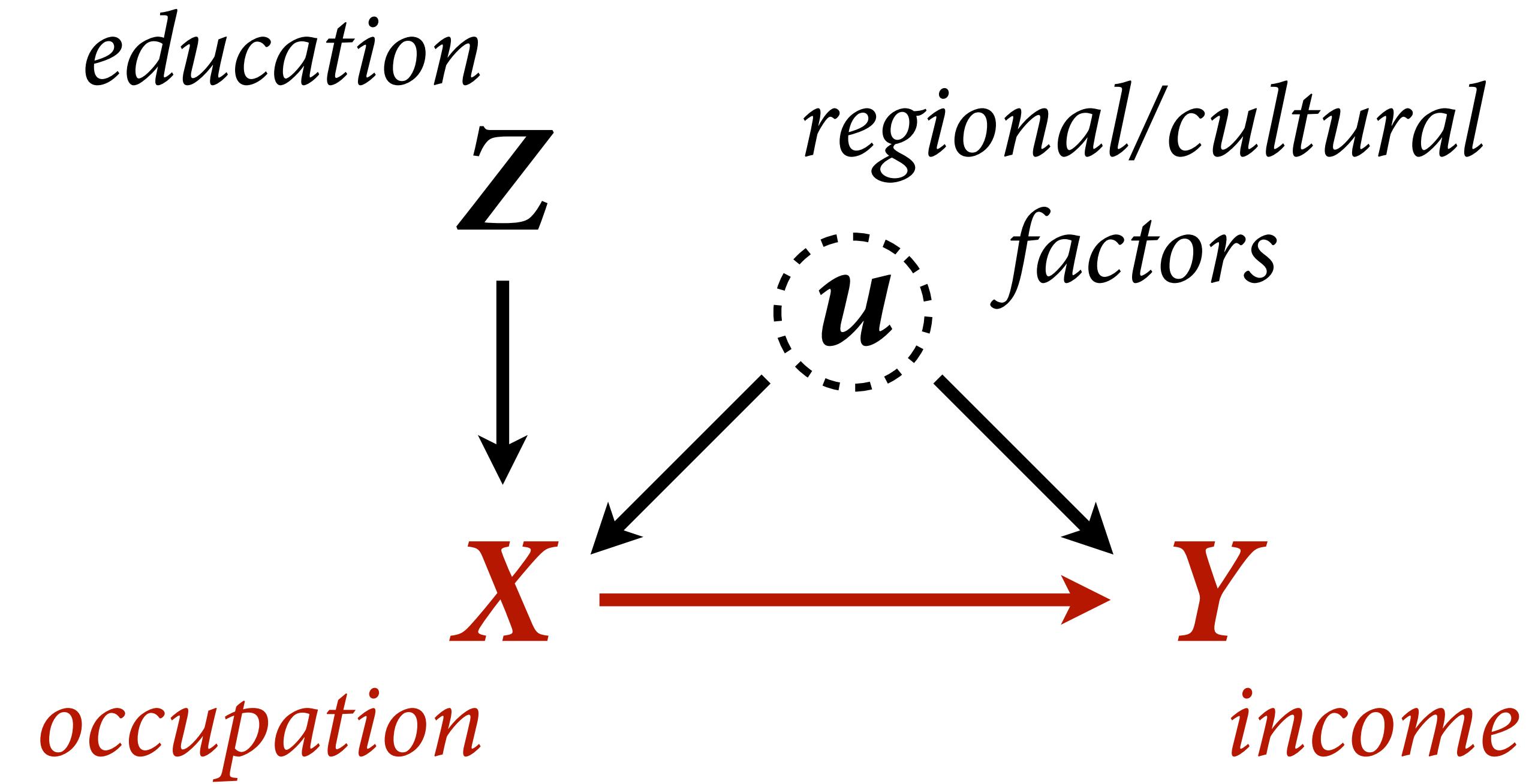


```

n <- 1000
Z <- rbern(n)
u <- rnorm(n)
X <- rnorm(n, 7*Z + u )
Y <- rnorm(n, 0*X + u )

```





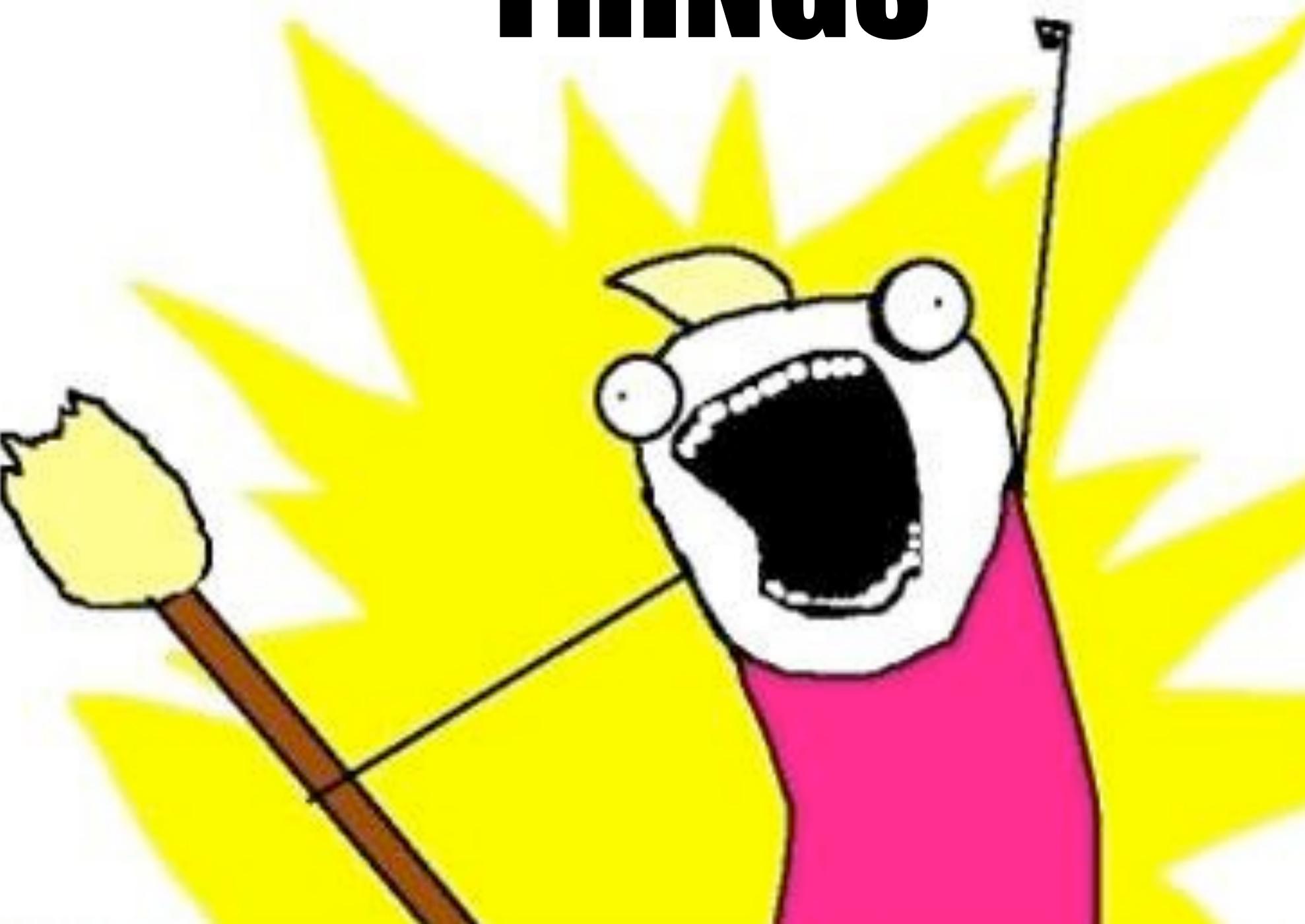
# Good & Bad Controls

**“Control” variable:** Variable introduced to an analysis so that a causal estimate is possible

Heuristics fail — adding control variables can be worse than omitting

Make assumptions explicit

**MODEL  
ALL THE  
THINGS**



# Course Schedule

Week 1	Bayesian inference	Chapters 1, 2, 3
Week 2	Linear models & Causal Inference	Chapter 4
Week 3	Causes, Confounds & Colliders	Chapters 5 & 6
Week 4	Overfitting / MCMC	Chapters 7, 8, 9
Week 5	Generalized Linear Models	Chapters 10, 11
Week 6	Integers & Other Monsters	Chapters 11 & 12
Week 7	Multilevel models I	Chapter 13
Week 8	Multilevel models II	Chapter 14
Week 9	Measurement & Missingness	Chapter 15
Week 10	Generalized Linear Madness	Chapter 16

[https://github.com/rmcelreath/stat\\_rethinking\\_2023](https://github.com/rmcelreath/stat_rethinking_2023)



# BONUS

TABLE 2—ESTIMATED PROBIT MODELS  
 FOR THE USE OF A SCREEN

	Preliminaries blind	Finals	
	(1)	(2)	blind
(Proportion female) <sub>t-1</sub>	2.744 (3.265) [0.006]	3.120 (3.271) [0.004]	0.490 (1.163) [0.011]
(Proportion of orchestra personnel with <6 years tenure) <sub>t-1</sub>	-26.46 (7.314) [-0.058]	-28.13 (8.459) [-0.039]	-9.467 (2.787) [-0.207]
“Big Five” orchestra		0.367 (0.452) [0.001]	
pseudo $R^2$	0.178	0.193	0.050
Number of observations	294	294	434

# Table 2 Fallacy

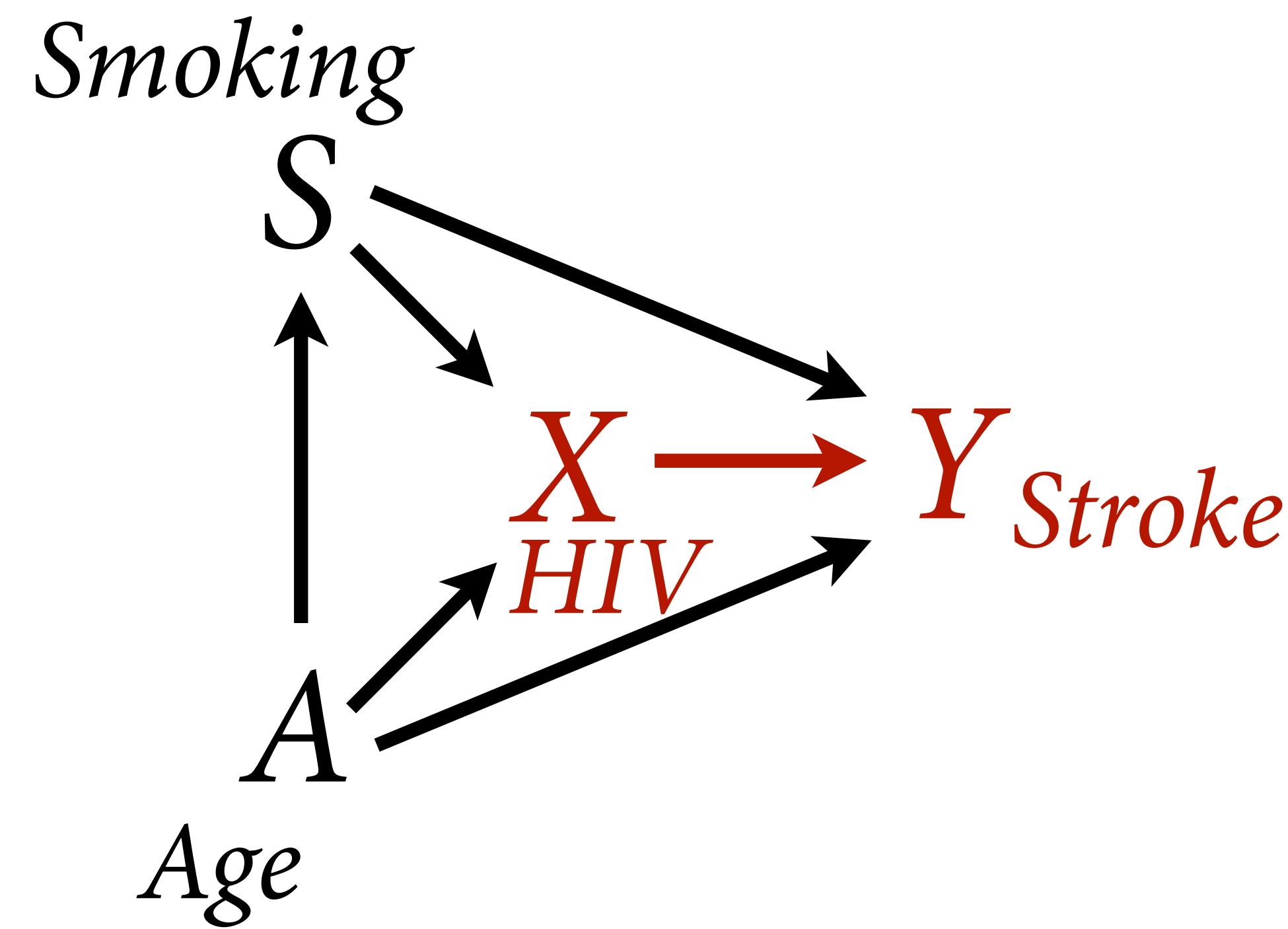
Not all coefficients are causal effects

Statistical model designed to identify  $X \rightarrow Y$  will not also identify effects of control variables

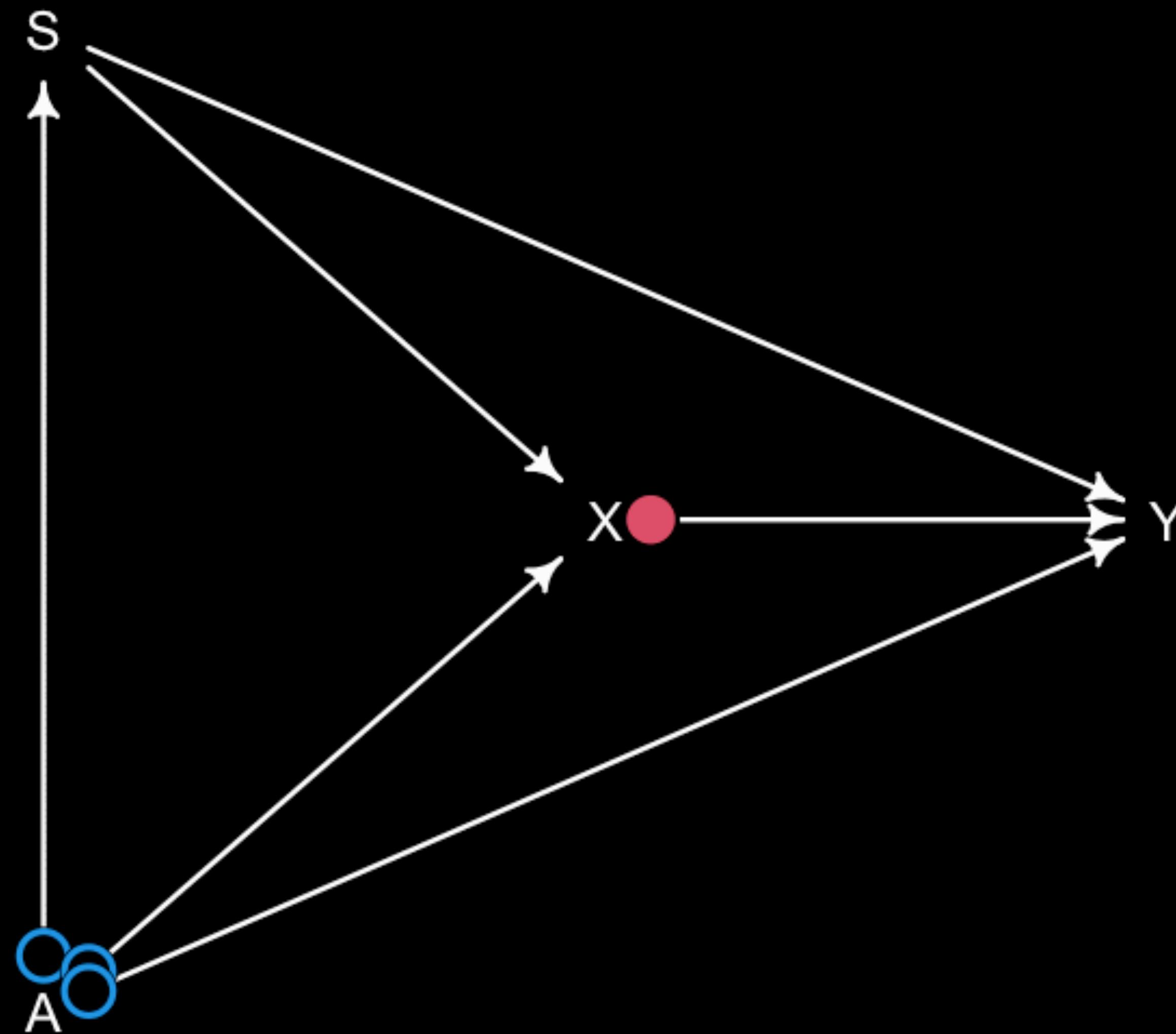
Table 2 is dangerous

TABLE 2—ESTIMATED PROBIT MODELS  
FOR THE USE OF A SCREEN

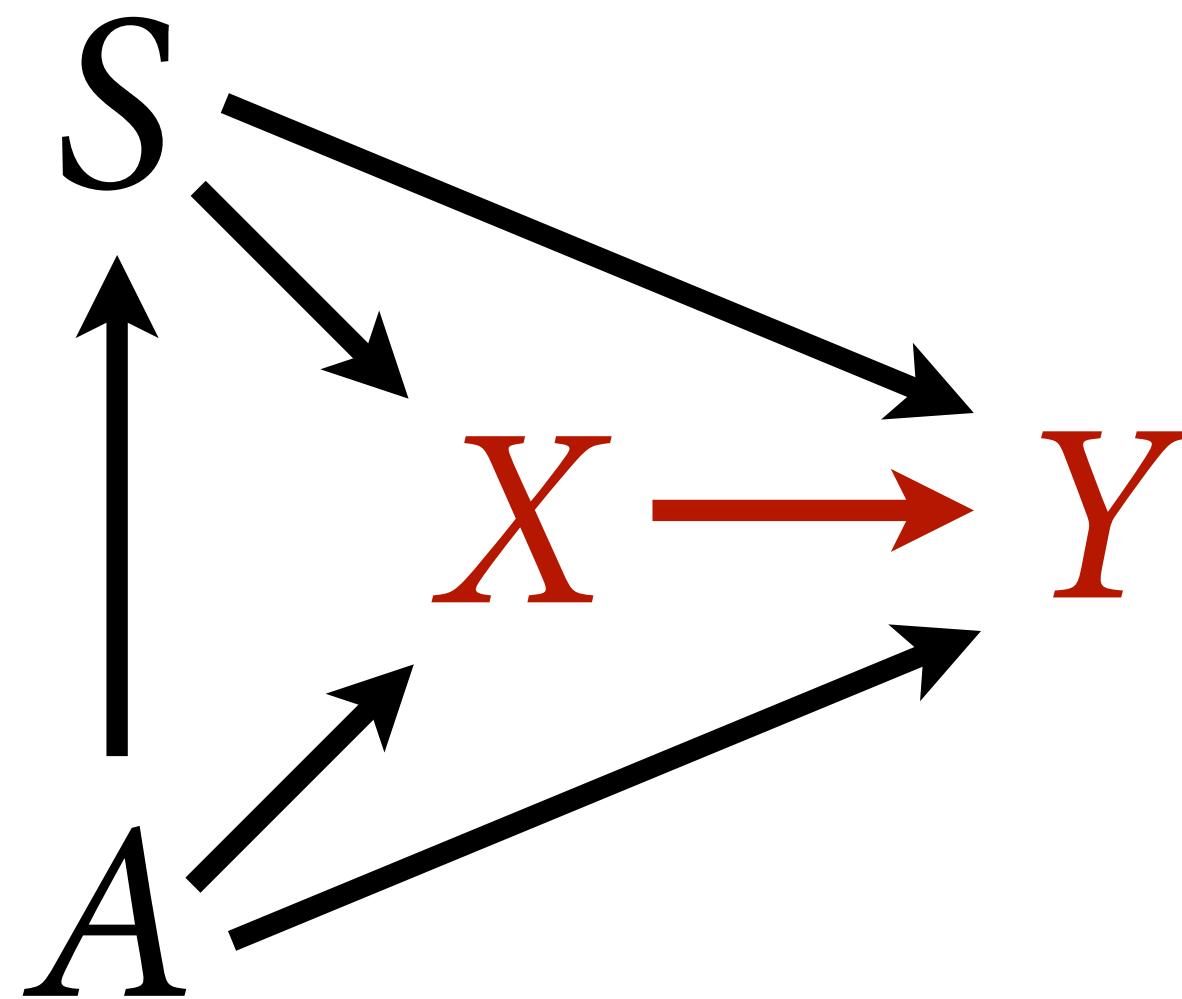
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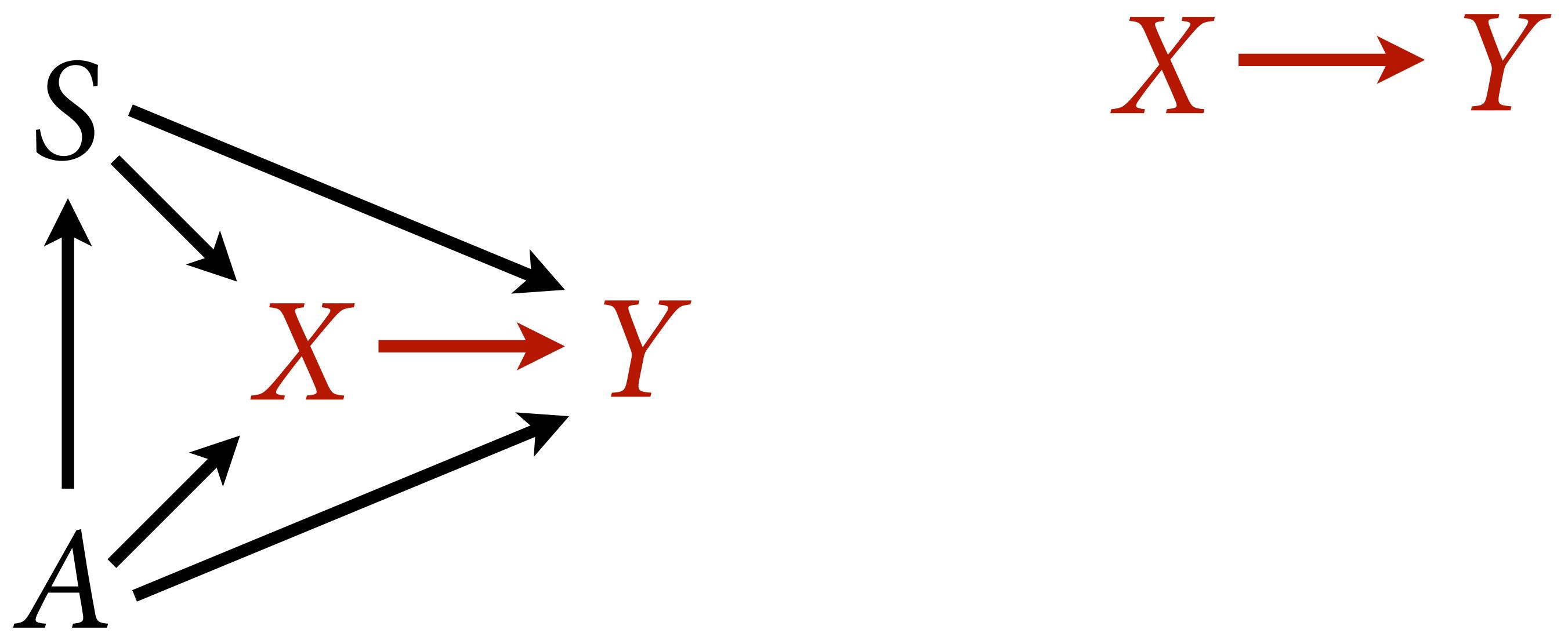
Westreich & Greenland 2013 The Table 2 Fallacy



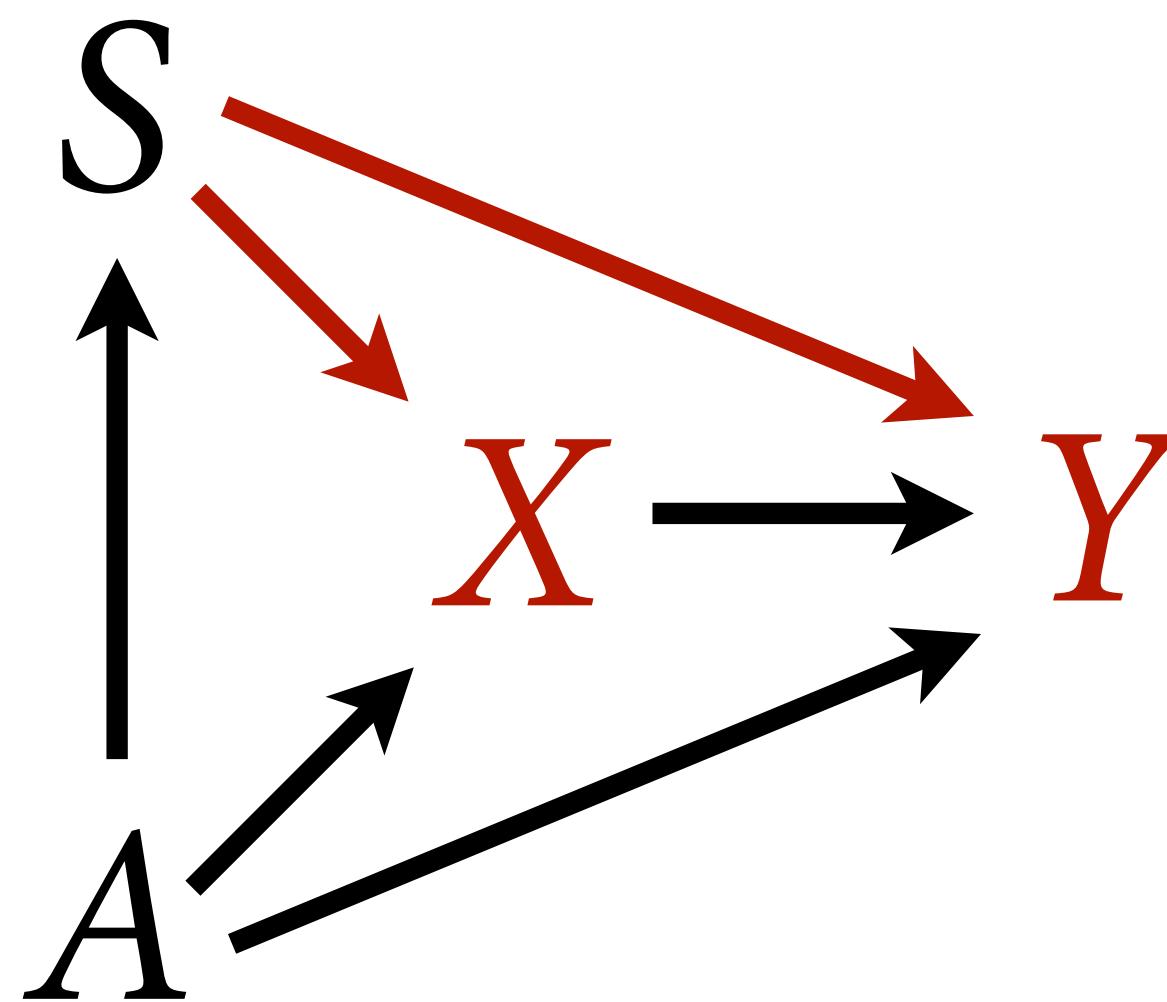
# Use Backdoor Criterion



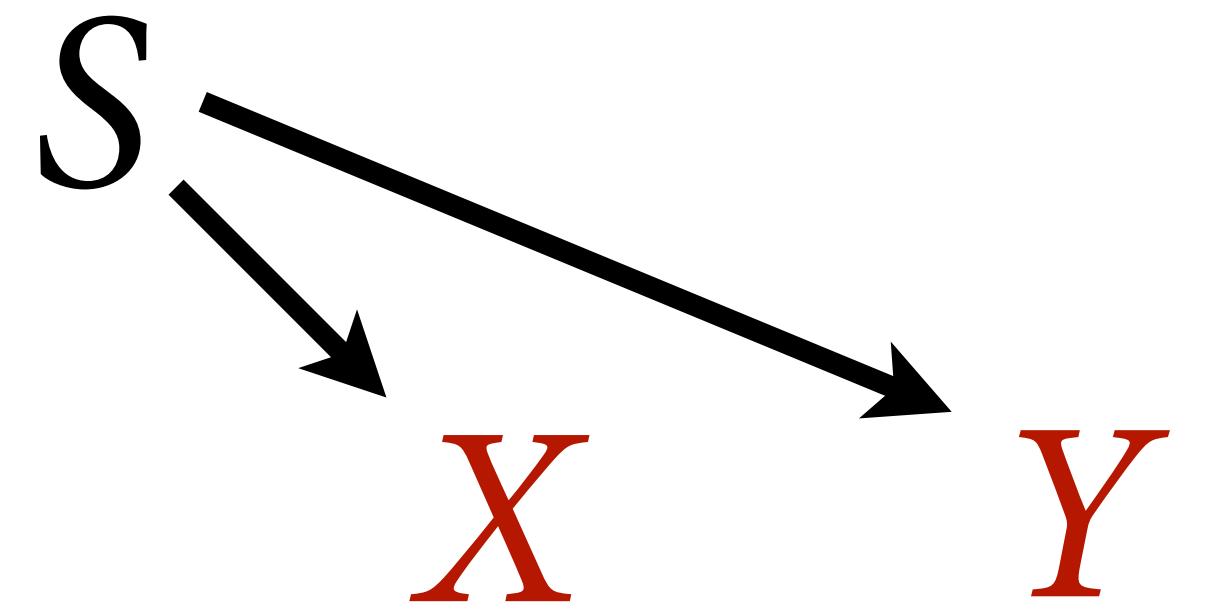
# Use Backdoor Criterion



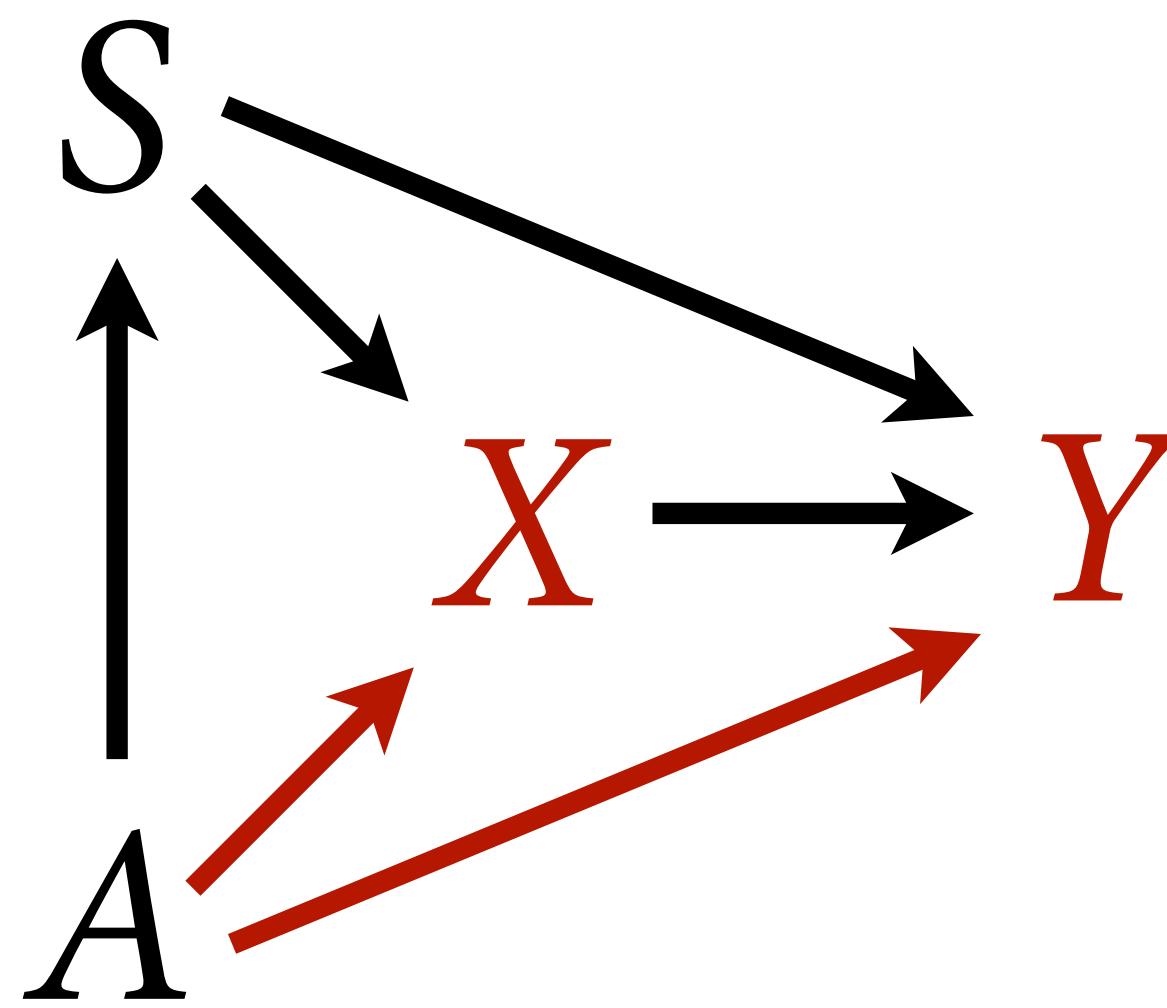
# Use Backdoor Criterion



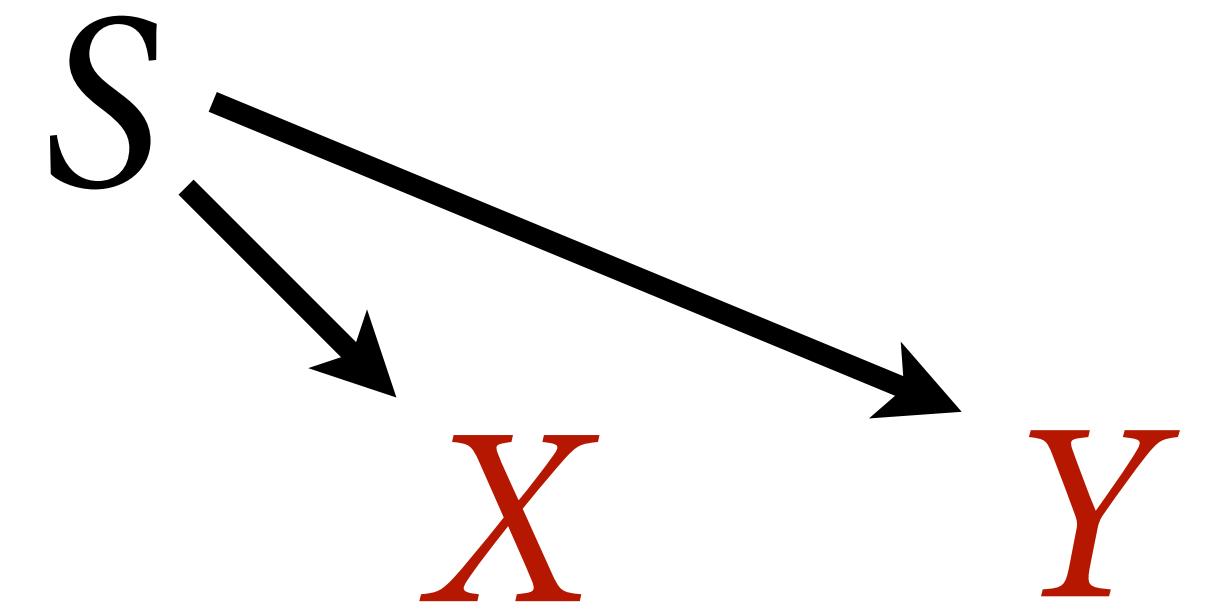
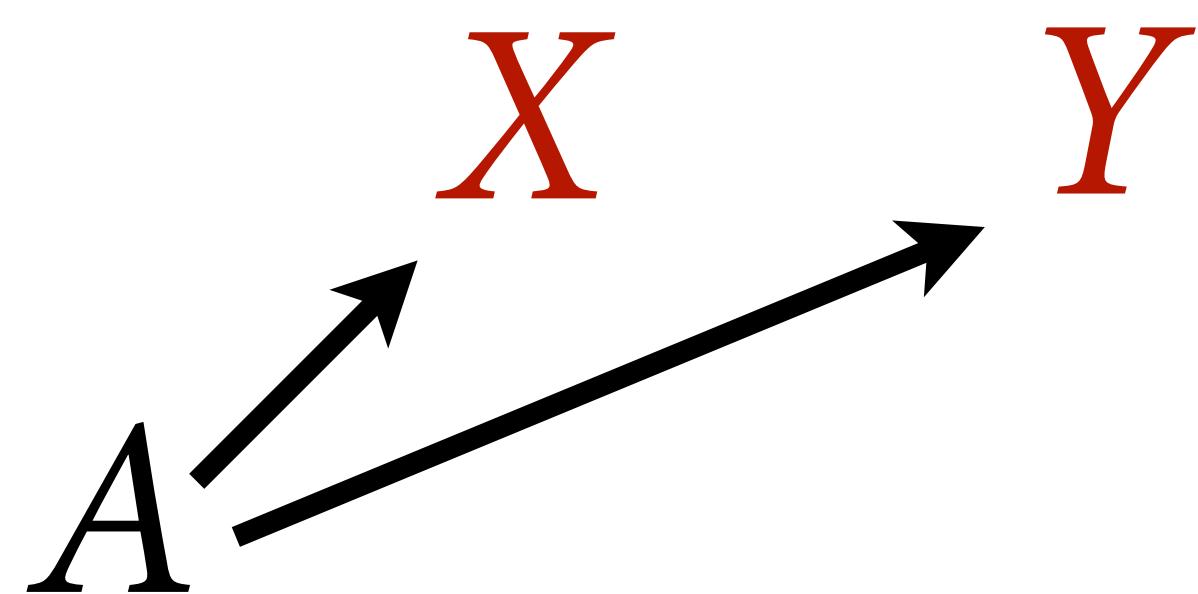
$X \rightarrow Y$



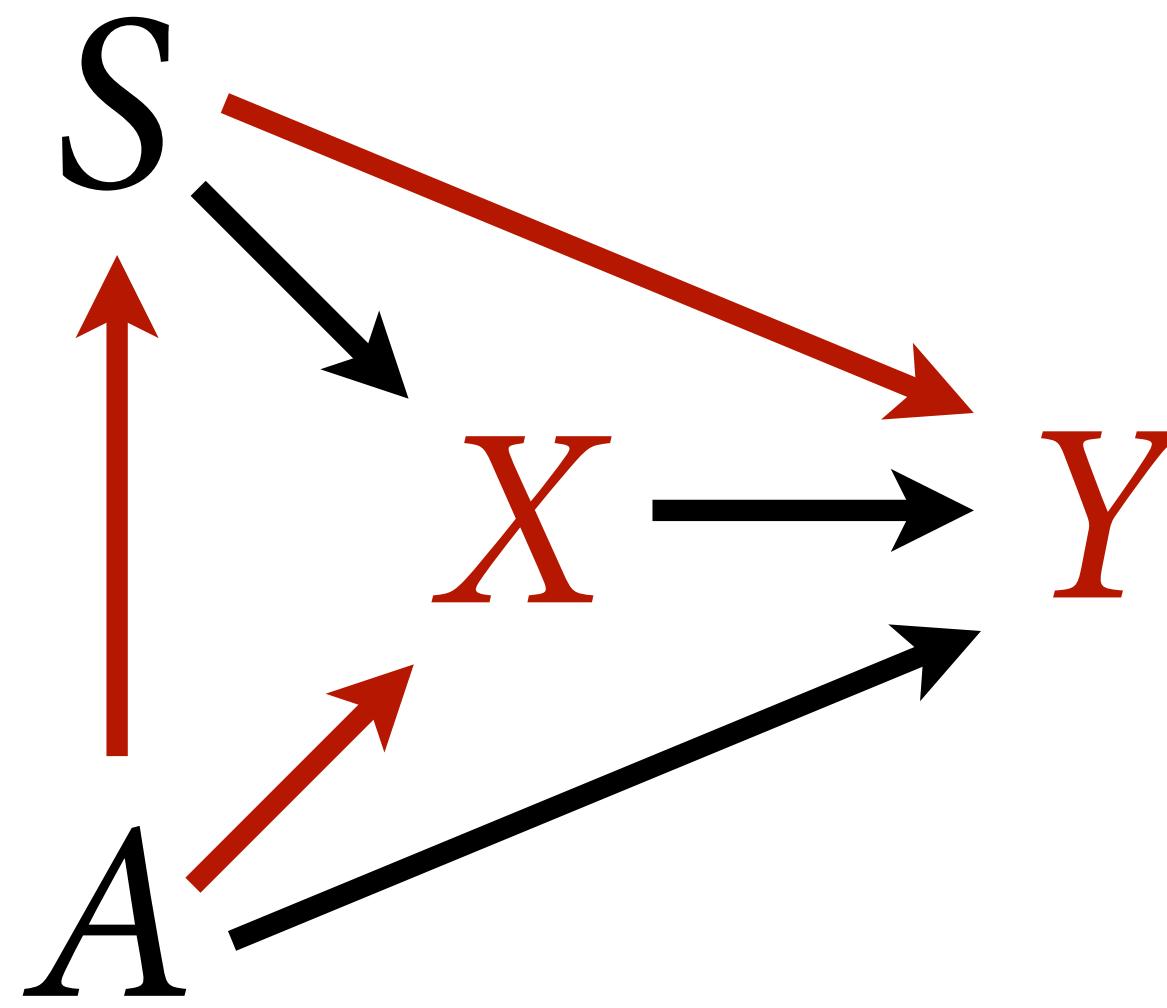
# Use Backdoor Criterion



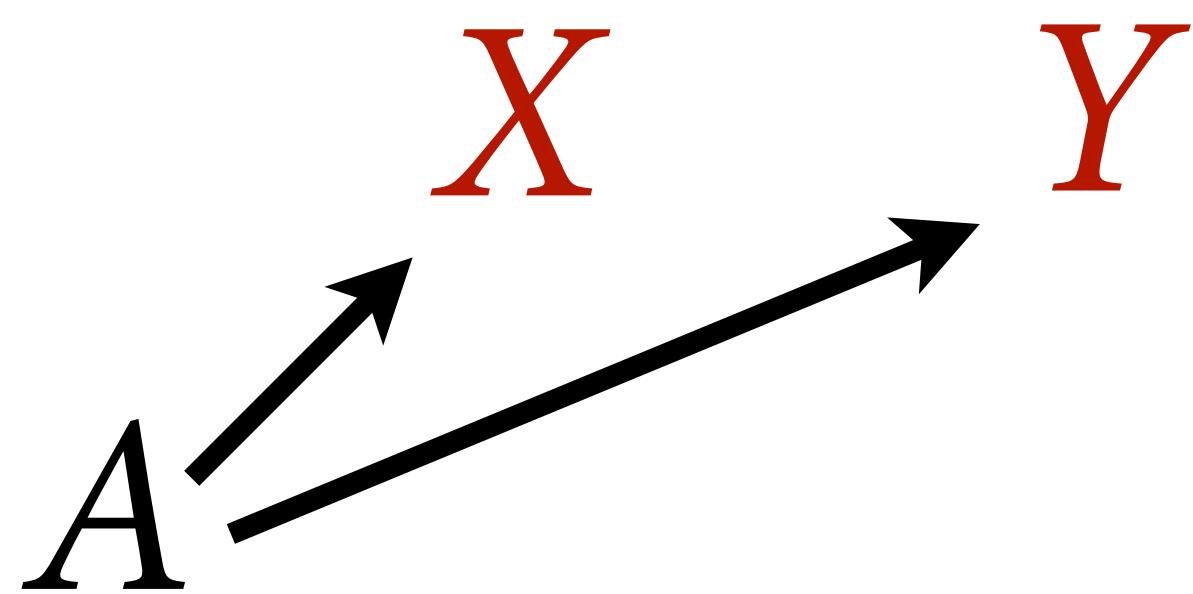
$$X \rightarrow Y$$



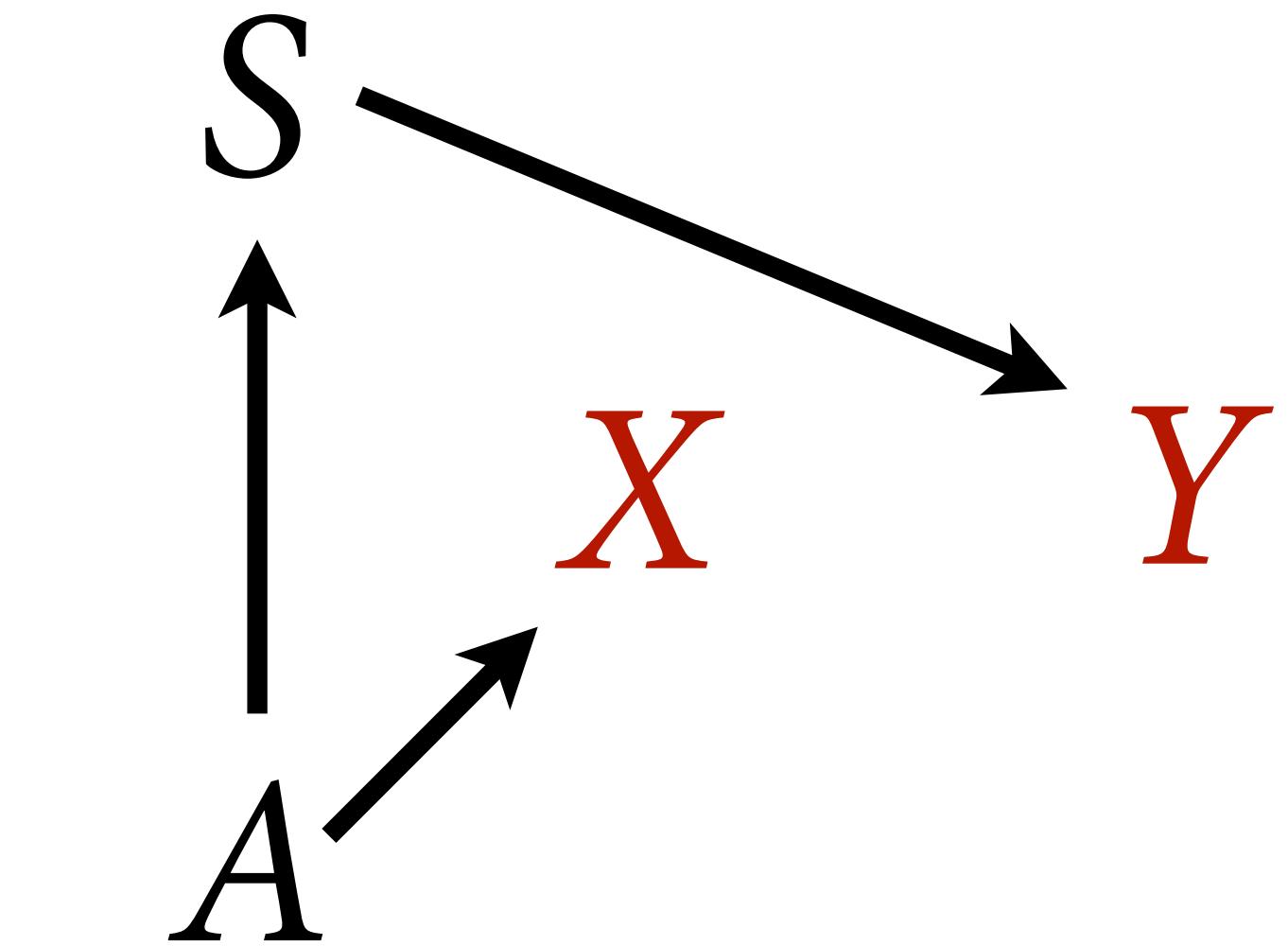
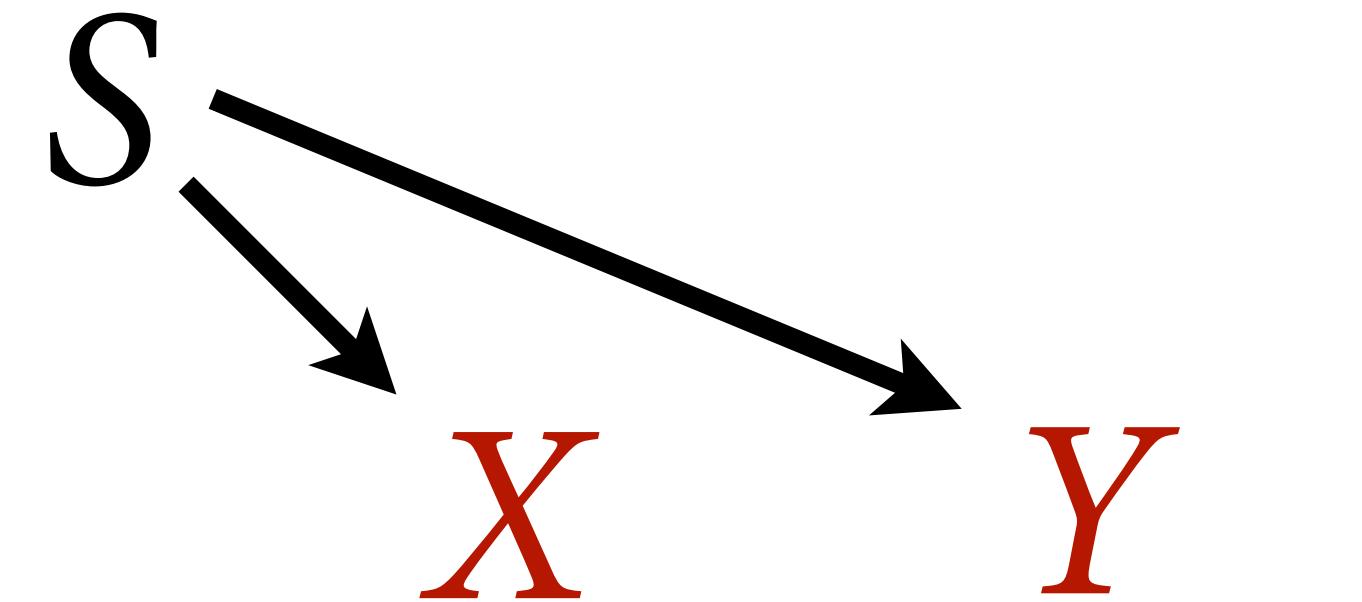
# Use Backdoor Criterion



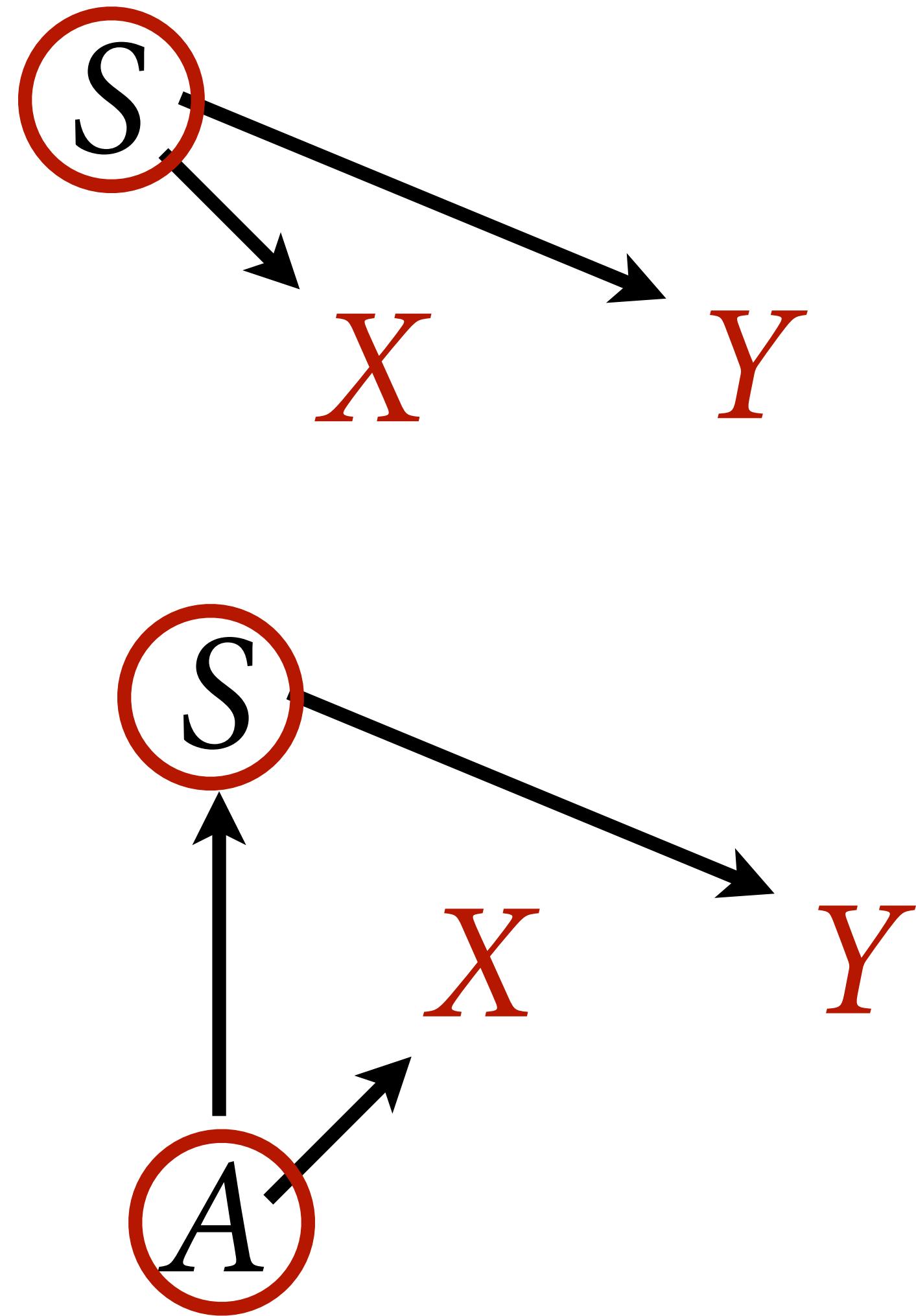
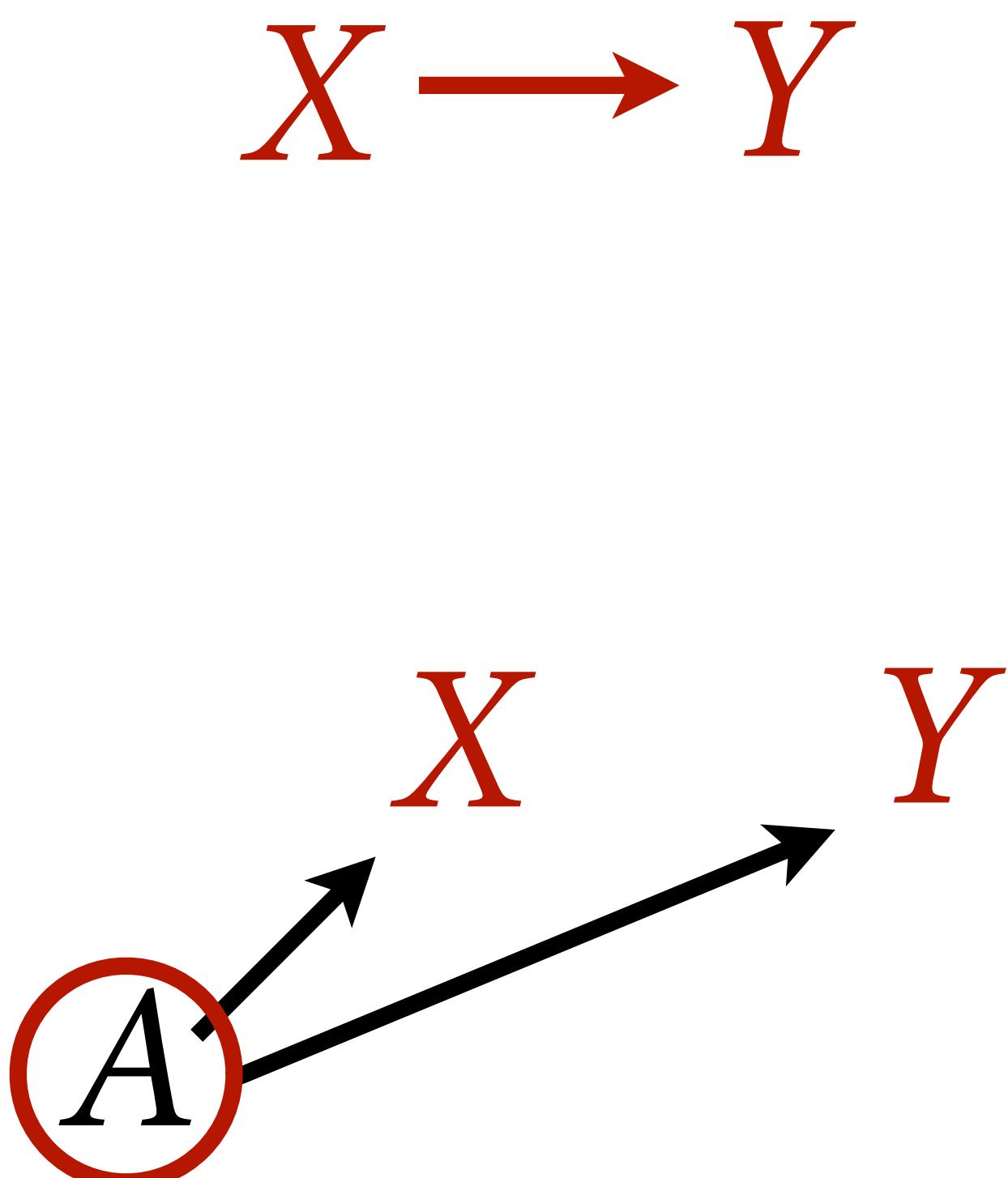
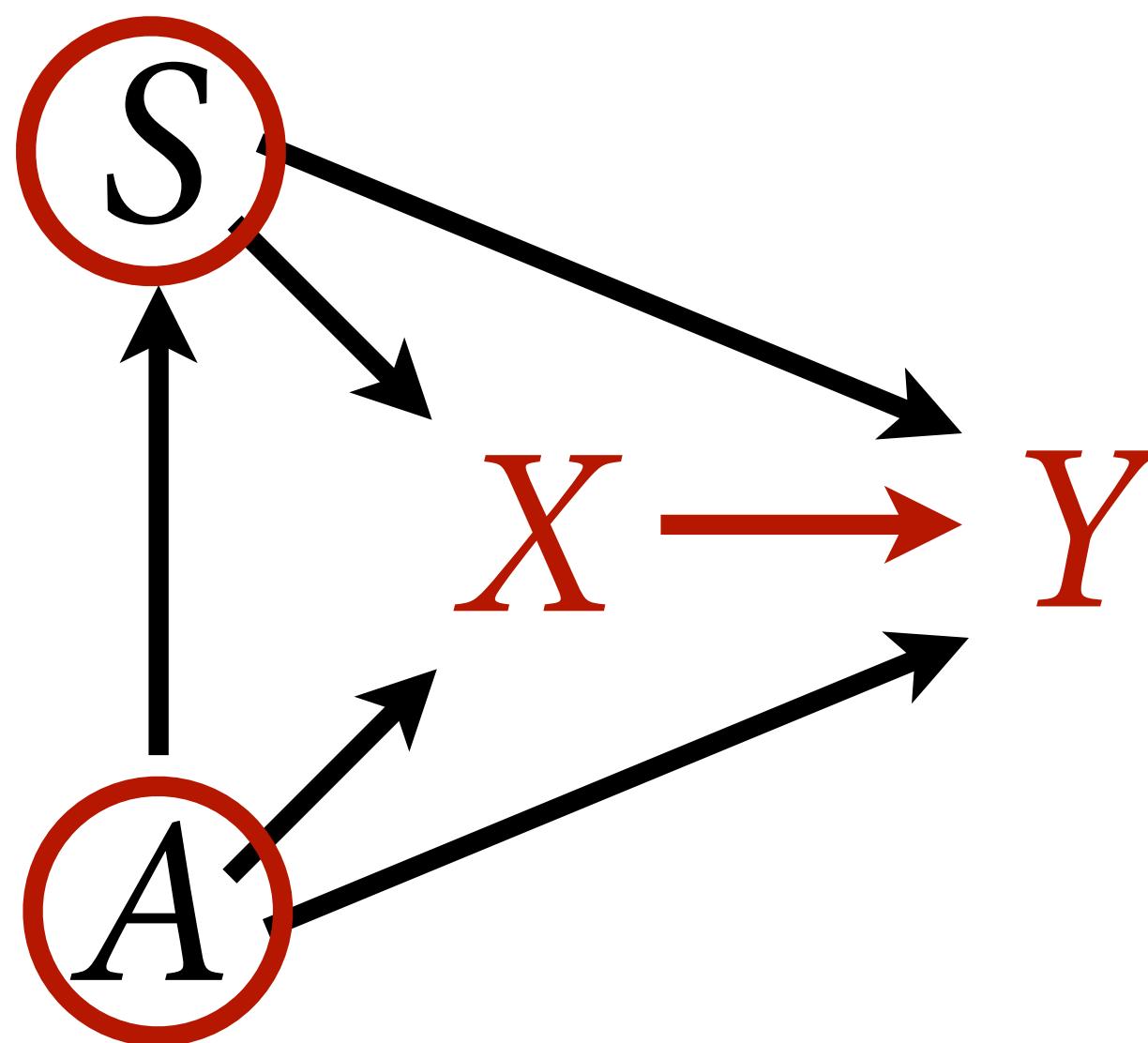
$$X \rightarrow Y$$

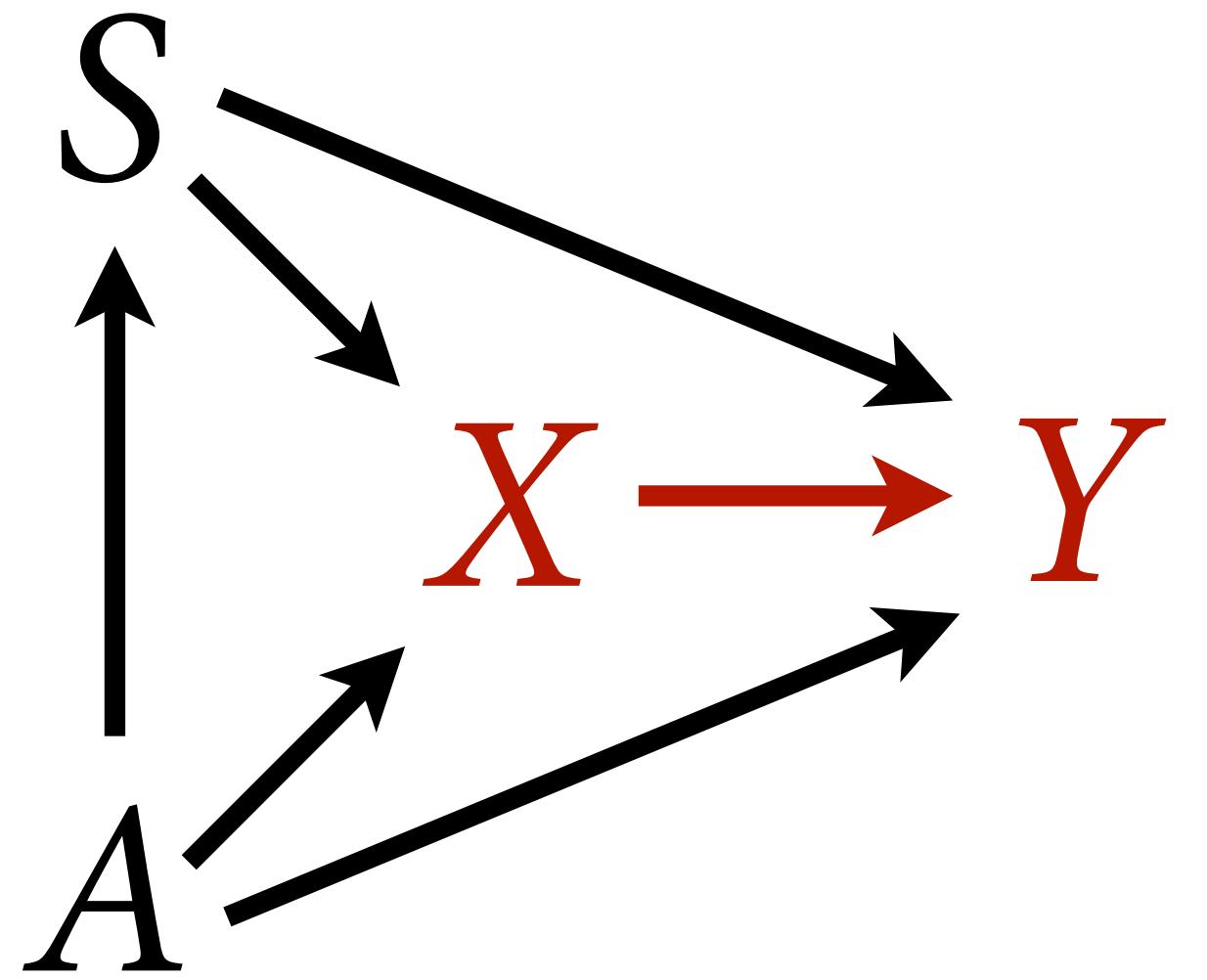


$$X \rightarrow Y$$



# Use Backdoor Criterion



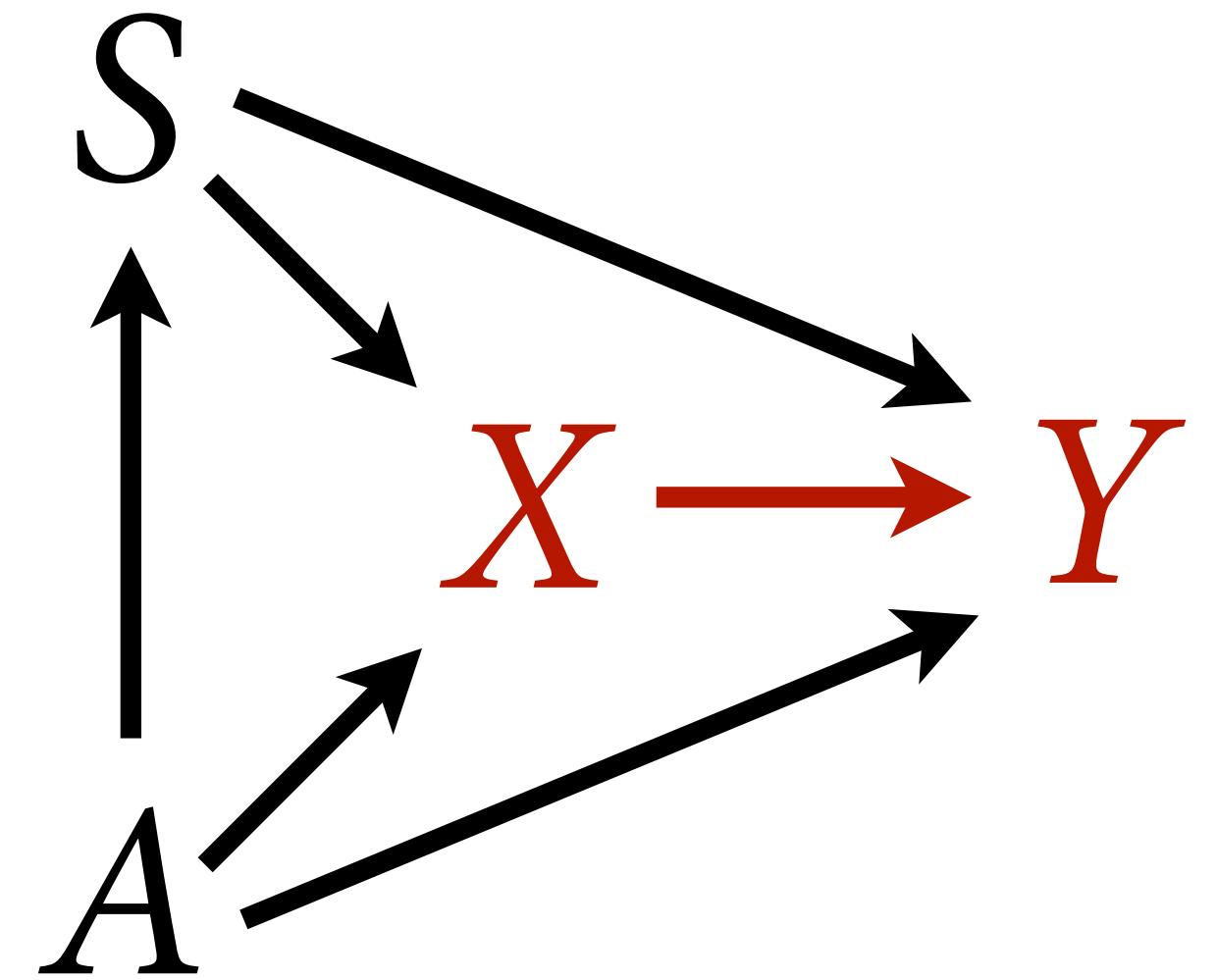


$$Y_i \sim \text{Normal}(\mu_i, \sigma)$$

$$\mu_i = \alpha + \beta_X X_i + \beta_S S_i + \beta_A A_i$$

**X**

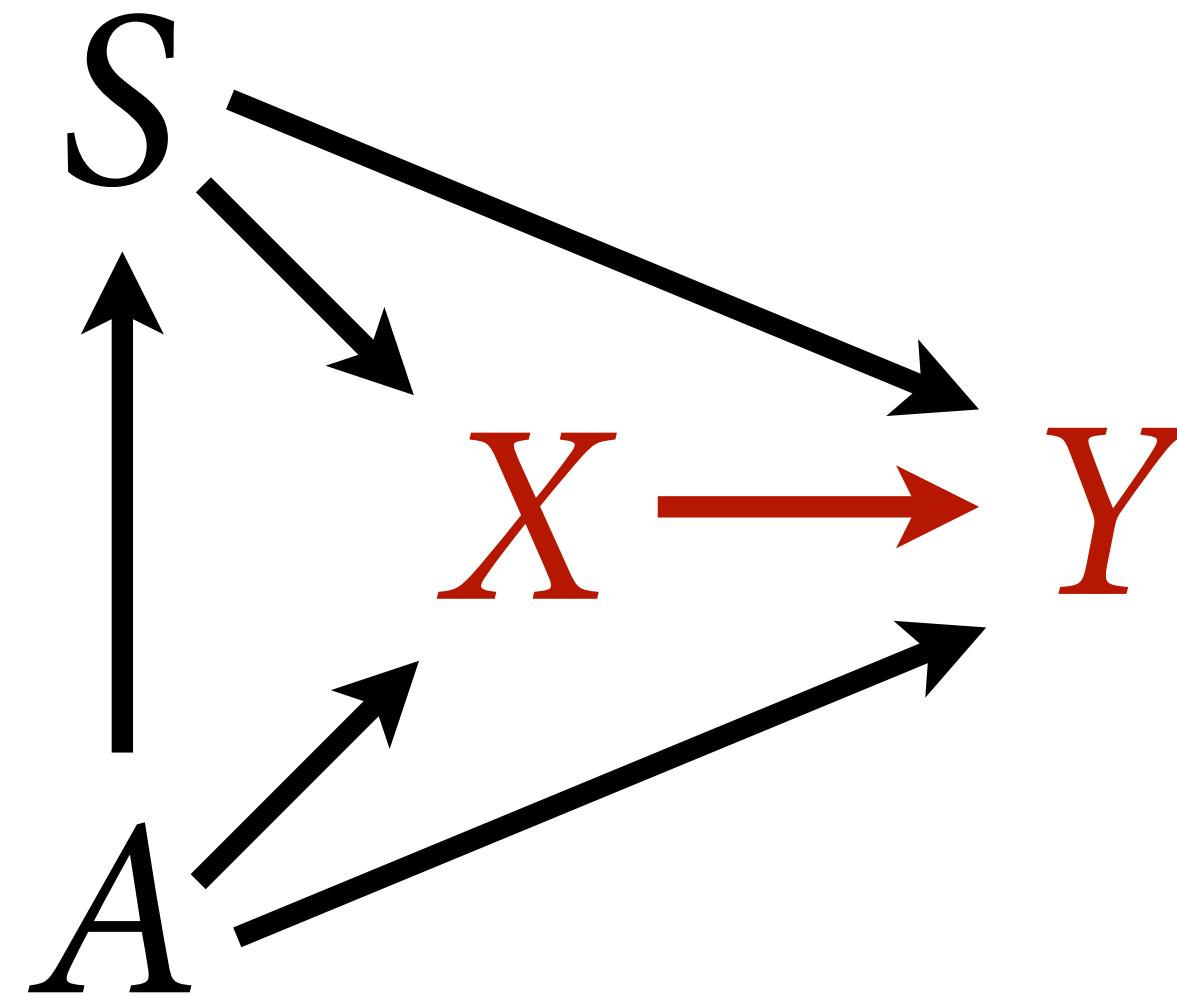
*Unconditional*



Confounded by  $A$   
and  $S$

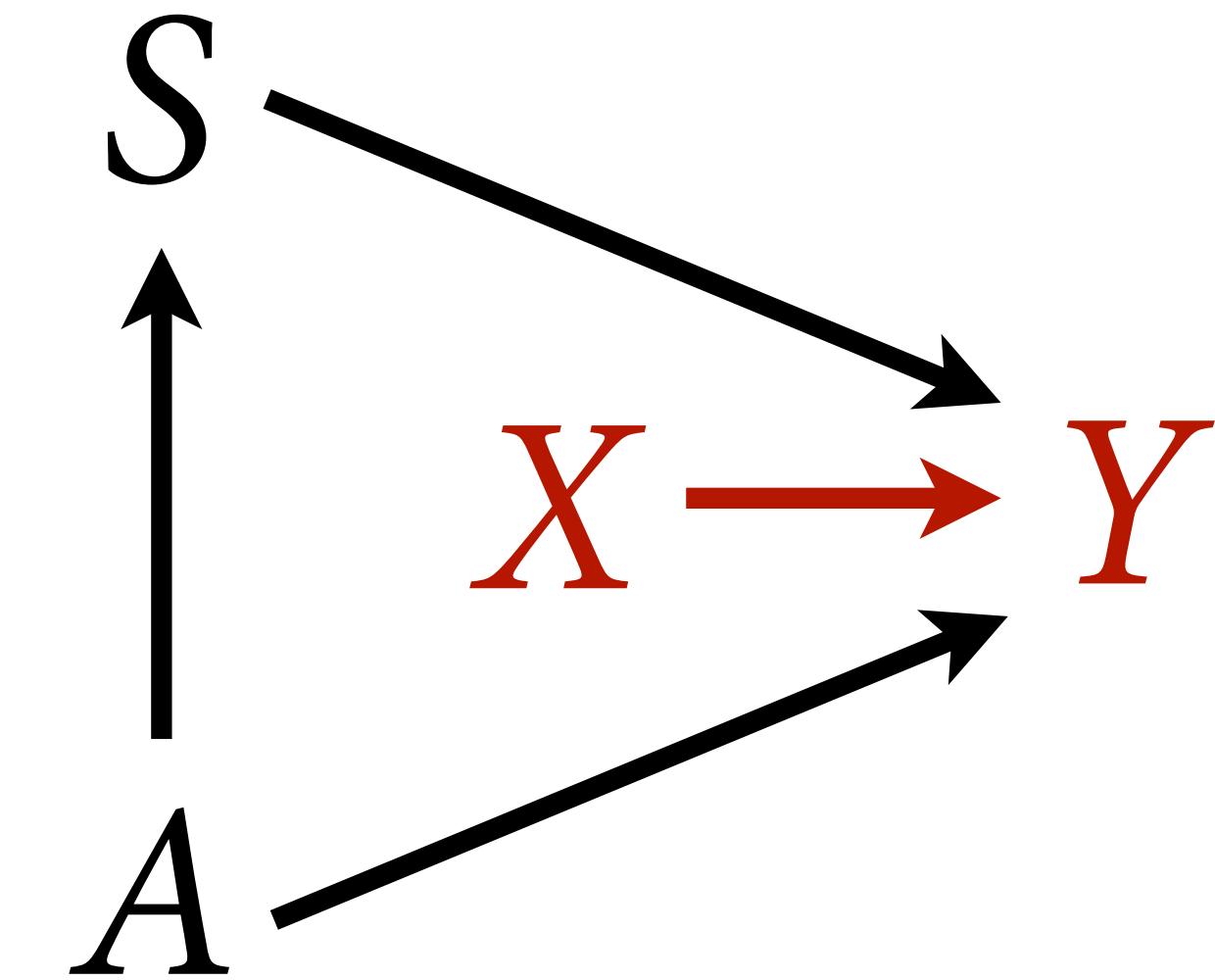
# X

*Unconditional*



Confounded by A  
and S

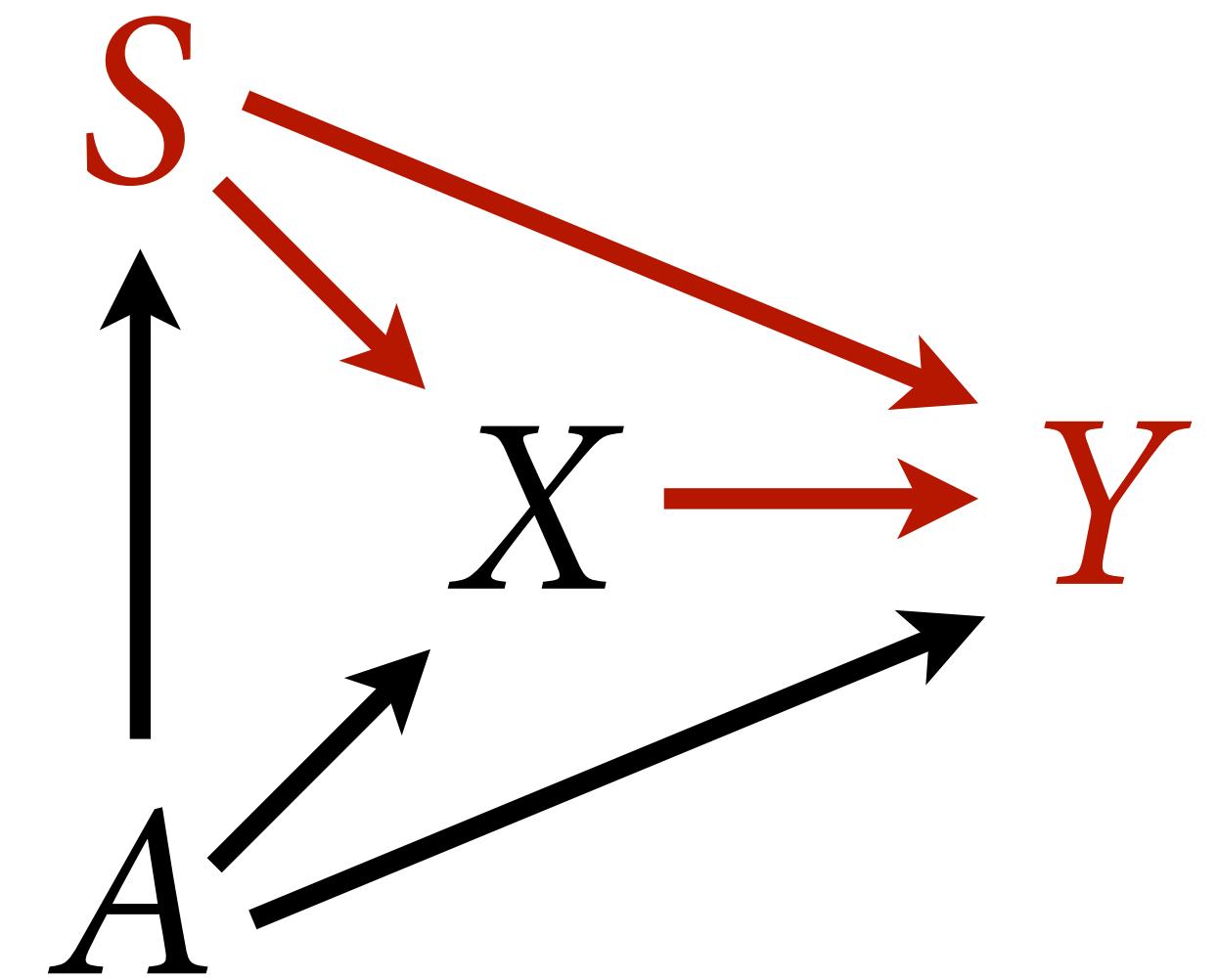
*Conditional on A and S*



Coefficient for **X**:  
Effect of X on Y  
(still must  
marginalize!)

# S

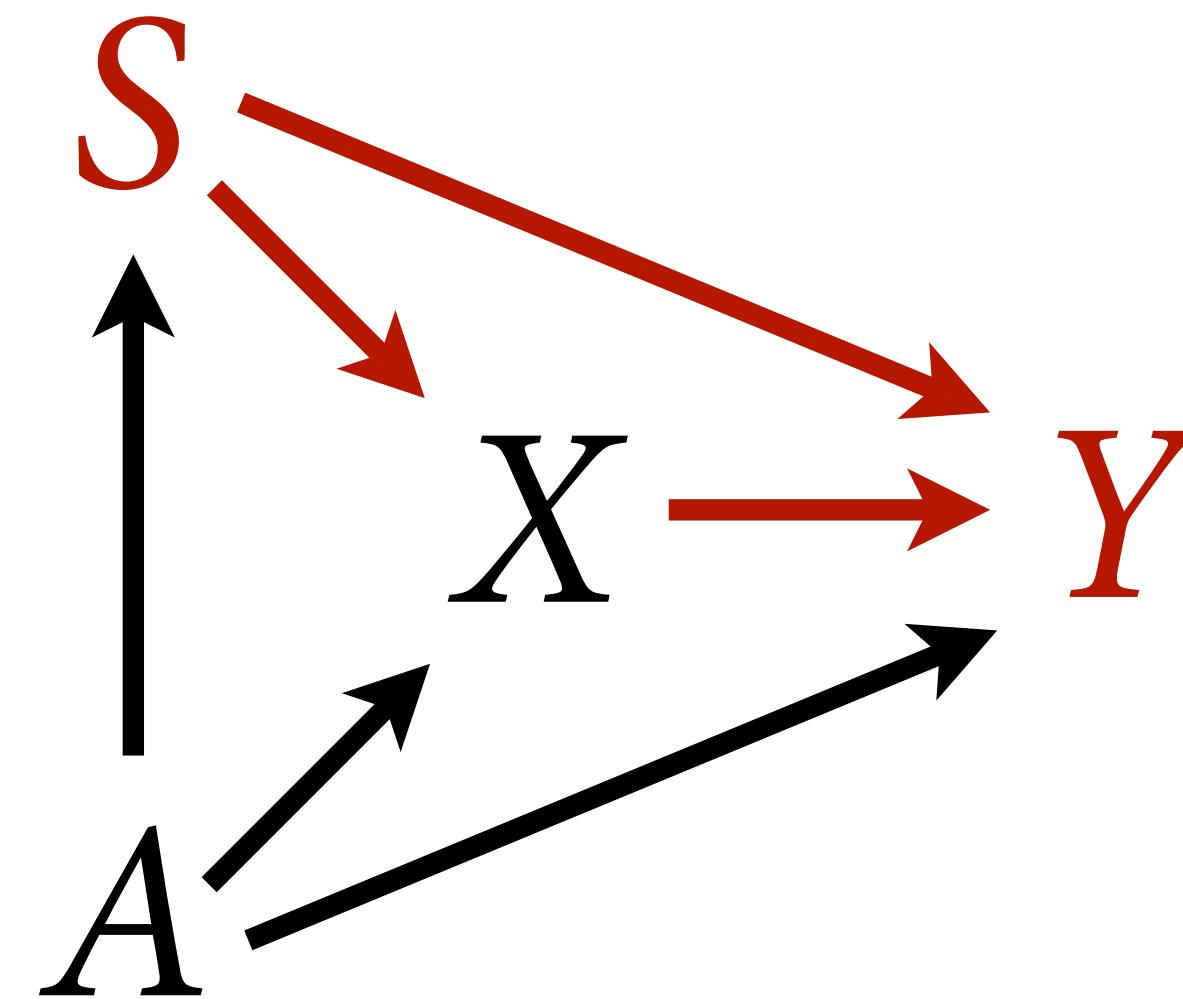
## *Unconditional*



Effect of  $S$   
confounded by  $A$

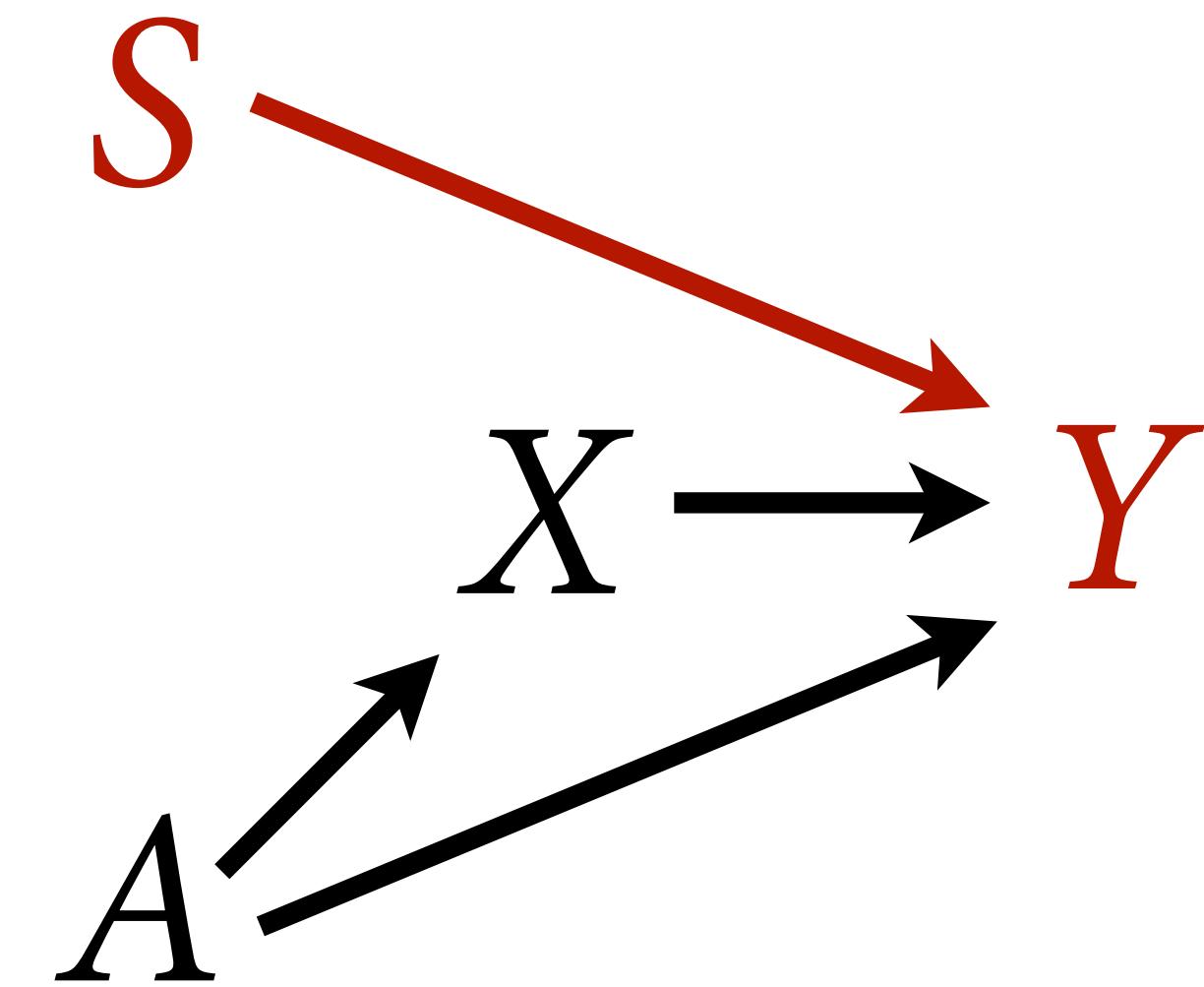
**S**

*Unconditional*



Effect of **S**  
confounded by **A**

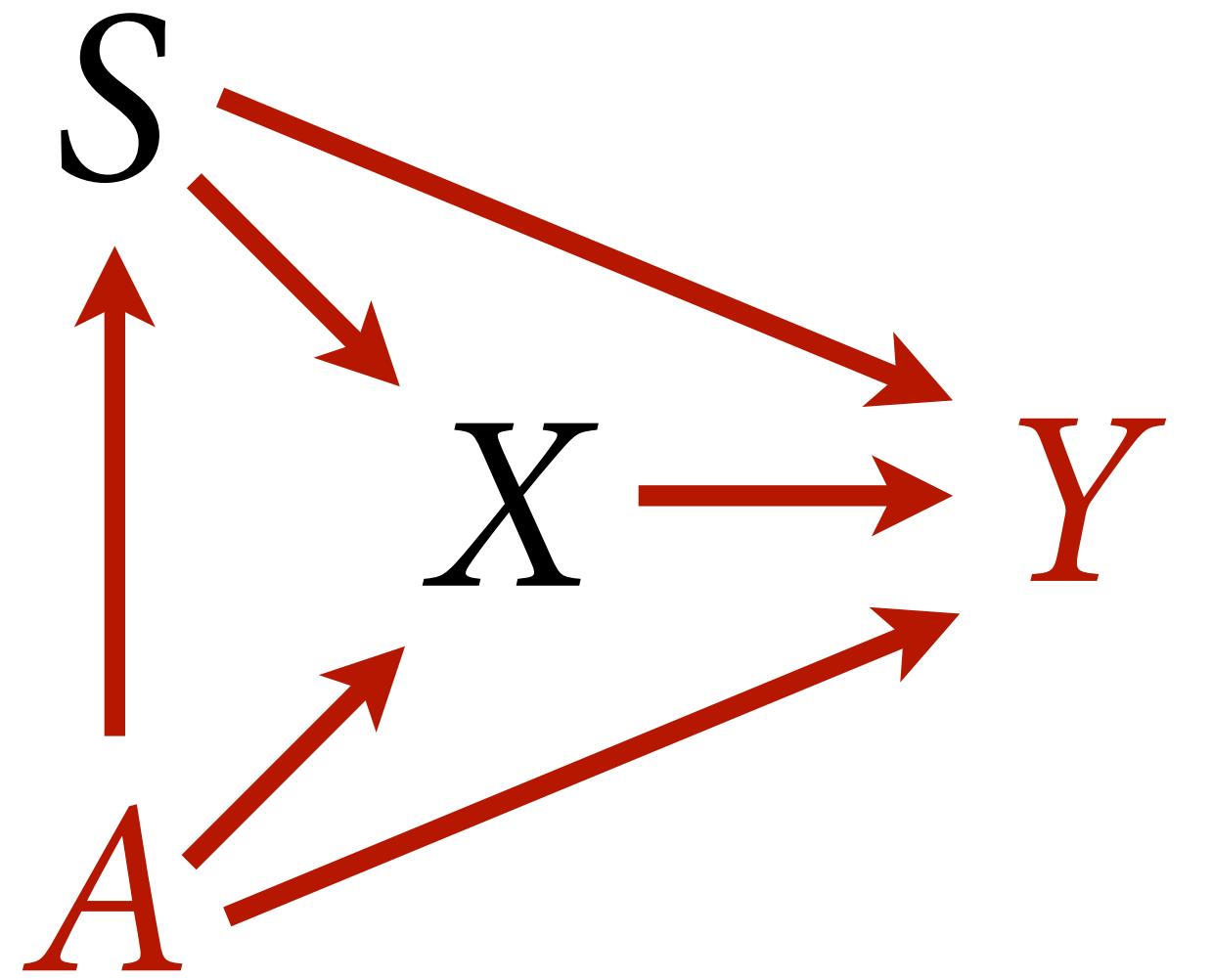
*Conditional on A and X*



Coefficient for **S**:  
**Direct** effect of **S** on **Y**

# A

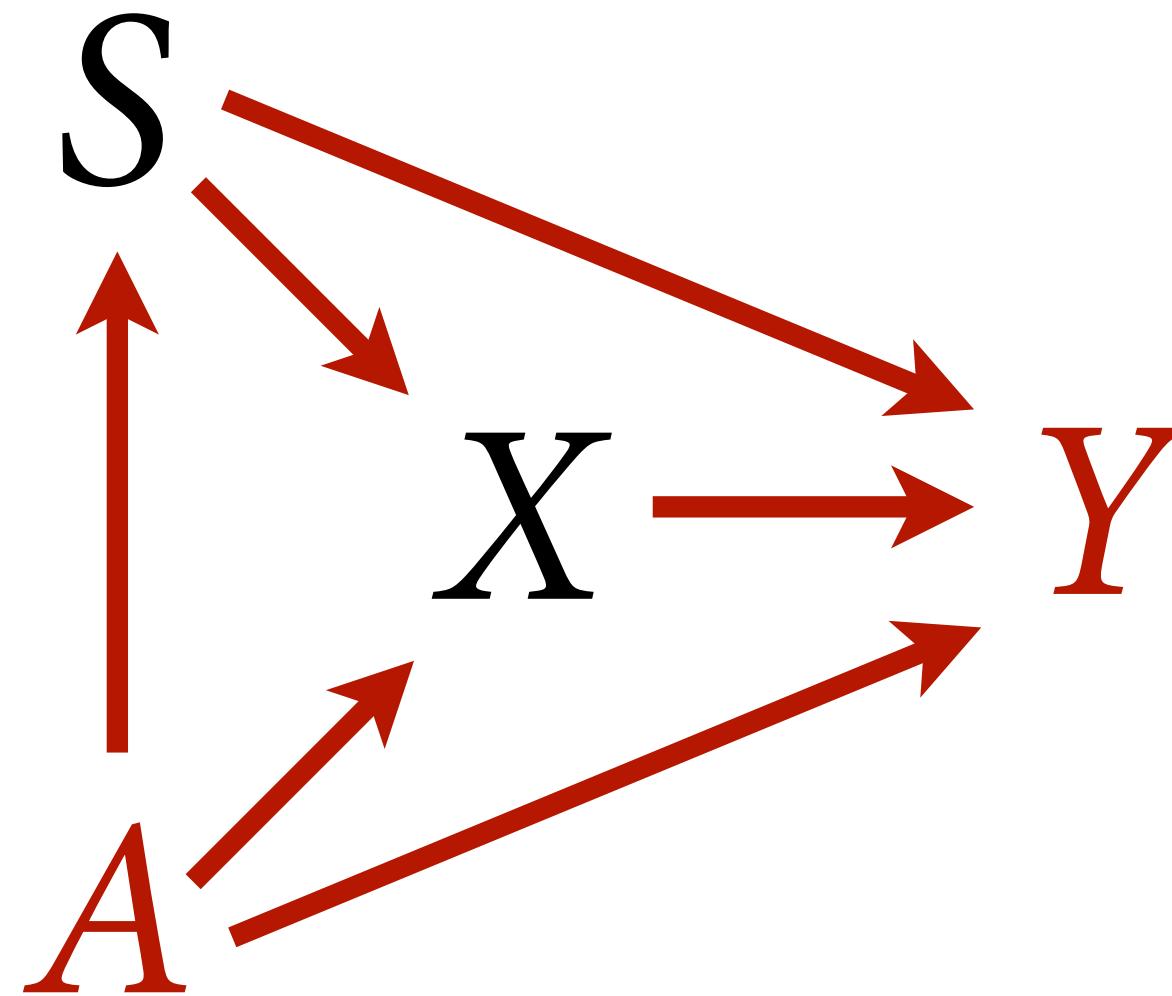
## *Unconditional*



Total causal effect  
of  $A$  on  $Y$  flows  
through all paths

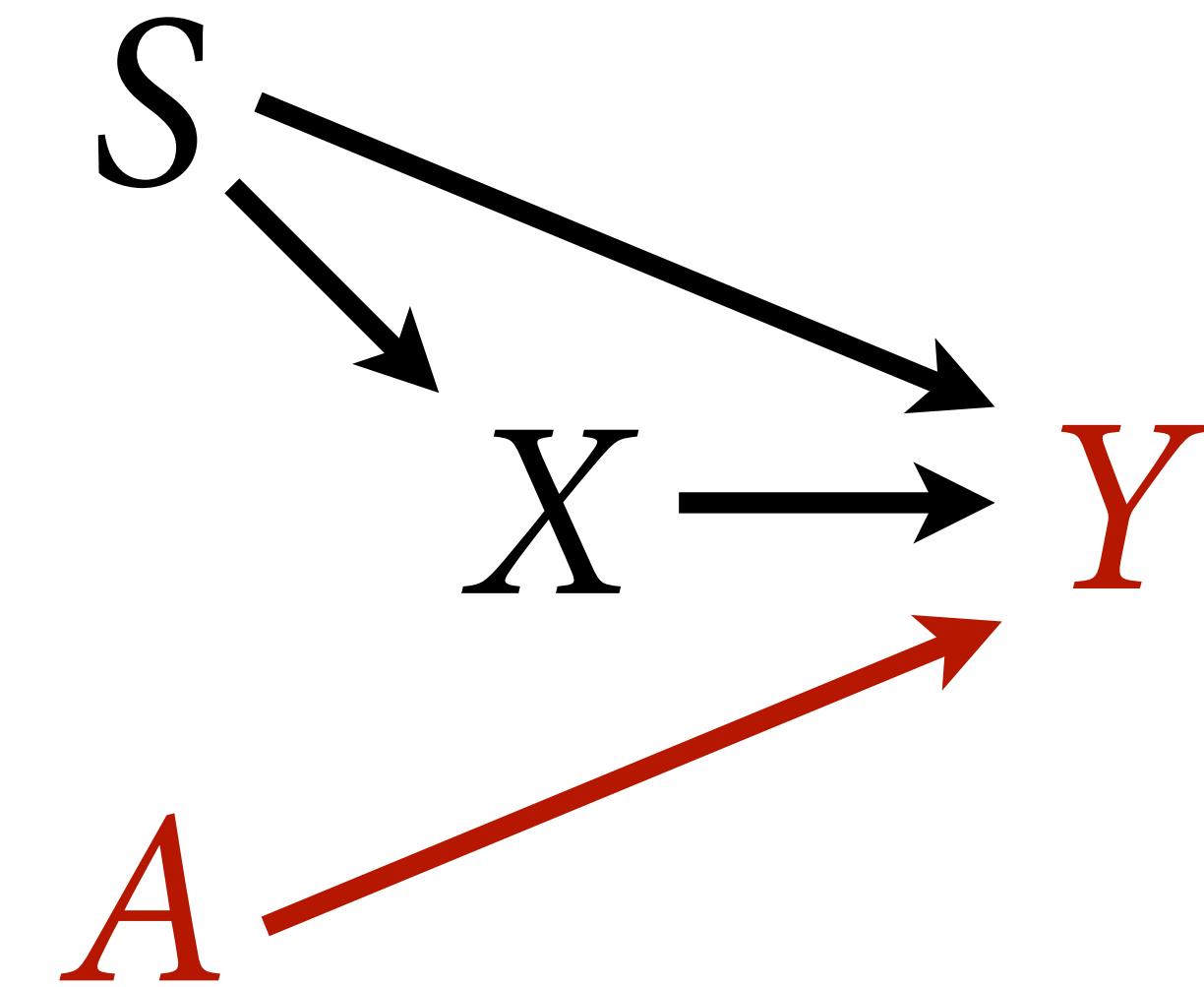
# A

*Unconditional*

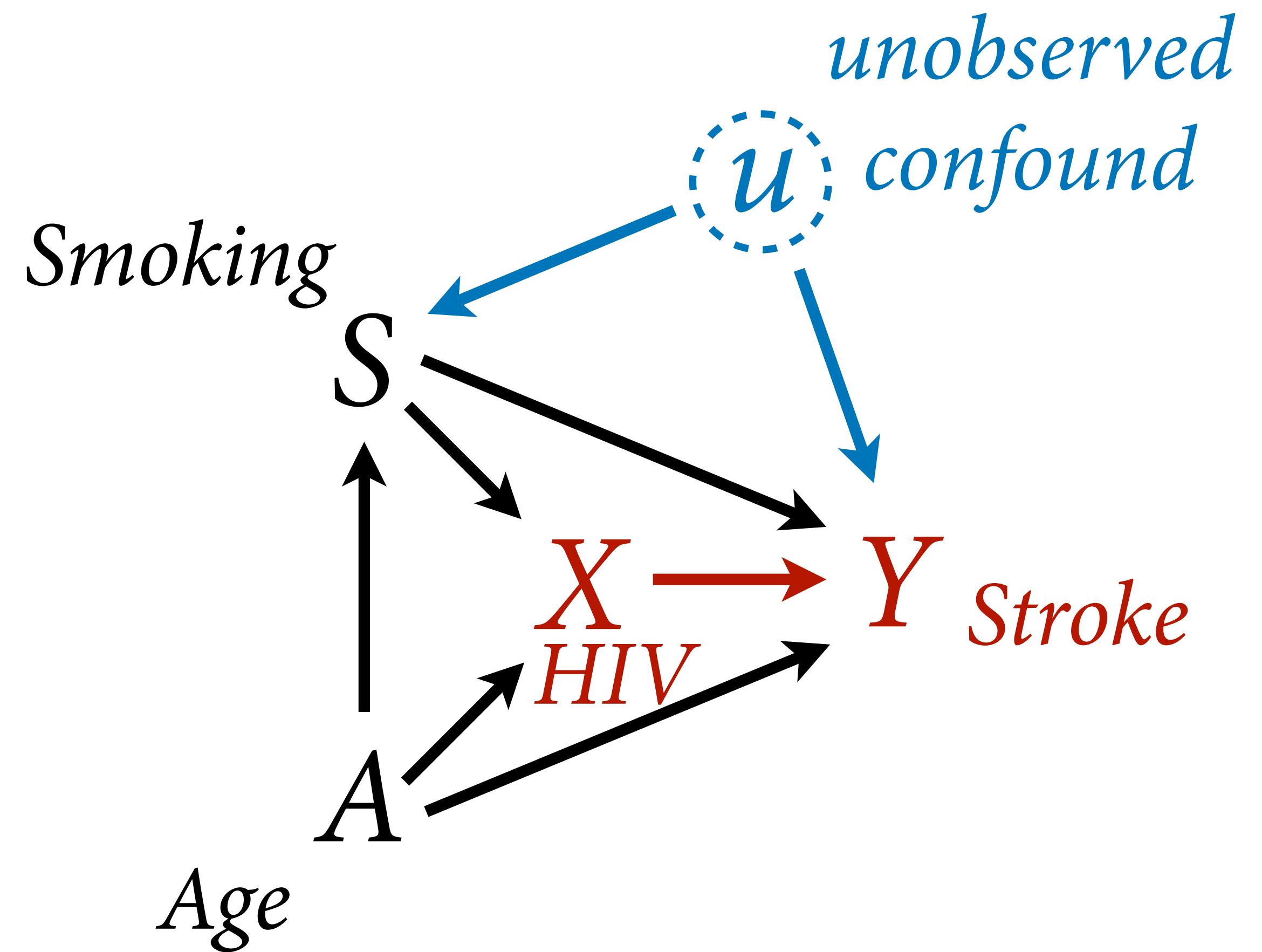


Total causal effect  
of  $A$  on  $Y$  flows  
through all paths

*Conditional on  $X$  and  $S$*



Coefficient for  $A$ :  
**Direct** effect of  $A$  on  $Y$



# Table 2 Fallacy

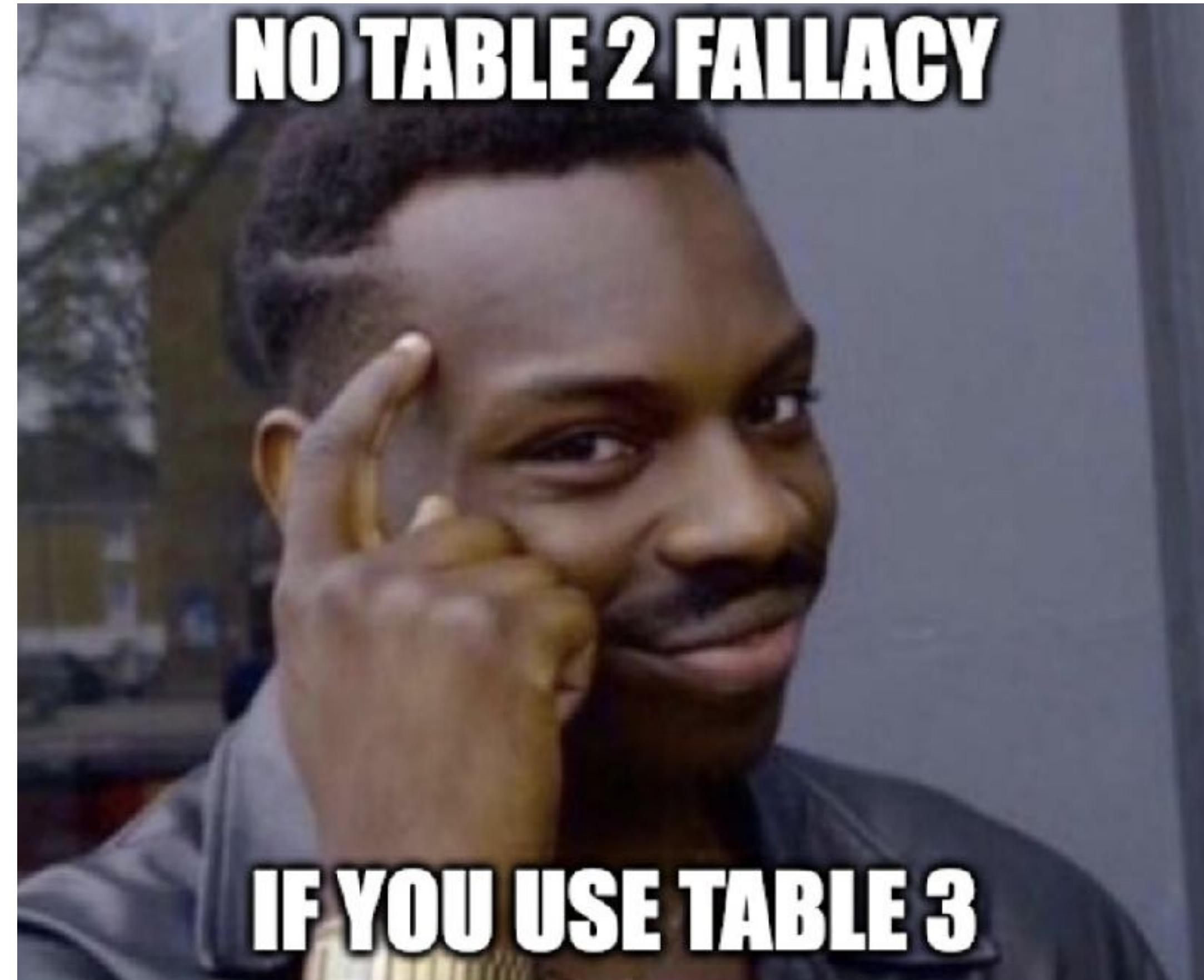
Not all coefficients created equal

So do not present them as equal

Options:

- Do not present control coefficients
- Give explicit interpretation of each

No interpretation without causal representation



**NO TABLE 2 FALLACY**

**IF YOU USE TABLE 3**

