

Chapter 7: Confounding

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In Chapter 3, we introduced exchangeability as a key identifiability condition. In Chapter 6, we learned to represent causal relationships using DAGs and introduced the backdoor criterion for identifying confounding. This chapter provides a detailed examination of **confounding**—the most common threat to validity in observational studies.

This chapter is based on Hernán and Robins (2020, chap. 7, pp. 77-92).

1 7.1 The Structure of Confounding (pp. 77-80)

Confounding occurs when a common cause of treatment and outcome creates a non-causal association between them.

Definition 1.1 (Confounding Structure). A variable L is a **confounder** of the effect of A on Y if:

1. L causes A (or shares a common cause with A)
2. L causes Y (or shares a common cause with Y)
3. L is not affected by A (not a consequence of treatment)

Causal diagram representation:

$L \rightarrow A \rightarrow Y$
 $L \rightarrow Y$

The path $A \leftarrow L \rightarrow Y$ is a **backdoor path** that creates non-causal association.

Why confounding creates bias:

Without confounding:

$$E[Y|A = 1] - E[Y|A = 0] = E[Y^{a=1}] - E[Y^{a=0}]$$

(association equals causation)

With confounding:

$$E[Y|A = 1] - E[Y|A = 0] \neq E[Y^{a=1}] - E[Y^{a=0}]$$

(association does not equal causation)

The observed association includes both:

- The causal effect of A on Y
- The confounding bias due to the common cause L

1.1 Common Confounding Scenarios

Example 1: Healthy worker bias

- Healthier individuals are more likely to be employed (employed \rightarrow more likely to be exposed at work)
- Healthier individuals have better outcomes
- Comparing employed vs. unemployed introduces confounding by health status

Example 2: Confounding by indication

- Sicker patients receive more aggressive treatment
- Sicker patients have worse outcomes
- Treatment appears harmful when in fact it may be beneficial

Confounding can go in either direction:

- **Positive confounding:** Makes treatment appear more beneficial (or more harmful) than it truly is
- **Negative confounding:** Makes treatment appear less beneficial (or less harmful) than it truly is
- **Zero confounding:** The magnitude depends on the strength and direction of the $L \rightarrow A$ and $L \rightarrow Y$ relationships

2 7.2 Confounding and Exchangeability (pp. 80-82)

Confounding is equivalent to lack of (conditional) exchangeability.

2.1 No Confounding = Exchangeability

No confounding means:

$$Y^a \perp\!\!\!\perp A \quad \text{for all } a$$

This is **marginal exchangeability**: the counterfactual outcomes are independent of treatment.

Confounding means exchangeability does not hold:

$$Y^a \not\perp\!\!\!\perp A$$

The treated and untreated differ with respect to their potential outcomes.

Example 2.1 (Confounding and Exchangeability). Suppose exercise (A) affects heart disease (Y), and both are affected by age (L):

Without confounding:

- Young and old people equally likely to exercise
- $E[Y^{a=1}|A = 1] = E[Y^{a=1}|A = 0]$ (exchangeable)

With confounding:

- Younger people more likely to exercise
- Younger people have lower baseline risk
- $E[Y^{a=1}|A = 1] \neq E[Y^{a=1}|A = 0]$ (not exchangeable)
- Those who exercise would have had better outcomes even without exercising

2.2 Conditional Exchangeability

Even when marginal exchangeability fails, we may achieve **conditional exchangeability** by adjusting for confounders:

$$Y^a \perp\!\!\!\perp A \mid L \quad \text{for all } a$$

Within levels of L , the treated and untreated are exchangeable.

Key insight: Confounding can be eliminated by conditioning on (adjusting for) the confounders.

Requirements: 1. We must **identify** all confounders based on subject-matter knowledge 2. We must **measure** them accurately 3. We must **adjust** for them appropriately in analysis

If these requirements are met, we can estimate causal effects from observational data.

3 7.3 Confounding and the Backdoor Criterion (pp. 82-85)

The backdoor criterion (Chapter 6) provides a graphical method for identifying confounding.

3.1 Backdoor Paths and Confounding

A **backdoor path** from A to Y :

- Starts with an arrow into A (i.e., $\cdot \rightarrow A$)
- Connects A to Y through any sequence of arrows

Confounding exists if backdoor paths are open (unblocked).

Example 3.1 (Identifying Confounders with the Backdoor Criterion). **Diagram 1:**

$L \rightarrow A \rightarrow Y$
 $L \rightarrow Y$

Backdoor path: $A \leftarrow L \rightarrow Y$ **Confounders:** L **Solution:** Adjust for L

Diagram 2:

$U \rightarrow L \rightarrow A \rightarrow Y$
 $L \rightarrow Y$

Backdoor paths: $A \leftarrow L \rightarrow Y$, $A \leftarrow L \leftarrow U \rightarrow Y$ (if U causes Y) **Confounders:** L (and U if it affects Y) **Solution:** Adjust for L (and U if measured)

Diagram 3:

$A \rightarrow M \rightarrow Y$
 $L \rightarrow A$
 $L \rightarrow Y$

Backdoor path: $A \leftarrow L \rightarrow Y$ **Confounders:** L **Do NOT adjust for M :** M is a mediator (on the causal path), not a confounder

Common mistakes:

1. **Adjusting for mediators:** Variables on the causal path from A to Y should NOT be adjusted for, as this blocks the causal effect we're trying to estimate
2. **Adjusting for colliders:** Variables caused by both A and Y should NOT be adjusted for, as this induces bias
3. **Failing to adjust for all confounders:** If even one confounder is unmeasured or unadjusted, bias remains
4. **Overadjustment:** Including unnecessary variables (especially descendants of treatment) can introduce bias

4 7.4 Confounding and Confounders (pp. 85-87)

The traditional definition of “confounder” in epidemiology differs slightly from the causal DAG perspective.

4.1 Traditional Confounder Definition

Traditionally, a variable L is considered a confounder if: 1. L is associated with treatment A 2. L is associated with outcome Y (among the untreated) 3. L is not affected by treatment A

4.2 DAG-Based Definition

From the DAG perspective, L is a confounder if:

- L opens a backdoor path from A to Y

Differences:

The traditional definition is based on **associations** (statistical relationships). The DAG definition is based on **causal structure** (graphical relationships).

Why this matters:

1. **A variable can be associated with both A and Y without being a confounder**
 - Example: A collider caused by both A and Y
 - Adjusting for it would induce bias, not remove it
2. **A variable can be a confounder without being associated with both A and Y in the data**
 - Example: A confounder whose effects cancel out, leaving no association
 - Failing to adjust for it would leave bias

Best practice: Use DAGs to identify confounders based on causal structure, not associations alone.

5 7.5 Single-World Intervention Graphs (pp. 87-89)

Single-World Intervention Graphs (SWIGs) are an extension of DAGs that explicitly represent interventions and counterfactual outcomes.

5.1 SWIGs vs. DAGs

- **Standard DAGs:** Represent relationships among observed variables
- **SWIGs:** Represent relationships among counterfactual variables under specified interventions

SWIG notation:

- Y_a : Counterfactual outcome under intervention $do(A = a)$
- A_a : Treatment value set to a by intervention
- Edges represent causal effects in the counterfactual world where $A = a$

Example SWIG: For the causal effect of A on Y with confounder L :

$L \rightarrow A_a \rightarrow Y_a$
 $L \rightarrow Y_a$

SWIGs make explicit:

- Which variables are set by intervention
- Which variables remain as observed
- Which counterfactual outcome we're interested in

Advantages:

- Clearer representation of counterfactuals
- Explicit about the intervention
- Useful for complex scenarios (time-varying treatments, mediation)

Disadvantages:

- More complex notation
- Require more assumptions to be specified
- Not yet as widely used as standard DAGs

For most purposes in this book, standard DAGs suffice. SWIGs are mentioned for completeness and for readers interested in advanced topics.

6 7.6 Confounding Adjustment (pp. 89-92)

Once confounders are identified, several methods can adjust for them.

6.1 Methods for Confounding Adjustment

1. **Stratification:** Estimate effects within strata of L , then combine (standardization)
2. **Regression adjustment:** Include L as covariates in a regression model
3. **Inverse probability weighting:** Weight by $1/Pr[A|L]$ to create a pseudo-population where A and L are independent (Chapter 12)
4. **Matching:** Match treated and untreated individuals on L

Example 6.1 (Comparing Adjustment Methods). **Data:** Effect of smoking (A) on lung cancer (Y), adjusting for age (L)

Stratification:

- Estimate effect separately for age = 40, 50, 60, 70
- Combine using weighted average

Regression:

```
glm(Y ~ A + L, family = binomial())
```

IP weighting (Chapter 12):

```
weight <- 1 / predict(glm(A ~ L, family = binomial()), type = "response")
glm(Y ~ A, weights = weight, family = binomial())
```

Matching:

- For each smoker, find non-smoker of same age
- Compare outcomes

Choosing an adjustment method:

Stratification:

- Transparent, easy to understand
- Allows checking for effect modification
- Limited to discrete confounders
- Requires large sample sizes for fine strata

Regression:

- Handles continuous confounders
- Efficient (uses all data)
- Relies on model assumptions
- Can obscure effect modification

IP weighting:

- Flexible, can handle complex confounding
- Estimates marginal (population-average) effects
- Can be unstable with extreme weights
- More complex to implement

Matching:

- Intuitive, creates comparable groups
- Discards unmatched data
- Largely superseded by other methods

Modern recommendation: Use IP weighting or doubly robust methods (combine regression and weighting) for flexibility and robustness.

7 Summary

This chapter provided a detailed examination of **confounding**.

Key concepts:

1. **Confounding structure:** Common causes of treatment and outcome create backdoor paths
2. **Exchangeability:** Confounding = lack of exchangeability; conditional exchangeability can be achieved by adjusting for confounders
3. **Backdoor criterion:** Provides a graphical method to identify which variables to adjust for
4. **DAG vs. traditional definitions:** DAG-based confounding identification is preferred over association-based criteria
5. **Adjustment methods:** Stratification, regression, IP weighting, and matching can all adjust for confounding
6. **Critical assumptions:**
 - All confounders must be identified (no unmeasured confounding)
 - All confounders must be measured accurately
 - Adjustment must be done correctly

Practical guidelines for dealing with confounding:

1. **Draw a DAG** based on subject-matter knowledge before analyzing data
2. **Identify confounders** using the backdoor criterion
3. **Measure confounders** as accurately as possible
4. **Choose an adjustment method** appropriate for your data and confounders
5. **Check assumptions:**
 - Positivity: Do all (A, L) combinations occur?
 - Model fit: Are regression model assumptions met?
 - Balance: After adjustment, are confounders balanced?
6. **Conduct sensitivity analyses:** How robust are findings to unmeasured confounding?
7. **Be transparent:** Report the DAG, adjustment set, and method clearly

Limitations:

Even with perfect adjustment for measured confounders, bias can remain if:

- Important confounders are unmeasured (Chapter 19 covers sensitivity analysis)
- Confounders are measured with error (Chapter 9)
- Adjustment methods are applied incorrectly
- Positivity is violated

Confounding control is necessary but not sufficient for valid causal inference.

Looking ahead:

- **Chapter 8:** Selection bias
- **Chapter 9:** Measurement bias
- **Chapters 12-15:** Advanced methods for confounding adjustment

8 References

Hernán, Miguel A, and James M Robins. 2020. *Causal Inference: What If*. Boca Raton: Chapman & Hall/CRC. <https://miguelhernan.org/whatifbook>.