



ECON526: Quantitative Economics with Data Science Applications

Directed Graphical Models and Causality

Phil Solimine

philip.solimine@ubc.ca

University of British Columbia

Table of contents

- Overview
- Directed Graphical Models
- Building a Causal Graph
- Aside: Cycles
- Dependence Flows
- Viualizing Bias
- Confounding
- Selection
- Choosing Covariates
- Factorizing the Joint Distribution

Overview

Summary

- Previously in the course, we talked at a high level about some of the barriers to causal inference
- We used the potential outcomes framework to discuss the idea of a treatment effect
- Then we discussed the idea of a randomized experiment as a way to identify a treatment effect
 - However, we mentioned that there are many situations where we cannot run a randomized experiment
- Today, we will discuss the idea of using a **graphical model** as a way to analyze whether you can truly identify a treatment effect

Directed Graphical Models

Conditional Independence

- Recall that two random variables X and Y are **conditionally independent** given a third random variable Z if and only if the following holds:
 - $P(X|Z) = P(X|Z \cap Y)$
 - Equivalently, $P(X \cap Y|Z) = P(X|Z)P(Y|Z)$
 - We will denote this as $X \perp Y|Z$
- In the context of potential outcomes, we require that $(Y_0, Y_1) \perp T|X$
 - This means that the potential outcomes are independent of the treatment assignment, given the covariates

Directed Graphical Models

- Complete independence is rare in complex systems. However, we can often find conditional independence relationships, that help inform our choice of statistical model.
- We can visualize conditional independence relationships using a **Bayesian** or **directed graphical model**
- A Bayesian graphical model is a directed graph where:
 - The nodes or vertices represent random variables
 - The links or edges represent conditional independence relationships

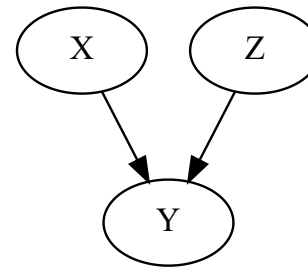
Directed Graphical Models

- Fundamentally, a graphical model is a way to represent how a joint probability distribution factorizes into a product of conditional distributions.
- For our purposes, we will usually interpret the edges as causal relationships.
- Furthermore, the graphs we draw will be **acyclic**, meaning that there are no loops in the graph.
 - In other words, there is no way to start at a node and follow the arrows to get back to the same node
 - This is because we are interested in **causal** relationships, and a cycle would imply that there is a feedback loop where a variable causes itself
- Since these graphs are both directed and acyclic, we call them **directed acyclic graphs** or **DAGs**

Directed Graphical Models

- A directed graphical model might look something like this:

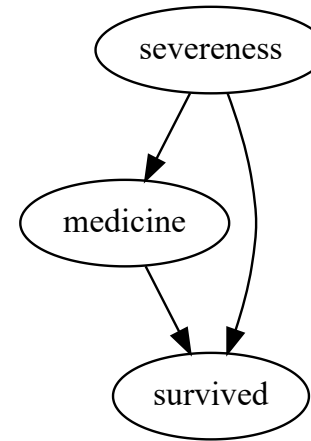
```
1 import graphviz as gr
2
3 g = gr.Digraph()
4 g.node('X')
5 g.node('Y')
6 g.node('Z')
7 g.edge('X', 'Y')
8 g.edge('Z', 'Y')
9 g
```



- Here, we have three random variables: X , Y , and Z
 - X and Z are independent
 - Y depends on both X and Z

Directed Graphical Models

```
1 g = gr.Digraph()
2 g.edge("medicine", "survived")
3 g.edge("severeness", "survived")
4 g.edge("severeness", "medicine")
5 g
```



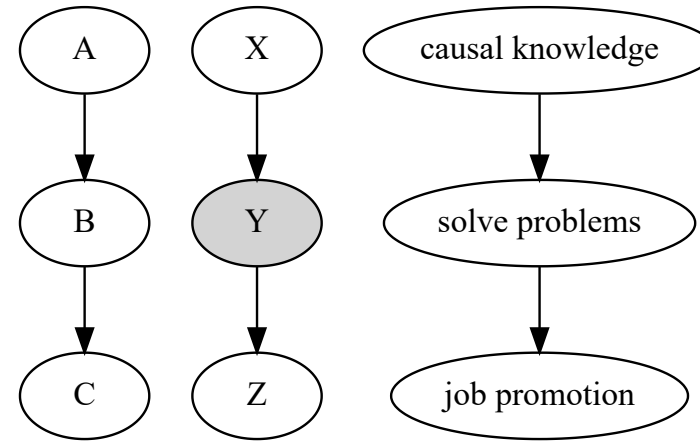
- We use arrows to indicate the direction of the conditional independence relationship
 - For example, *medicine* → *survived* means that *survived* depends on *medicine*, but not the other way around

Directed Graphical Models

- Directed graphical models are useful because they allow us to visualize conditional independence relationships, which can often be difficult to keep track of
- However, they are also useful because they allow us to determine whether or not we can identify a treatment effect.
- There are three very common sub-structures that appear in graphical models, that inform how dependence will flow through the model.

Directed Graphical Models

```
1 g = gr.Digraph()
2 g.edge("A", "B")
3 g.edge("B", "C")
4
5 g.edge("X", "Y")
6 g.edge("Y", "Z")
7 g.node("Y", "Y", style="filled")
8
9
10 g.edge("causal knowledge", "solve problems")
11 g.edge("solve problems", "job promotion")
12
13 g
```



- In this stylized example of a **directed path**, we are postulating that knowing causal inference is the only way to solve business problems
 - This is obviously not true, but it is a useful example for our purposes

Directed Graphical Models

- This model highlights the following statistical process:
 - Knowing causal inference gives you the ability to solve business problems
 - Solving business problems makes you more likely to get a job promotion
- Notice that this does not imply that causal knowledge is independent of job promotion
 - That is, if we know the value of job promotion, we can still learn something about causal knowledge
 - If we observe that a promotion happens in this model, this tells us that it is more likely that the person knows causal inference

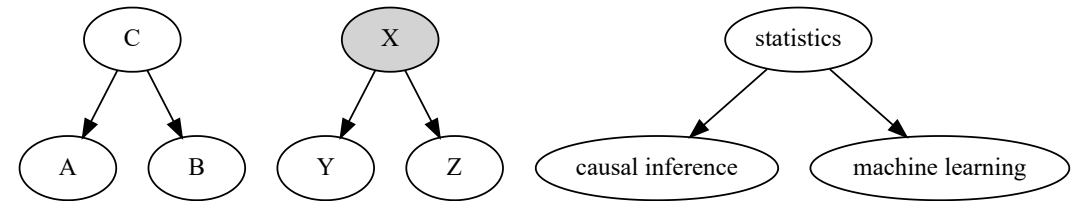
Directed Graphical Models

- Now let's condition on the intermediate variable Y
 - This is the variable that represents the ability to solve business problems
 - In the graph, we have colored this variable grey to indicate that we are conditioning on it
- Conditioning on Y means that we are assuming that we know whether or not the person solved a business problem
- In this case, conditioning on Y breaks the dependence relationship between X and Z
 - That is, X and Z are now independent, given Y
 - Mathematically, $A \perp C$, but $X \not\perp Z|Y$

Directed Graphical Models

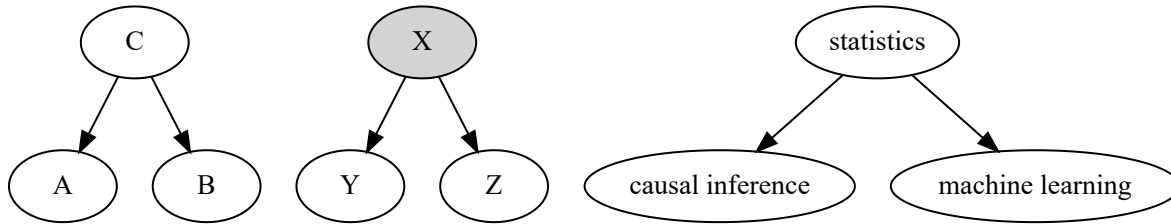
Now let's look at another common structure:

```
1 g = gr.Digraph()
2 g.edge("C", "A")
3 g.edge("C", "B")
4
5 g.edge("X", "Y")
6 g.edge("X", "Z")
7 g.node("X", "X", style="filled")
8
9 g.edge("statistics", "causal inference")
10 g.edge("statistics", "machine learning")
11
12 g
```



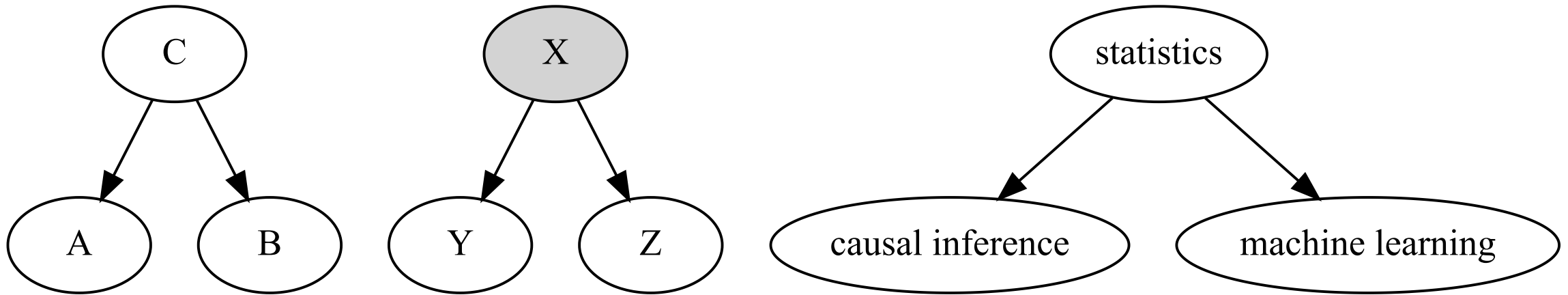
- In this model, we are postulating that statistics is a prerequisite for both causal inference and machine learning.

Directed Graphical Models



- This model highlights the following statistical process:
 - Knowing statistics gives you the ability to do causal inference
 - Knowing statistics gives you the ability to do machine learning
- When we don't condition for the root node, C , there is still a dependence relationship between A and B
 - That is, if we know that an individual has causal knowledge, this tells us that they are more likely to know statistics, and thus to also know machine learning

Directed Graphical Models

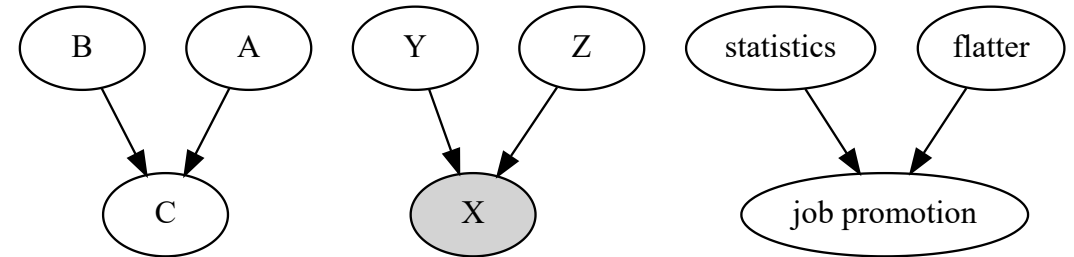


- By conditioning on X , we break the dependence relationship between Y and Z
 - That is, $A \not\perp B$, but $Y \perp Z|X$
- We would call this a **fork** structure

Directed Graphical Models

Finally, let's look at a third common structure, called a **collider**:

```
1 g = gr.Digraph()
2 g.edge("B", "C")
3 g.edge("A", "C")
4
5 g.edge("Y", "X")
6 g.edge("Z", "X")
7 g.node("X", "X", style="filled")
8
9 g.edge("statistics", "job promotion")
10 g.edge("flatter", "job promotion")
11
12 g
```



- In this model, we are postulating that statistics and flattery are both determinants of getting a job promotion.

Building a Causal Graph

Building a Causal Graph

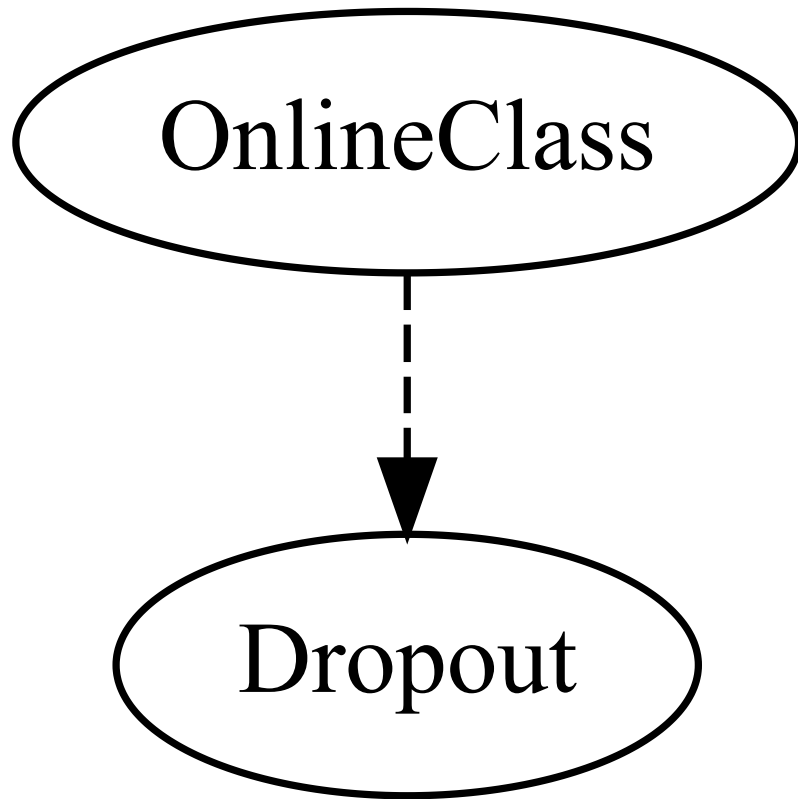
- Now that we have seen some examples of causal graphs, we should think about how to draw a causal graph for a problem in practice.
- Drawing a causal graph is a way of encoding the assumptions that you are making about the data generating process.
- The first step is to identify the variables that are relevant to the problem at hand.
 - Do your research!

Building a Causal Graph

- In reality, there are many variables that are relevant to a problem.
- Real data-generating processes are therefore extremely large and complex.
- Think about what the data generating process might look like for something similar the online class example we have been using
 - (if we didn't have a randomized experiment)

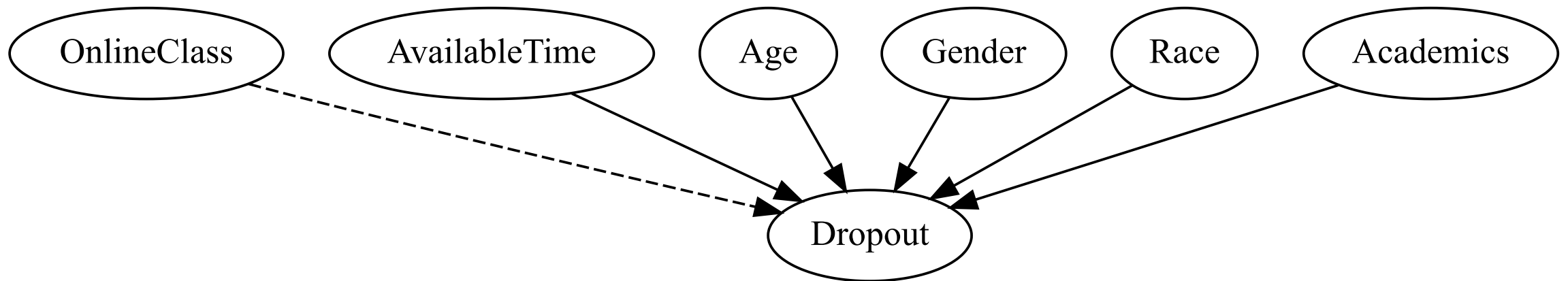
Building a Causal Graph

- Let's say we were interested in the effect of online classes on college dropout rates. The treatment effect we want to identify is:



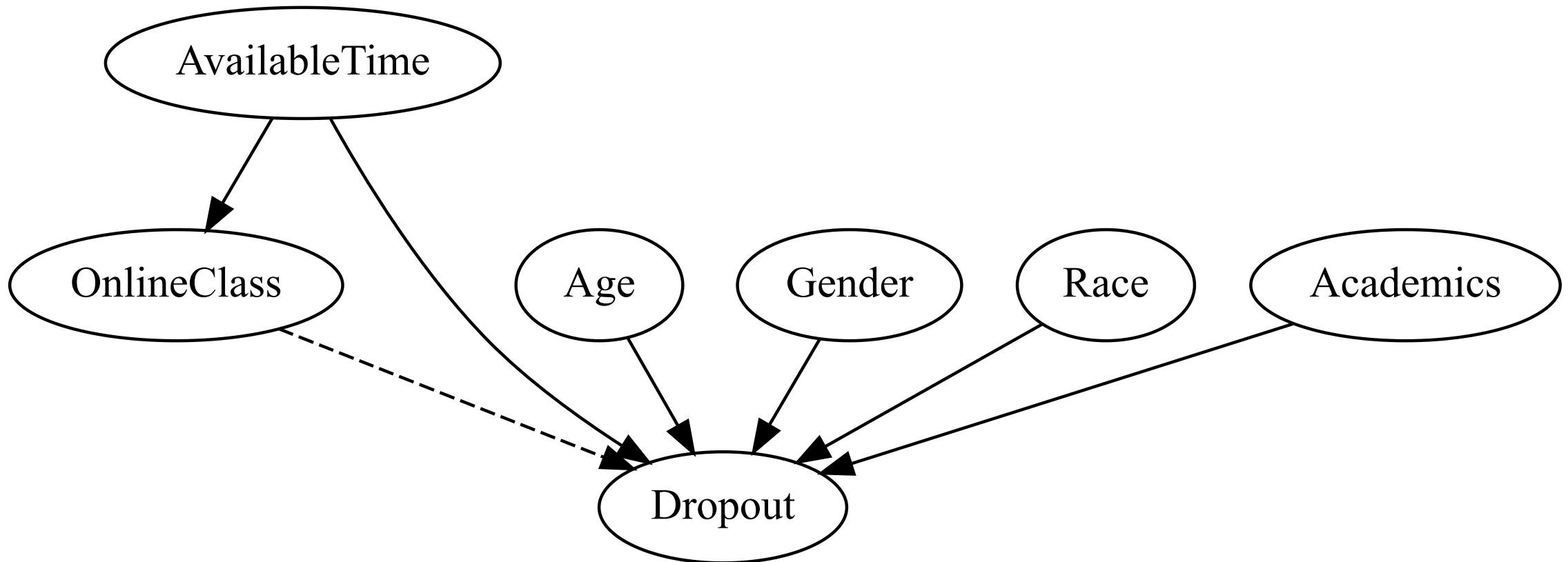
Building a Causal Graph

- In addition to **OnlineClass**, there are **many** variables that might cause a student to drop out of college.



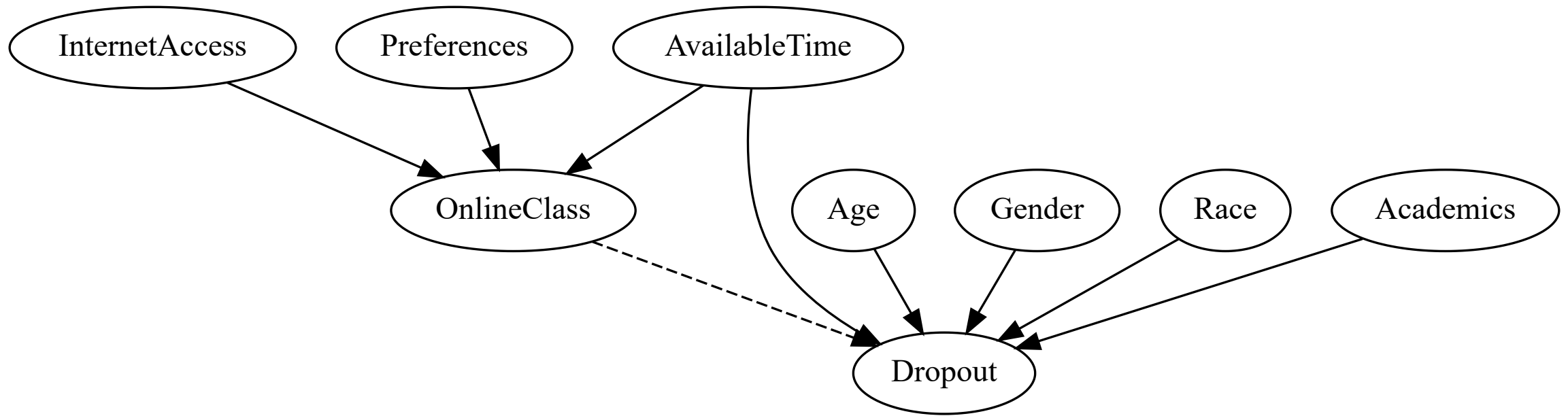
Building a Causal Graph

- Some of these variables might **also** determine whether or not a student takes an online class.



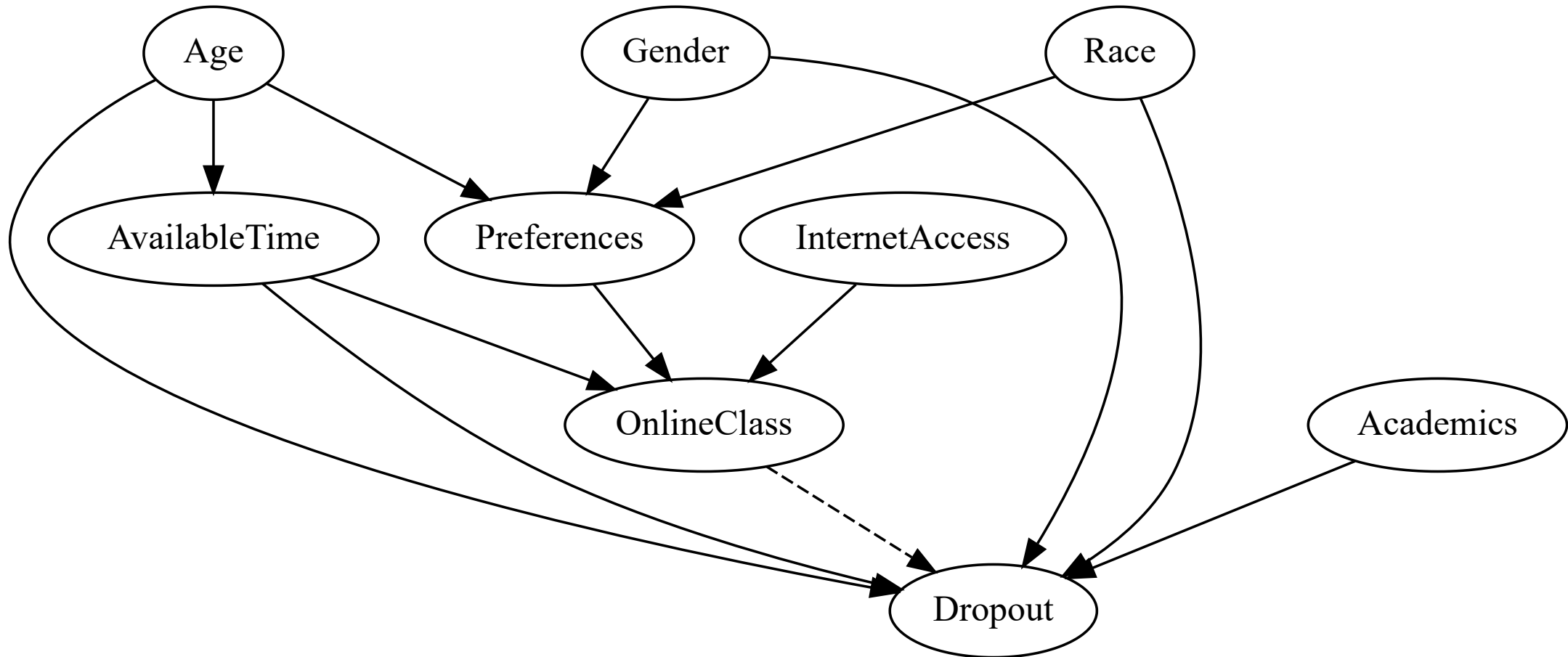
Building a Causal Graph

- Then there are some other variables that affect just **OnlineClass** and not **Dropout**



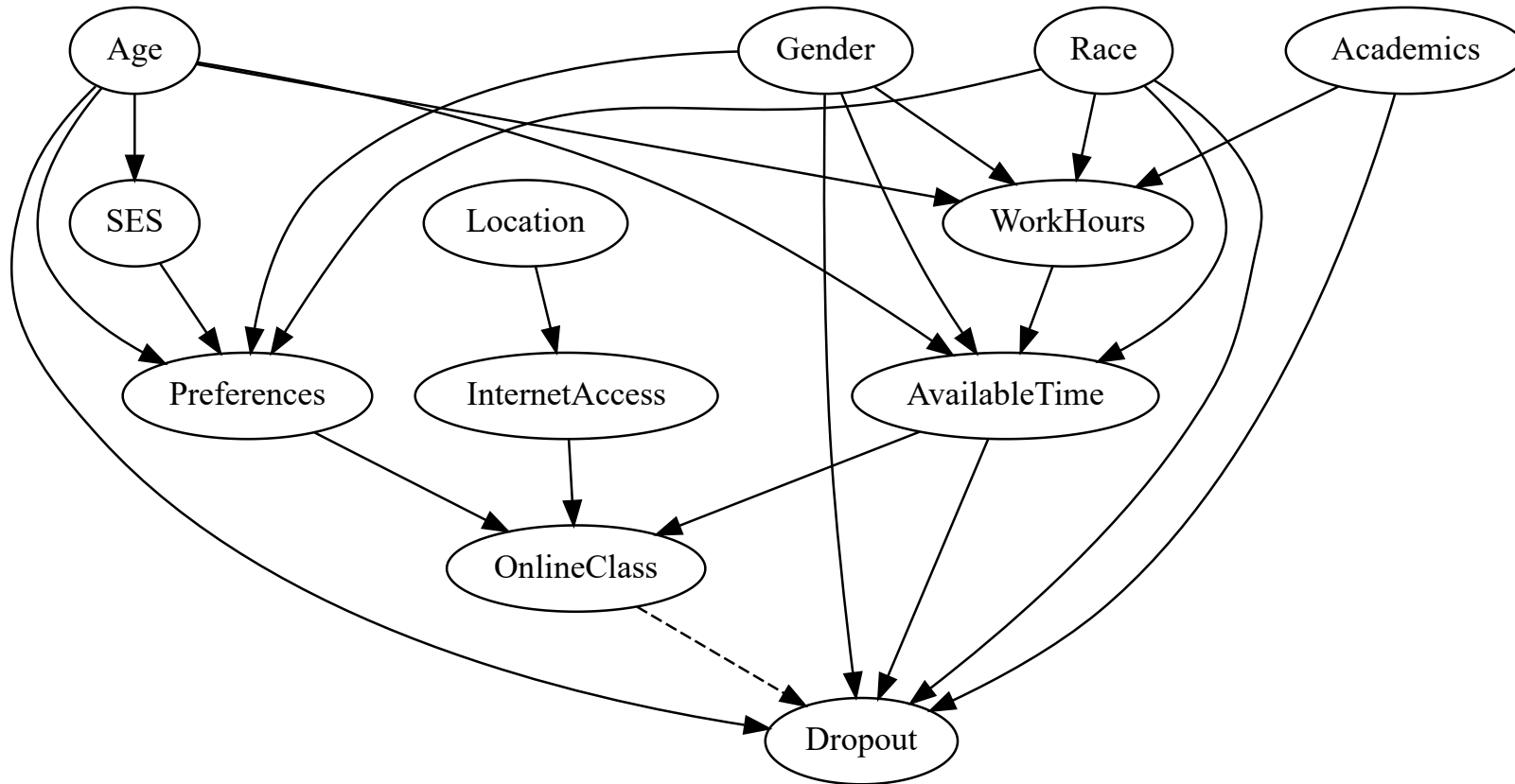
Building a Causal Graph

- There are relationships between some of these variables as well



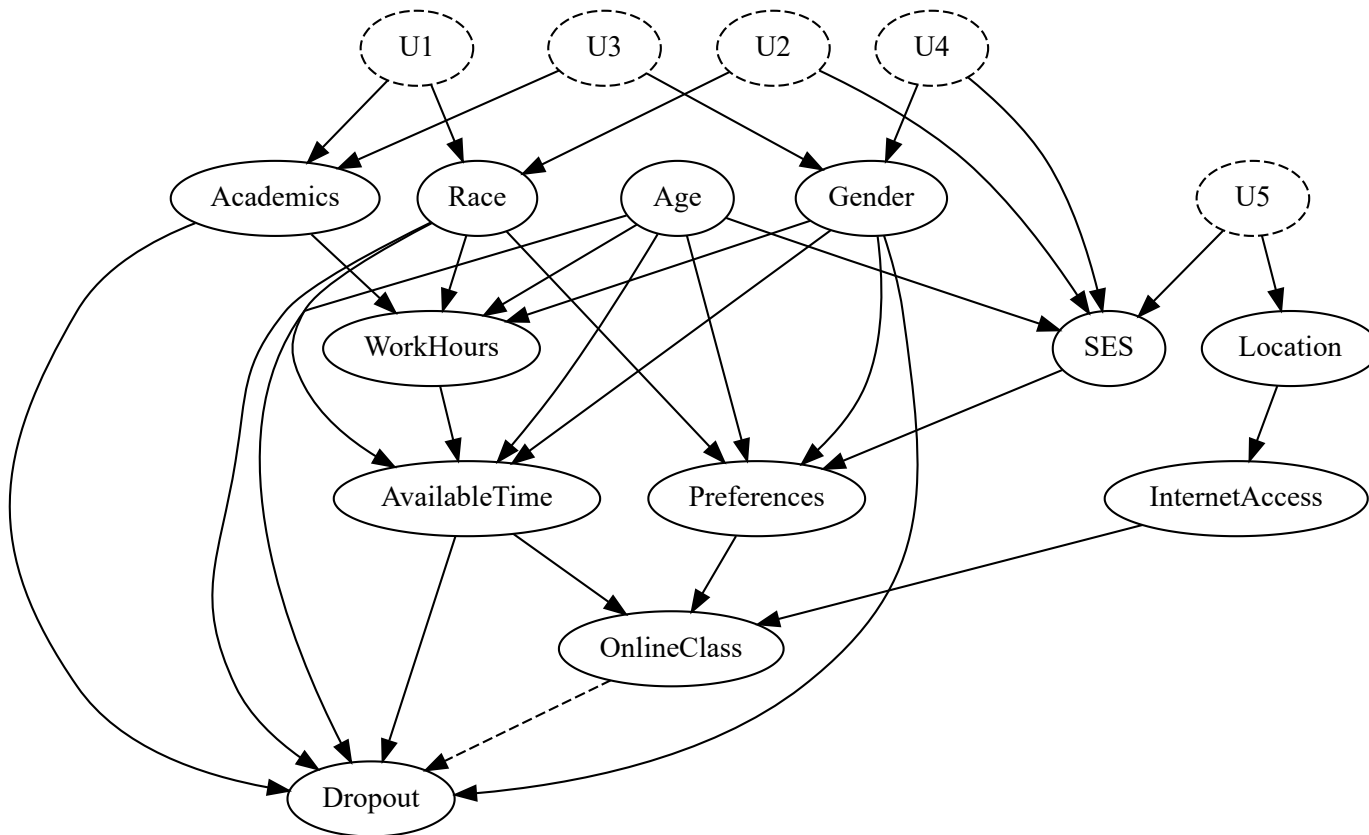
Building a Causal Graph

- Even if there is not a direct causal relationship between two variables, there might be an indirect relationship through a third variable

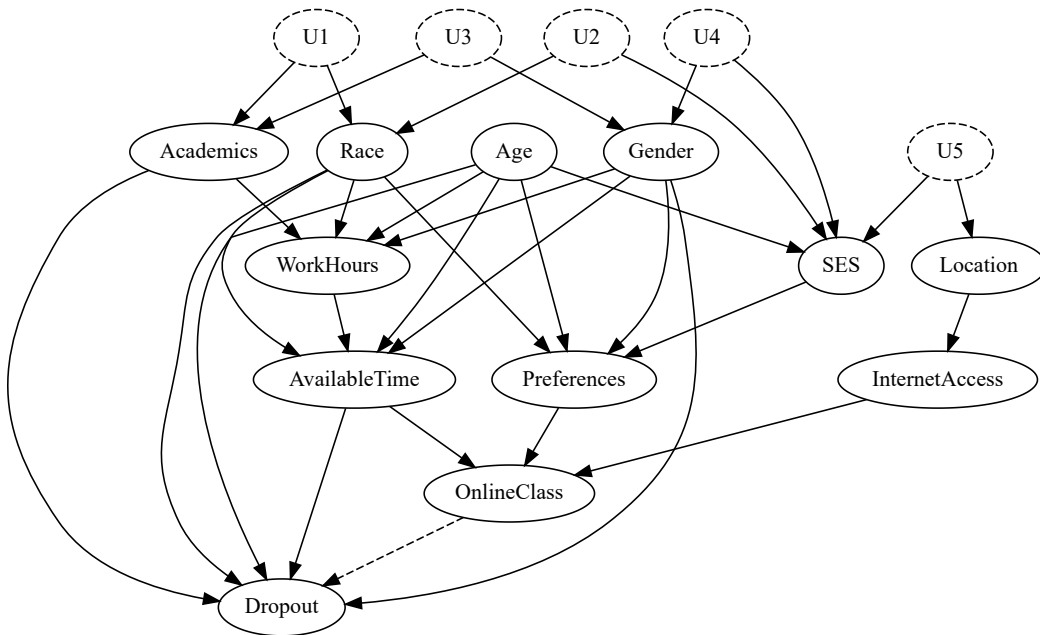


Building a Causal Graph

- And then there are some variables that we know are correlated, but due to some other combination of unknown factors

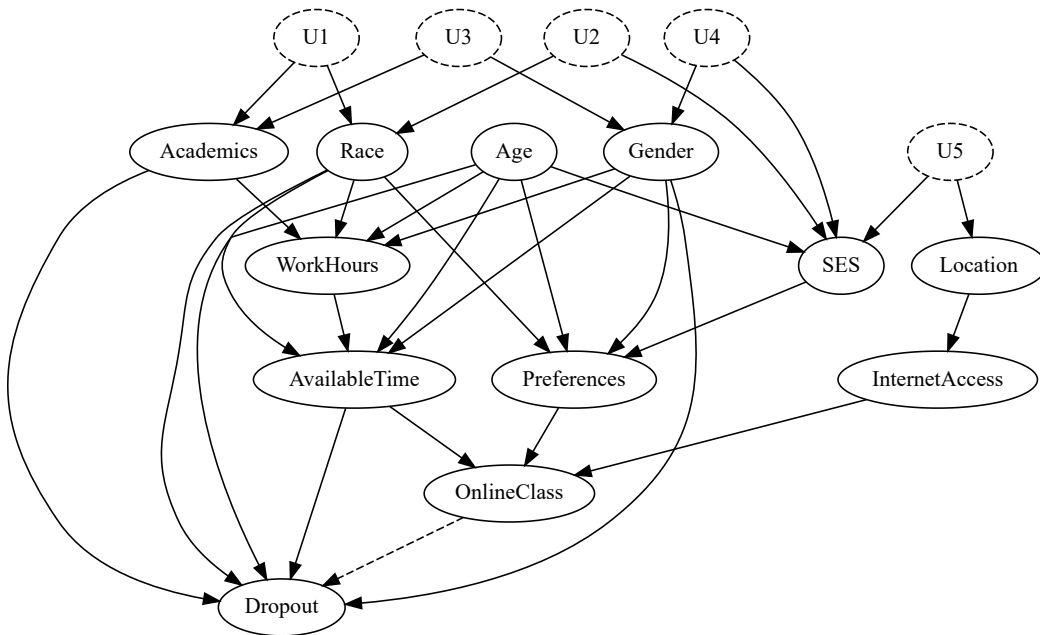


Building a Causal Graph



- As we can see, this graph got very complex, very quickly.
- It is clearly not capturing all of the relevant variables
- What else might be missing?

Building a Causal Graph

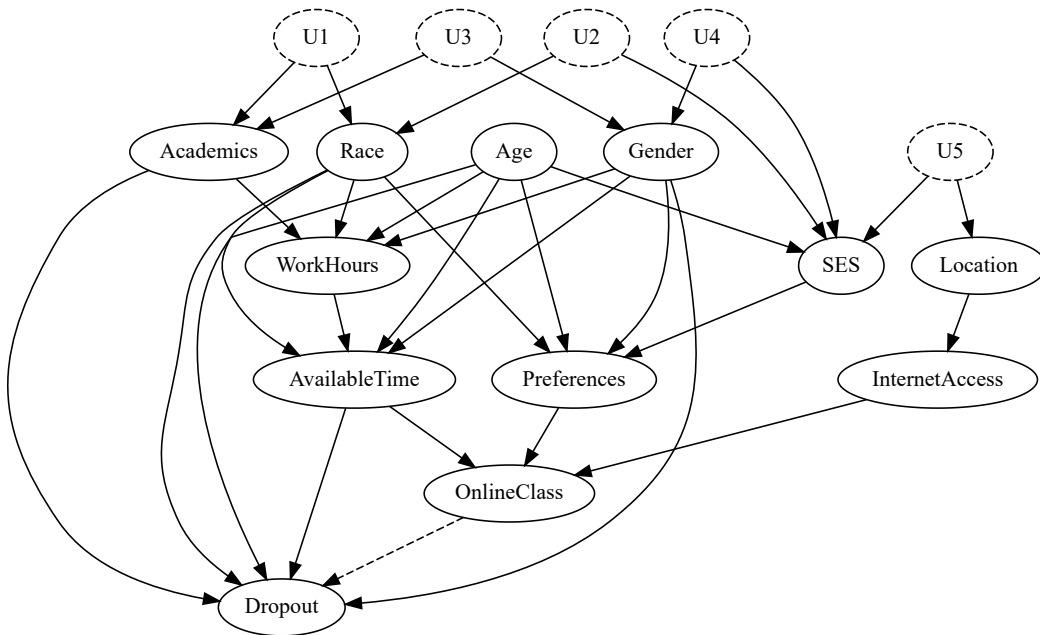


- As we can see, this graph got very complex, very quickly.
- It is clearly not capturing all of the relevant variables
- What else might be missing?
 - CommunityCollege vs. University
 - *Income*

Simplifying the Graph

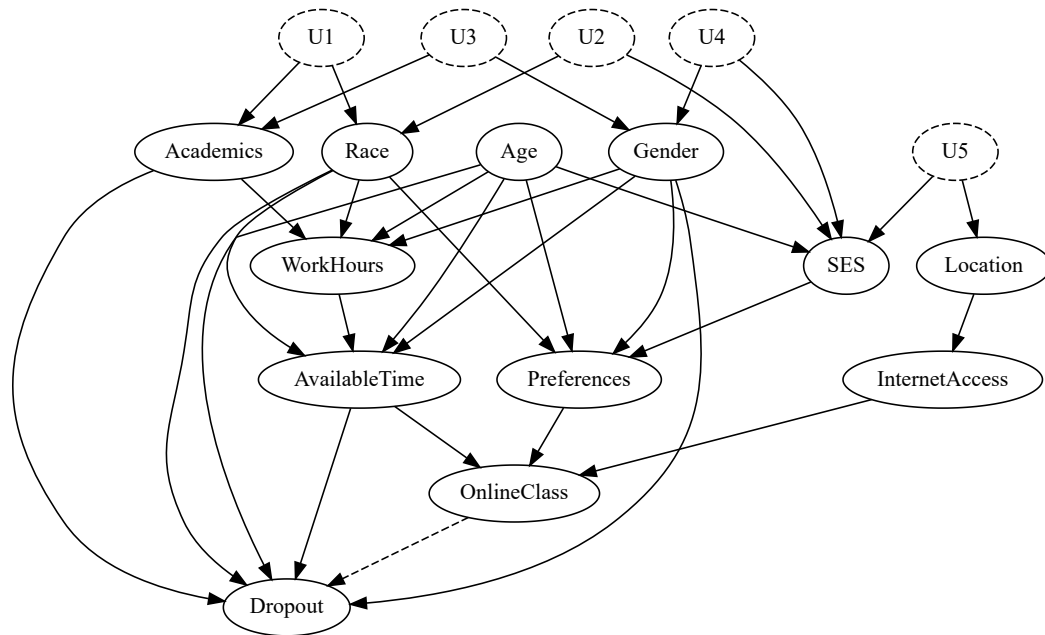
- In practice, we will not be able to draw a graph that captures everything. How can we choose which relationships to include, and which to ignore?
1. **Unimportance** - If the arrows coming in and out of a variable all represent small or negligible effects, we can ignore them.
 2. **Redundancy** - If there are variables on the diagram that occupy the same space (i.e. they both have incoming and outgoing links from the same variables) then we can probably combine them into a single variable.
 3. **Mediators** - If a variable is **only** included as a way to connect two other variables, we can probably remove it.
 4. **Irrelevance** - If a variable is important to the DGP, but isn't part of a **dependence path** between the treatment and the outcome, we can ignore it.

Building a Causal Graph



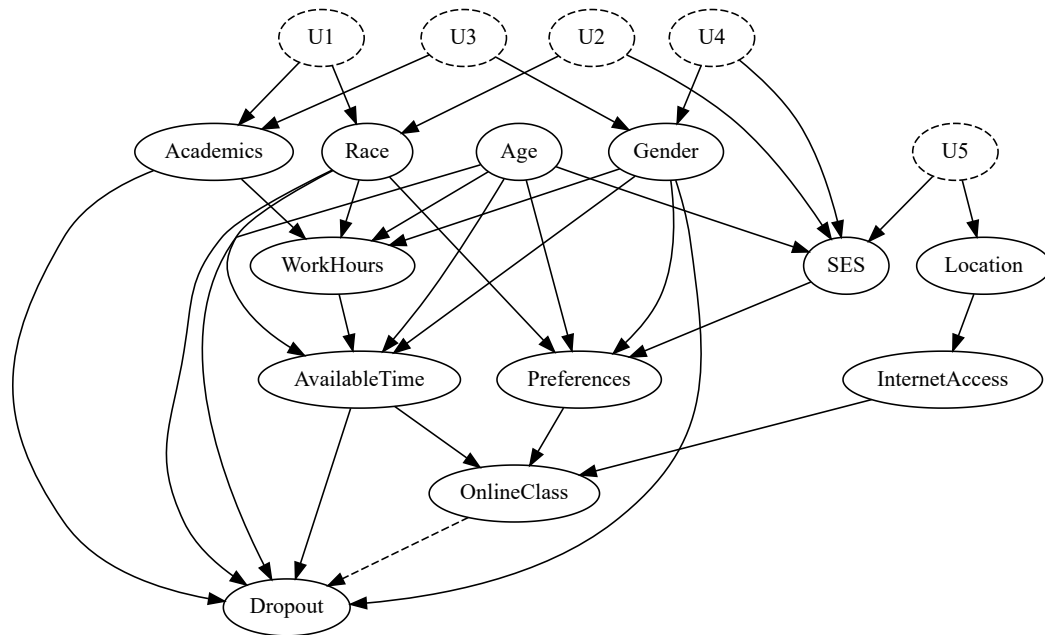
- **Unimportance** was already necessarily applied in the creation of the graph.
- We can also apply **redundancy** by combining **Race** and **Gender** into a single variable, **Demographics**

Building a Causal Graph



- There are also quite a few **Mediators**
 - The most prevalent is **Preferences**, (we can just have **Demographics**, etc. directly affect **OnlineClass**).
 - We can throw out **InternetAccess** without losing anything.
- There is one more. Can you find it?

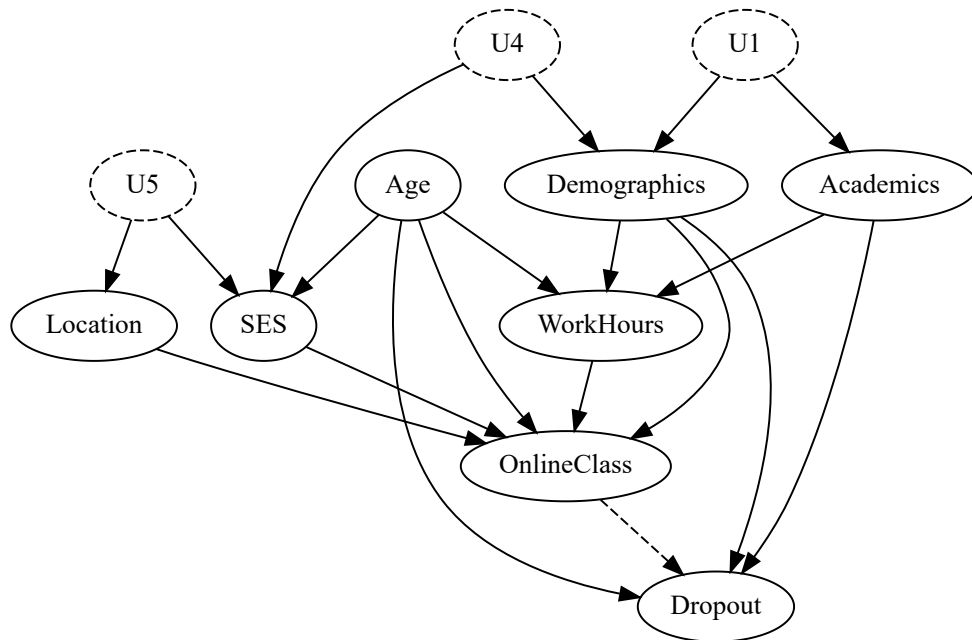
Building a Causal Graph



- Notice that all of the variables that cause **AvailableTime** are also causes of **WorkHours**.
- We can therefore throw out **AvailableTime**, and just have **WorkHours** directly affect **OnlineClass**, rather than through **AvailableTime**.
- **AvailableTime** is also a mediator.

Building a Causal Graph

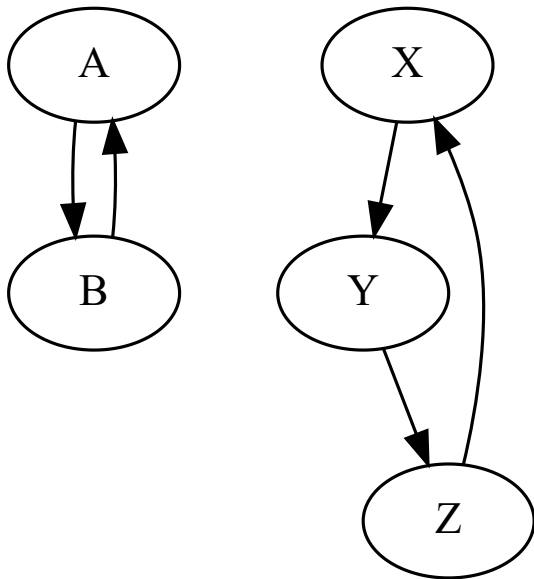
- We are left with a much simpler (although still messy) model, that still captures most of the relevant relationships.
- Even though the simplified model is nicer to work with, these rules are just heuristics and shouldn't be applied blindly.



Aside: Cycles

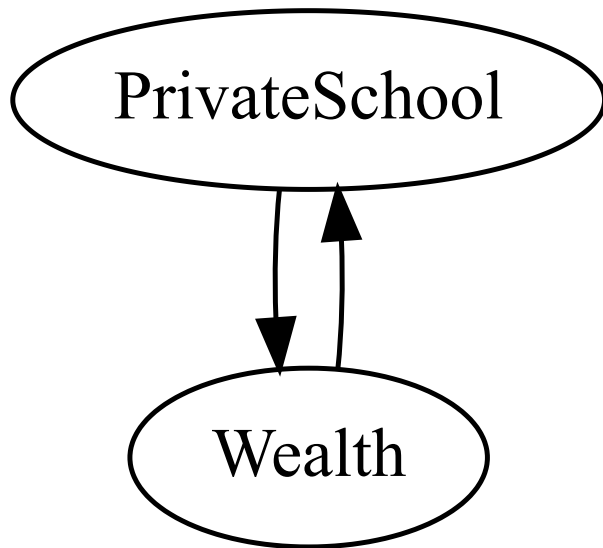
Cycles

- There is one thing that a causal diagram cannot have: a **cycle**.
- A cycle is a directed path that starts and ends at the same node.
 - This would mean that a variable causes itself, which is impossible.
- The simplest cycles look like this:



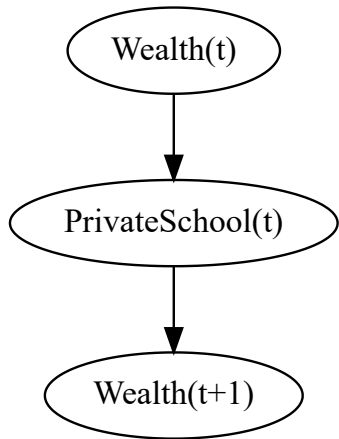
Cycles

- Why can't we have cycles? Surely there are feedback loops in the real world.



- If we have a cycle, then our causal problem is ill-specified.
 - We can't identify the effect of **PrivateSchool** on **Wealth**, because **Wealth** also causes **PrivateSchool**... or does it?

Cycles



- We can solve this problem by introducing the **time** dimension.
- By creating a new variable to represent the lagged value, we can break the cycle.
 - This also loosely corresponds with the physicists view of causality, where the arrow of time is fundamental.
 - Similar to the time-series concept of **Granger Causality**.

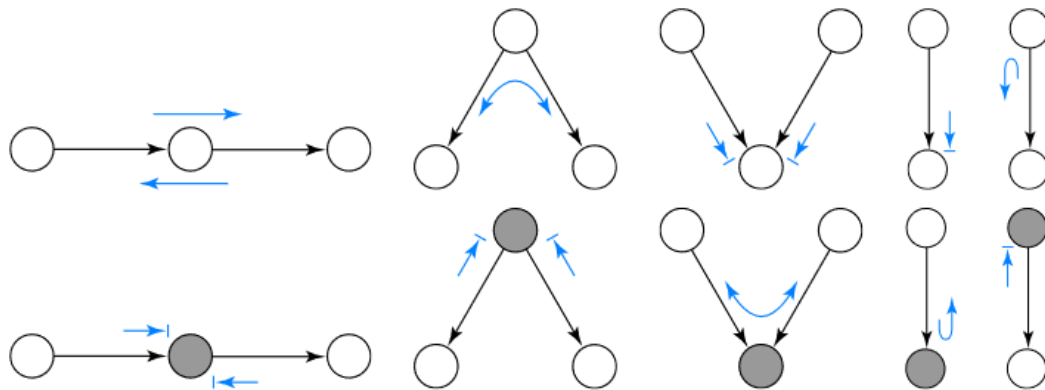
Dependence Flows

Dependence Flows

- In order to determine whether or not we can identify a treatment effect, we need to understand how dependence **flows** through a graphical model.
- We have seen that conditioning on a node can either make or break the dependence relationship between two other nodes
- To identify the treatment effect, we want the *link between the treatment and the outcome to be unblocked*.
- However, we also need to make sure that there is *no other dependence path* between the treatment and the outcome that is unblocked.

The Rules of Bayes-Ball

- We can think about the flow of dependence as a game of “Bayes-ball”
- The rules of Bayes-ball are reasonably simple. A dependence path is blocked if and only if:
 1. It contains a *non-collider* that is conditioned on
 2. It contains a *collider* that is not conditioned on, and neither are *any of its descendants*

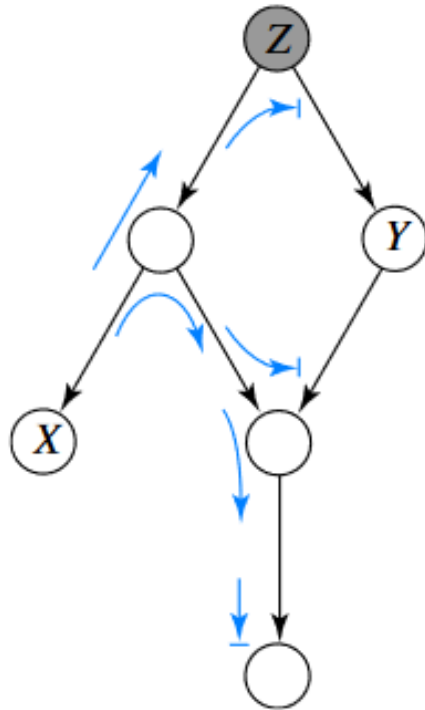


Directed Graphical Models

Turning back to our **collider** example:

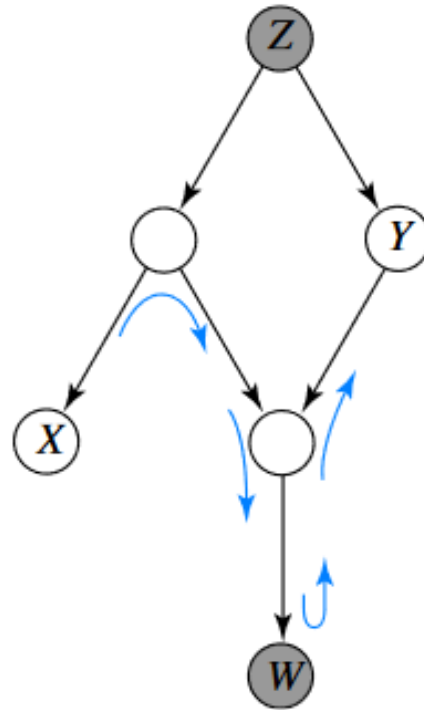
- Notice that when we do not condition on C , A and B are independent.
- However, somewhat unintuitively, when we condition on X , Y and Z become dependent.
 - This is because conditioning on X opens the flow of dependence from Y to Z .
 - In any case that is *not* a collider, conditioning on a node *blocks* the flow of dependence.

Two Games of Bayes Ball



no active paths

$$X \perp\!\!\!\perp Y \mid Z$$



one active path

$$X \not\perp\!\!\!\perp Y \mid \{W, Z\}$$

Viualizing Bias

Bias and Causality

- In a causal inference framework, we can use graphical models to determine whether or not we can identify a treatment effect, and which covariates we need to condition on.
- Typically, drawing out a graphical model is not necessary, but it can be a useful exercise to help you think through the problem.
 - The links you draw represent the assumptions you are making about the data generating process

Bias and Causality

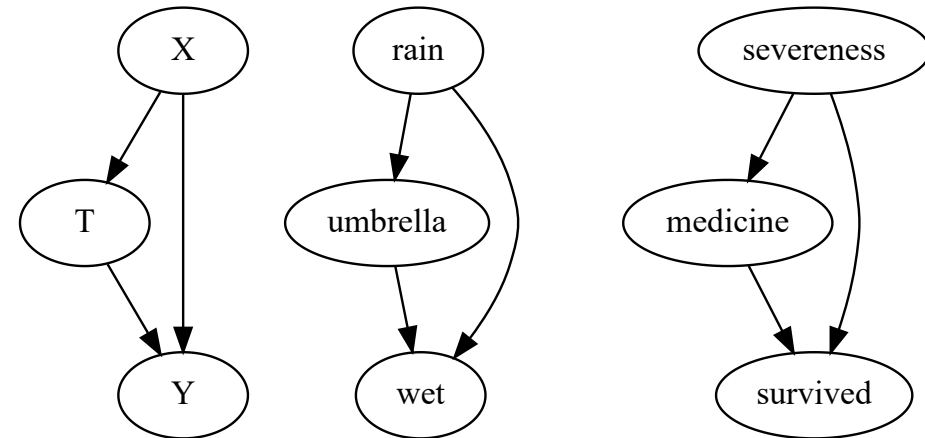
- There are two major types of bias that we need to worry about in causal inference:
 - **Confounding**: When there is an unobserved variable that is a common cause of both the treatment and the outcome
 - **Selection**: When there is an unobserved variable that is a common cause of both the treatment and the selection into the sample
- Both of these types of bias can be represented using a graphical model

Confounding

Confounding

- Let's look at an example of confounding:

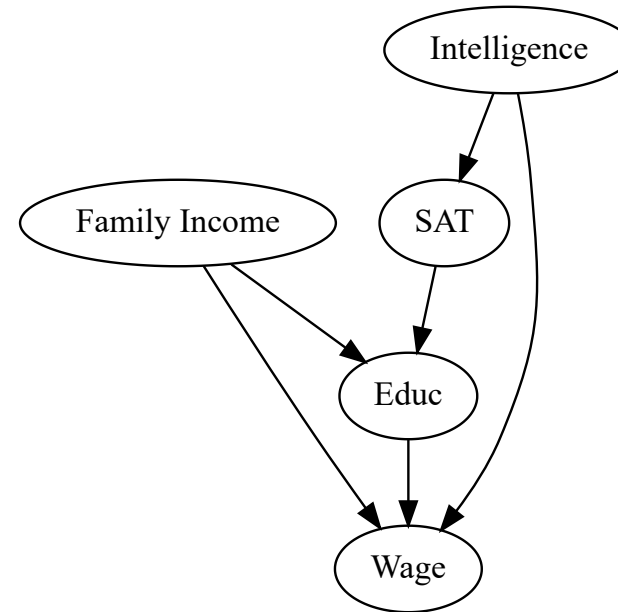
```
1 g = gr.Digraph()
2 g.edge("X", "T")
3 g.edge("X", "Y")
4 g.edge("T", "Y")
5
6 g.edge("rain", "umbrella")
7 g.edge("rain", "wet")
8 g.edge("umbrella", "wet")
9
10 g.edge("severeness", "medicine")
11 g.edge("severeness", "survived")
12 g.edge("medicine", "survived")
13 g
```



- To control for confounding, we need to condition on all of the common causes of the treatment and the outcome.

Confounding

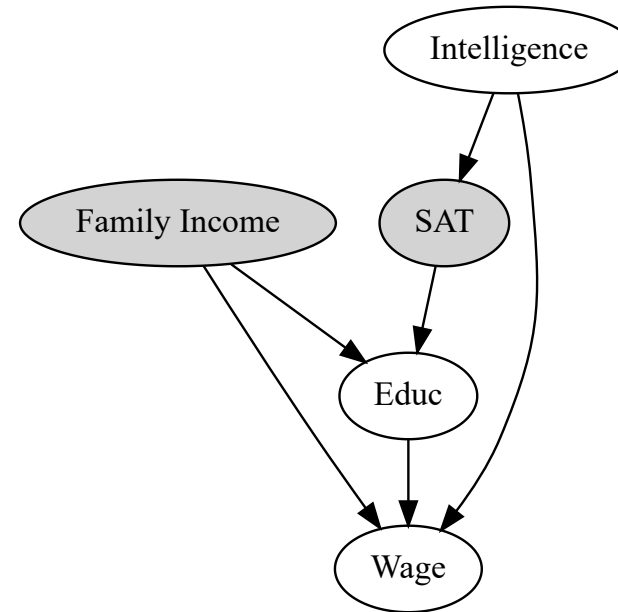
```
1 g = gr.Digraph()
2
3 g.node("Family Income")
4 g.edge("Family Income", "Educ")
5 g.edge("Educ", "Wage")
6
7 g.node("SAT")
8 g.edge("SAT", "Educ")
9
10 g.node("Intelligence")
11 g.edge("Intelligence", "SAT")
12 g.edge("Intelligence", "Wage")
13
14 g
```



- Often, there are confounding variables that we cannot observe
 - For example, we cannot observe intelligence, but it is a common cause of both education (the treatment) and wages

Confounding

```
1 g = gr.Digraph()
2
3 g.node("Family Income", style="filled")
4 g.edge("Family Income", "Educ")
5 g.edge("Educ", "Wage")
6
7 g.node("SAT", style="filled")
8 g.edge("SAT", "Educ")
9
10 g.node("Intelligence", style="filled")
11 g.edge("Intelligence", "SAT")
12 g.edge("Intelligence", "Wage")
13
14 g
```



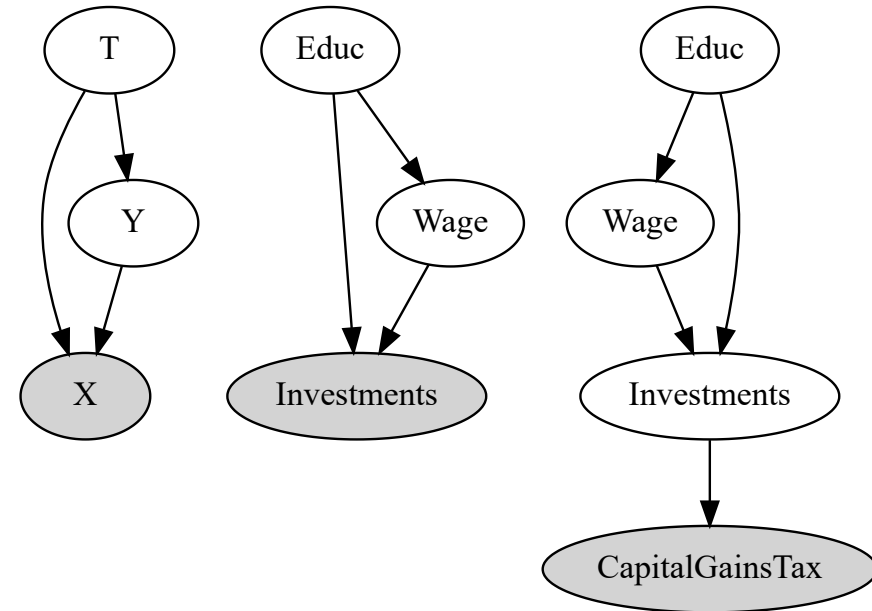
- Often, there are confounding variables that we cannot observe
 - For example, we cannot observe intelligence, but it is a common cause of both education (the treatment) and wages
 - But we can use SAT as a **surrogate** or **proxy** for intelligence.

Selection

Selection

- Selection bias often occurs when there is an unobserved variable that is a common cause of both the treatment and the selection into the sample, in other words, by conditioning on a variable that you shouldn't.

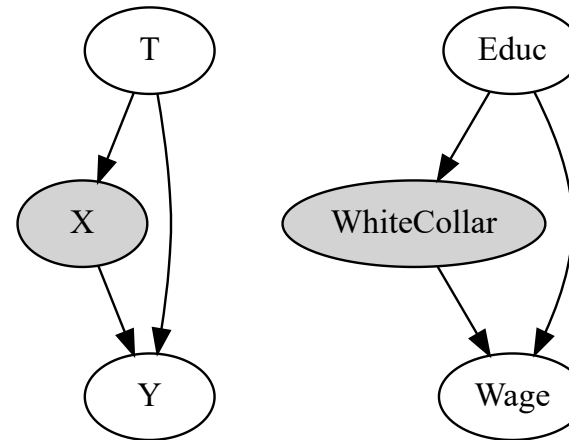
```
1 g = gr.Digraph()
2 g.node("X", style="filled")
3 g.edge("T", "X")
4 g.edge("T", "Y")
5 g.edge("Y", "X")
6 g.node("Investments", "Investments", style="filled")
7 g.edge("Educ", "Investments")
8 g.edge("Educ", "Wage")
9 g.edge("Wage", "Investments")
10 g.node("Educ2", "Educ")
11 g.node("Wage2", "Wage")
12 g.node("Investments2", "Investments")
13 g.edge("Educ2", "Wage2")
14 g.edge("Wage2", "Investments2")
15 g.edge("Educ2", "Investments2")
16 g.node("CapitalGainsTax", "CapitalGainsTax", style="filled")
```



Selection

- Selection bias can also occur when controlling for a **mediator** between the treatment and the outcome

```
1 g = gr.Digraph()
2
3 g = gr.Digraph()
4 g.edge("T", "X")
5 g.edge("T", "Y")
6 g.edge("X", "Y")
7 g.node("X", "X", style="filled")
8
9 g.edge('Educ', 'WhiteCollar')
10 g.edge('Educ', 'Wage')
11 g.edge('WhiteCollar', 'Wage')
12 g.node('WhiteCollar', style="filled")
13
14 g
```



Choosing Covariates

Choosing Covariates

- With these types of bias in mind, we can think about how to choose which covariates to condition on.
- Notice that some controls will reduce bias, as in the case of confounding,
- But *not all controls are good!* Some controls will actually create bias, as in the case of selection.
- This means that we don't want to just “throw the kitchen sink” at the problem, and include every variable we can think of.

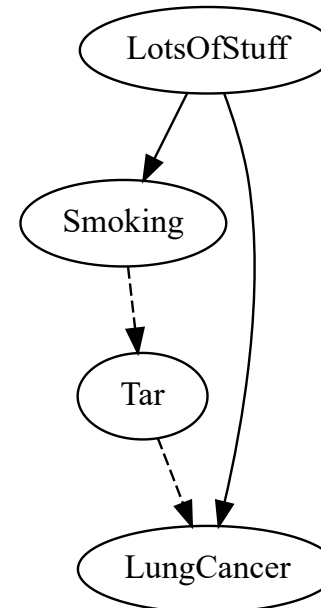
Choosing Covariates

- We want to choose covariates that will close any unblocked secondary dependence paths between the treatment and the outcome, but not open any new ones.
- To do this, we can use the “front door criterion” and the “back door criterion”

The Front Door Criterion

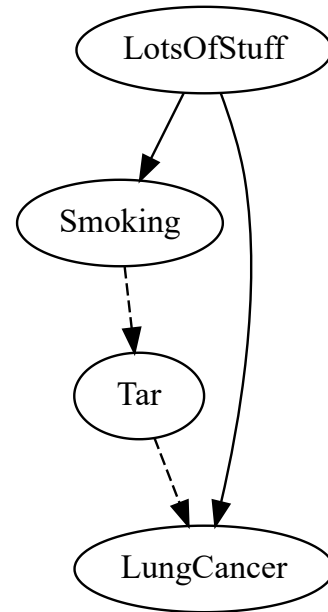
- The **front door criterion** is one way to isolate the effect of a treatment on an outcome, when there is a confounder and a mediator between the treatment and the outcome.

```
1 g = gr.Digraph()
2 g.edge("LotsOfStuff", "Smoking")
3 g.edge("LotsOfStuff", "LungCancer")
4 g.edge("Smoking", "Tar", style="dashed")
5 g.edge("Tar", "LungCancer", style="dashed")
6
7 g
```



The Front Door Criterion

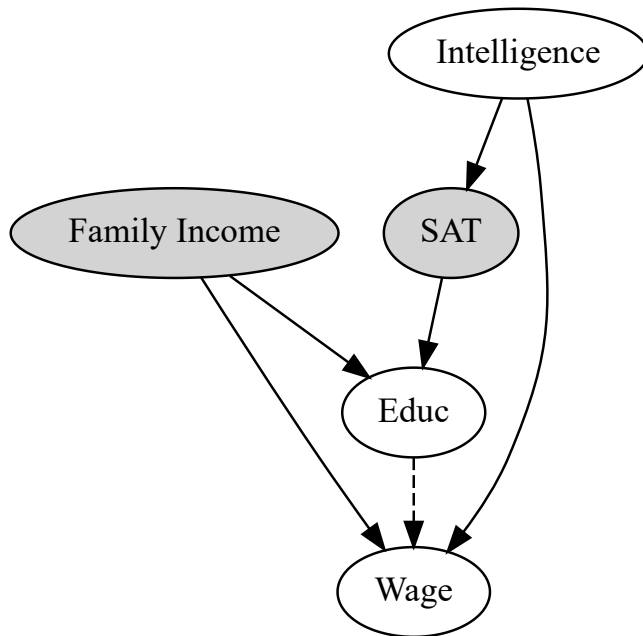
```
1 g = gr.Digraph()
2 g.edge("LotsOfStuff",
3 g.edge("LotsOfStuff",
4 g.edge("Smoking", "Tar
5 g.edge("Tar", "LungCa
6
7 g
```



- In this example, we want to know the effect of smoking on lung cancer, but we also know that there are lots of variables (like stress), that cause both the treatment and the outcome.
- However, stress does not cause tar, so we can condition on tar.
 - This works by first measuring the effect of tar on lung cancer, and *then* the effect of smoking on tar.

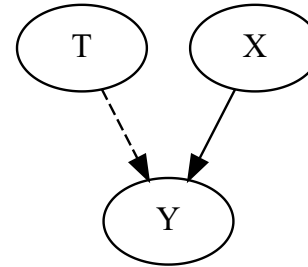
The Back Door Criterion

- It's pretty rare that we'll actually be able to use the front door criterion, because we usually don't have a mediator that we can condition on.
- Instead, we can close all of the “back door paths” from treatment to outcome. We have already seen an example of this.



Choosing Covariates: Two Examples

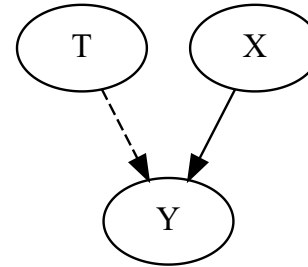
```
1 g = gr.Digraph()
2 g.edge("T", "Y", style="dashed")
3 g.edge("X", "Y")
4 g
```



- Do we need to condition on X ?

Choosing Covariates: Two Examples

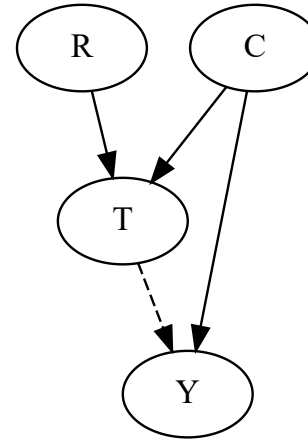
```
1 g = gr.Digraph()
2 g.edge("T", "Y", style="dashed")
3 g.edge("X", "Y")
4 g
```



- Do we need to condition on \mathbf{X} ?
 - No, not necessarily. There is no unblocked dependence path between \mathbf{T} and \mathbf{Y} that goes through \mathbf{X} .
 - However, conditioning on \mathbf{X} will reduce the variance of our estimate!
 - If we didn't condition on \mathbf{X} , it would become part of our estimation error. But since $\mathbf{X} \perp \mathbf{T}$, it won't bias our estimate.

Choosing Covariates: Two Examples

```
1 g = gr.Digraph()
2 g.edge("R", "T")
3 g.edge("C", "T")
4 g.edge("C", "Y")
5 g.edge("T", "Y", style="dashed")
6 g
```



- Suppose we don't have any data on C . Can we identify the treatment effect?
 - Yes! If we look at the effect of R on Y , we can see that it is unblocked.
 - Furthermore, since R only affects Y through T , the effect of R on Y is the same as the effect of T on Y .
 - This is called an **instrumental variable**.

Factorizing the Joint Distribution

Factorizing the Joint Distribution

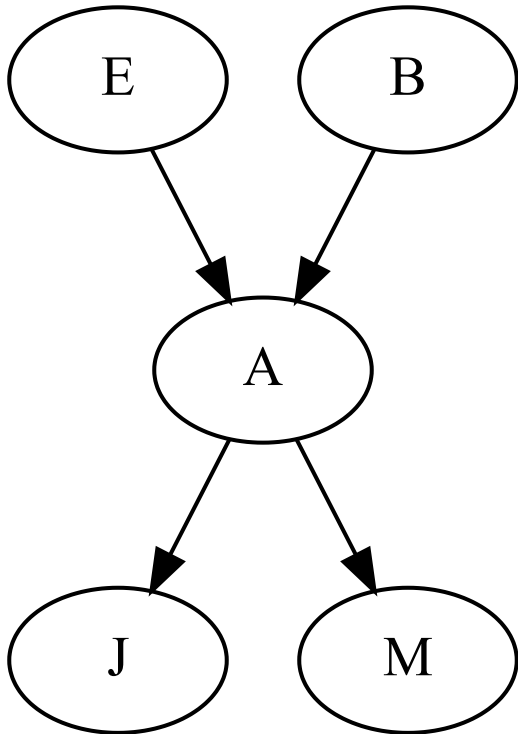
- We have seen how these graphical models can be used to determine whether or not we can identify a treatment effect.
- However, we can also use them as a computational tool, to help us find the simplest representation of the joint distribution of the data.
- This is useful because it allows us to find the simplest model that is consistent with our assumptions about conditional independence.

Factorizing the Joint Distribution

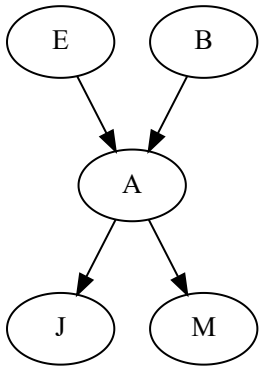
- To understand why this is useful, we'll follow this example (from [Mark Paskin](#)).
- Suppose we have a DGP with five variables:
 1. $E \in \{true, false\}$ - Has an earthquake happened? *Earthquakes are unlikely*
 2. $B \in \{true, false\}$ - Has a burglary happened? *Burglaries are unlikely, but more likely than earthquakes*
 3. $A \in \{true, false\}$ - Is the alarm going off? *The alarm is triggered by both earthquakes and burglaries*
 4. $J \in \{true, false\}$ - Is John calling? *John calls when he hears the alarm, but he often misses it*
 5. $M \in \{true, false\}$ - Is Mary calling? *Mary calls when she hears the alarm, but she also calls to chat*

Factorizing the Joint Distribution

- The goal is to compute $P(B|J = \textit{true})$ from the joint distribution $P(E, B, A, J, M)$. We'll start by drawing the graphical model, to understand the conditional independence relationships.



Factorizing the Joint Distribution



- In order to represent the full joint distribution, we could use the chain rule of probabilities:

$$P(E, B, A, J, M) = P(E)P(B|E)P(A|E, B)P(J|E, B, A)P(M|E, B, A, J)$$

- Q: How many probabilities would we need to compute to represent the joint distribution this way?

Factorizing the Joint Distribution



- In order to represent the full joint distribution, we could use the chain rule of probabilities:

$$\underbrace{P(E, B, A, J, M)}_{31} = \underbrace{P(E)}_1 \underbrace{P(B|E)}_2 \underbrace{P(A|E, B)}_4 \underbrace{P(J|E, B, A)}_8 \underbrace{P(M|E, B, A, J)}_{16}$$

- Q: How many probabilities would we need to store to represent the joint distribution this way?
 - A: There are 2^5 possible outcomes. We need 31 probabilities. (why not 32?)

Factorizing the Joint Distribution

88

- However, we can use the conditional independence relationships to simplify this representation.
 - For example, we know that $P(B|E) = P(B)$, because B and E are independent.
 - We also know that $P(A|E, B) = P(A|B)$, because A is independent of E , given B .
- This means that we can simplify the joint distribution to:

$$\underbrace{P(E, B, A, J, M)}_{10} = \underbrace{P(E)}_1 \underbrace{P(B)}_1 \underbrace{P(A|E, B)}_4 \underbrace{P(J|A)}_2 \underbrace{P(M|A)}_2$$

Factorizing the Joint Distribution

- Beyond causal inference, graphical models are a useful tool for computing all kinds of conditional probabilities.
- This is useful in many inference problems, and in structural econometrics
 - The **likelihood function** is the conditional probability of the data, given the parameter values
 - In Bayesian inference the **posterior** is the conditional probability of the parameters, given the data (and our priors)

Variable Elimination

- In order to simplify a conditional distribution, we can use **variable elimination**.
- For example, let's say we want to compute the probability of a burglary, given that John calls.

$$\begin{aligned} p_{B|J}(b, \text{true}) &\propto \sum_e \sum_a \sum_m p_{EBAJM}(e, b, a, \text{true}, m) \\ &= \sum_e \sum_a \sum_m p_E(e) \cdot p_B(b) \cdot p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a) \cdot p_{M|A}(m, a) \end{aligned}$$

- Then, we can reduce the computational complexity by eliminating variables one at a time, exploiting the distributive property of multiplication

$$\rightarrow xy + xz = x(y + z).$$

Variable Elimination

- Variable elimination works like this:
 - Repeat the following steps:
 1. choose a variable to eliminate
 2. push in its sum as far as possible
 3. compute the sum, resulting in a new factor

Variable Elimination

$$\begin{aligned} & \sum_e \sum_a \sum_m p_E(e) \cdot p_B(b) \cdot p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a) \cdot p_{M|A}(m, a) \\ &= \sum_e \sum_a p_E(e) \cdot p_B(b) \cdot p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a) \cdot \sum_m p_{M|A}(m, a) \\ &= \sum_e \sum_a p_E(e) \cdot p_B(b) \cdot p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a) \cdot \psi_A(a) \\ &= \sum_e p_E(e) \cdot p_B(b) \cdot \sum_a p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a) \cdot \psi_A(a) \\ &= \sum_e p_E(e) \cdot p_B(b) \cdot \psi_{EB}(e, b) \\ &= p_B(b) \cdot \sum_e p_E(e) \cdot \psi_{EB}(e, b) \\ &= p_B(b) \cdot \psi_B(b) \end{aligned}$$

Variable Elimination

- That's a lot of math! But let's focus on the first step:

$$\begin{aligned}
 & p_{B|J}(b, \text{true}) \propto \\
 & \underbrace{\sum_e \sum_a \sum_m}_{2^3=8 \text{ iterations}} \underbrace{p_E(e) \cdot p_B(b) \cdot p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a) \cdot p_{M|A}(m, a)}_{4 \text{ multiplications}}
 \end{aligned}$$

8*4=32 multiplications + 7 additions = 39 total operations

Variable Elimination

- That's a lot of math! But let's focus on the first step:

$$\begin{aligned}
 & p_{B|J}(b, \text{true}) \propto \\
 & \underbrace{\sum_e \sum_a \sum_m}_{2^3=8 \text{ iterations}} \underbrace{p_E(e) \cdot p_B(b) \cdot p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a) \cdot p_{M|A}(m, a)}_{4 \text{ multiplications}} \\
 & \underbrace{8 \cdot 4 = 32 \text{ multiplications} + 7 \text{ additions} = 39 \text{ total operations}}
 \end{aligned}$$

$$\begin{aligned}
 & = \underbrace{\sum_e \sum_a}_{2^2=4 \text{ iterations}} \underbrace{p_E(e) \cdot p_B(b) \cdot p_{A|EB}(a, e, b) \cdot p_{J|A}(\text{true}, a)}_{4 \text{ multiplications}} \cdot \underbrace{\sum_m p_{M|A}(m, a)}_{1 \text{ addition}} \\
 & \underbrace{4 \cdot (4 \text{ multiplications} + 1 \text{ addition}) + 3 \text{ additions} = 23 \text{ total operations}}
 \end{aligned}$$

Variable Elimination

- Variable elimination is an algorithm that exploits our conditional independence assumptions to reduce the computational complexity of conditional probabilities.
- In this case, we were able to reduce the number of operations from 39 to 23 operations in just one step, each further step will continue to reduce the complexity.
- This is a very simple example, because we only have 5 variables. In practice, we might have hundreds or thousands of variables, and the computational complexity can become very large.
- Systematic patterns of conditional independence can be exploited to reduce the complexity of inference problems in some large models.

Credits

This lecture draws heavily from [Causal Inference for the Brave and True: Chapter 04 - Graphical Causal Models](#) by Matheus Facure.

There is also material from [A Short Course on Graphical Models Chapter 2: Structured Representations](#) by Mark Paskin.

As well as [The Effect: Chapter 7 - Drawing Graphical Diagrams](#) by Nick Huntington-Klein.