Partial Identification

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Roadmap for this Lecture

The Limits of Causal Inference

- \triangleright (Y_0, Y_1) never observed for the same person; can't learn their joint distribution.
- ▶ Quantities like $Var(Y_1 Y_0)$ or $\mathbb{P}(Y_1 Y_0 > 0)$ are **not identifiable.**

Partial Identification

 \blacktriangleright Even if we can't pin θ down exactly, we may be able to **rule out** many values.

Outline

- 1. Simplest example of partial identification.
- 2. Bounds on ATE while allowing for selection bias.
- 3. Bound the distribution of treatment effects.

Simple Example: Reverse Regression Bounds

Population Linear Regression

- lacktriangleq lpha and eta are intercept and slope from population linear regression of Y on X
- ▶ Thus we can write $Y = \alpha + \beta X + U$ where we define

$$\beta \equiv \frac{\mathsf{Cov}(X,Y)}{\mathsf{Var}(X)}, \quad \alpha \equiv \mathbb{E}[Y] - \beta \mathbb{E}[X], \quad U \equiv Y - \alpha - \beta X$$

b By **construction** we have $\mathbb{E}(XU) = \mathbb{E}(U) = 0$.

Point Identification

- ▶ If we could observe the whole population from which our sample was drawn, could we uniquely determine the parameters of interest?
- ▶ Suppose we observe the joint distribution of (X, Y)
- \blacktriangleright This is enough information to calculate (α, β) explicitly: they are **point identified**.

Classical Measurement Error

- ▶ Suppose we observe (Y, \widetilde{X}) rather than (Y, X), where $\widetilde{X} = X + W$
- ▶ W is classical measurement error: $Cov(W, X) = Cov(W, U) = \mathbb{E}(W) = 0$
- ▶ Are α and β still point identified?

The Good News

$$\mathbb{E}(\widetilde{X}) = \mathbb{E}(X+W) = \mathbb{E}(X)$$

$$\mathsf{Cov}(\widetilde{X},Y) = \mathsf{Cov}(X+W,Y) = \mathsf{Cov}(X,Y) + \mathsf{Cov}(W,Y)$$

$$= \mathsf{Cov}(X,Y) + \mathsf{Cov}(W,\alpha + \beta X + U)$$

$$= \mathsf{Cov}(X,Y) + \mathsf{Cov}(W,U) + \beta \mathsf{Cov}(W,X)$$

$$= \mathsf{Cov}(X,Y)$$

Are α and β still point identified?

The Bad News

• Because Var(W) is not point identified, neither are α and β .

$$\mathsf{Var}(\widetilde{X}) = \mathsf{Var}(X + W) = \mathsf{Var}(X) + \mathsf{Var}(W) \ge \mathsf{Var}(X)$$

$$\beta \equiv \frac{\mathsf{Cov}(X,Y)}{\mathsf{Var}(X)} = \frac{\mathsf{Cov}(X,Y)}{\mathsf{Var}(\widetilde{X}) - \mathsf{Var}(W)}, \quad \alpha \equiv \mathbb{E}[Y] - \beta \mathbb{E}[X] = \mathbb{E}[Y] - \beta \mathbb{E}[\widetilde{X}].$$

Partial Identification

 \blacktriangleright We can still **bound** β and hence α : the so-called **reverse regression bounds**

A Lower Bound for β

▶ Since $Cov(X, Y) = Cov(\widetilde{X}, Y)$,

$$\frac{\mathsf{Cov}(\widetilde{X},Y)}{\mathsf{Var}(\widetilde{X})} = \frac{\mathsf{Cov}(X,Y)}{\mathsf{Var}(X) + \mathsf{Var}(W)} = \frac{\mathsf{Cov}(X,Y)/\mathsf{Var}(X)}{1 + \mathsf{Var}(W)/\mathsf{Var}(X)} = \frac{\beta}{1 + \mathsf{Var}(W)/\mathsf{Var}(X)}.$$

ightharpoonup Since Var(W)/Var(X) is non-negative, $Cov(\widetilde{X},Y)/Var(\widetilde{X})$ has same sign as β and

$$\left| \frac{\mathsf{Cov}(\widetilde{X}, Y)}{\mathsf{Var}(\widetilde{X})} \right| \leq |\beta|.$$

An Upper Bound for β

ightharpoonup Run the **reverse regression** \widetilde{X} on Y

$$\frac{\mathsf{Cov}(\widetilde{X},Y)}{\mathsf{Var}(Y)} = \frac{\mathsf{Cov}(X,Y)}{\beta^2 \mathsf{Var}(X) + \mathsf{Var}(U)} = \frac{\beta \mathsf{Var}(X)}{\beta^2 \mathsf{Var}(X) + \mathsf{Var}(U)}.$$

► Take the reciprocal:

$$\frac{\mathsf{Var}(Y)}{\mathsf{Cov}(\widetilde{X},Y)} = \beta + \frac{\mathsf{Var}(U)}{\beta \mathsf{Var}(X)} = \beta \left[1 + \frac{\mathsf{Var}(U)}{\beta^2 \mathsf{Var}(X)} \right].$$

▶ Factor in brackets greater than one, so $Var(Y)/Cov(\widetilde{X},Y)$ has same sign as β and

$$\left| \frac{\mathsf{Var}(Y)}{\mathsf{Cov}(\widetilde{X},Y)} \right| \geq |\beta|.$$

Reverse Regression Bounds

Terminology

- A bound is sharp if it cannot be improved, under our assumptions.
- A bound is **tight** if it is short enough to be useful in a practical example.

Assumptions

- $Y = \alpha + \beta X + U$ where $\mathbb{E}(XU) = \mathbb{E}(U) = 0$.
- ightharpoonup Observe (\widetilde{X}, Y)
- $\widetilde{X} = X + W$ with $\mathbb{E}(W) = \text{Cov}(W, X) = \text{Cov}(W, U) = 0$

Sharp Bounds for β

How tight are the reverse regression bounds?

Let r denote the correlation between \widetilde{X} and Y. Then:

$$r^2 \equiv \frac{\mathsf{Cov}(\widetilde{X},Y)^2}{\mathsf{Var}(\widetilde{X})\mathsf{Var}(Y)} = \frac{\mathsf{Cov}(\widetilde{X},Y)}{\mathsf{Var}(\widetilde{X})} \cdot \frac{\mathsf{Cov}(\widetilde{X},Y)}{\mathsf{Var}(Y)}.$$

► Re-arranging, it follows that:

$$r^2 \cdot \frac{\mathsf{Var}(Y)}{\mathsf{Cov}(\widetilde{X},Y)} = \frac{\mathsf{Cov}(\widetilde{X},Y)}{\mathsf{Var}(\widetilde{X})}.$$

▶ All else equal, bounds for β are *tighter* when \widetilde{X} and Y are strongly correlated:

$$\mathsf{Width} = \left| \frac{\mathsf{Var}(Y)}{\mathsf{Cov}(\widetilde{X},Y)} - \frac{\mathsf{Cov}(\widetilde{X},Y)}{\mathsf{Var}(\widetilde{X})} \right| = (1-r^2) \left| \frac{\mathsf{Var}(Y)}{\mathsf{Cov}(\widetilde{X},Y)} \right|.$$

```
library(tidyverse)
library(broom) # for tidy()
set.seed(1066)
n < -5000
X \leftarrow rnorm(n)
U \leftarrow rnorm(n)
W <- rnorm(n)
alpha <- 0.5
beta <- 1
Y \leftarrow alpha + beta * X + U
Xtilde <- X + W
```

```
c(forward = cov(Xtilde, Y) / var(Xtilde),
 truth = beta.
 reverse = var(Y) / cov(Xtilde, Y)) %>% round(2)
## forward truth reverse
     0.51 1.00 1.95
##
# The regression we can't run in practice!
lm(Y \sim X) \%\% tidy()
## # A tibble: 2 x 5
    term estimate std.error statistic p.value
##
                           <dbl>
## <chr>
                  <dbl>
                                   <dbl>
                                             < dbl>
## 1 (Intercept) 0.489 0.0140 34.8 9.56e-238
                  1.02
## 2 X
                          0.0138 73.9 0
```

```
# Reduce the correlation between X and Y, hence Xtilde and Y
Y <- alpha + beta * X + 3 * U
c(forward = cov(Xtilde, Y) / var(Xtilde),
 truth = beta.
 reverse = var(Y) / cov(Xtilde, Y)) %>% round(2)
## forward truth reverse
## 0.52 1.00 9.31
# The regression we can't run in practice!
lm(Y ~ X) %>% tidy()
## # A tibble: 2 \times 5
## term estimate std.error statistic p.value
## <chr> <dbl> <dbl> <dbl> <dbl>
## 1 (Intercept) 0.466 0.0421 11.1 3.95e- 28
```

1.07 0.0414 25.7 7.45e-137

2 X

Review of Potential Outcomes Framework

- ► See https://expl.ai/QHUAVRV and https://expl.ai/DWVNRZU for more details.
- ▶ Binary **Treatment** $D \in \{0,1\}$
- **Description** Outcome Y depends on Potential Outcomes (Y_0, Y_1) via

$$Y = (1 - D)Y_0 + DY_1 = Y_0 + D(Y_1 - Y_0)$$

- \triangleright Only one of (Y_0, Y_1) is observed for any given person at any given time.
- The unobserved potential outcome is a counterfactual, i.e. a what if?
- ▶ Average Treatment Effect: ATE $\equiv \mathbb{E}(Y_1 Y_0)$.
- ▶ Treatment on the Treated: TOT $\equiv \mathbb{E}(Y_1 Y_0|D=1)$.

Example: Y is Wage, D is Attend University

Counterfactuals

- $ightharpoonup D = 1 \implies Y_0$ is the wage you would have earned if you hadn't attended.
- $ightharpoonup D = 0 \implies Y_1$ is the wage you would have earned if you had attended.

Treatment Effects

- ▶ ATE = $\mathbb{E}(Y_1 Y_0)$ is the average effect of *forcing* a randomly-chosen person to attend university.
- ▶ TOT = $\mathbb{E}(Y_1 Y_0 | D = 1)$ is the average effect of attending university for the sort of people who choose to attend.

Problem: Selection Bias

- ▶ We don't force randomly-chosen people to attend university!
- People who choose to attend are likely different in many ways

Selection Bias

Naïve Comparison of Means

$$\begin{split} \mathbb{E}(Y|D=1) - \mathbb{E}(Y|D=0) &= \mathbb{E}(Y_1|D=1) - \mathbb{E}(Y_0|D=0) \\ &= \mathbb{E}(Y_1|D=1) - \mathbb{E}(Y_0|D=0) + \mathbb{E}(Y_0|D=1) - \mathbb{E}(Y_0|D=1) \\ &= \underbrace{\mathbb{E}(Y_1 - Y_0|D=1)}_{\mathsf{TOT}} + \underbrace{\left[\mathbb{E}(Y_0|D=1) - \mathbb{E}(Y_0|D=0)\right]}_{\mathsf{Selection Bias}} \end{split}$$

How does selection matter?

- 1. TOT is probably different from ATE: selection on gains.
- 2. Average value of Y_0 ("outside option") probably varies with D.

How to solve the problem of selection bias?

Randomized Controlled Trial

- ▶ Hence: TOT = ATE and Selection Bias = 0.

Other Approaches

- ► Selection-on-observables (chapter 4, video 1, video 2)
- ► Instrumental Variables (chapter 5, tomorrow's lecture)
- Regression Discontinuity (chapter 7)
- Difference-in-differences (chapter 8)

Partial Identification

Bound the ATE without using the above approaches while allowing for selection bias.

Bounding the ATE when Y and D are Binary

- lacktriangle Example: Y=1 if you earn a PhD, D=1 if you attend an Ivy League University
- ▶ We know that *D* is *not* randomly assigned, and expect selection bias.

Starting point

- ightharpoonup Assume that (Y, D) are observed.
- ▶ Since Y is binary we know that $-1 \le ATE \le 1$ without observing any data!

$$0 \leq \mathit{Y}_0 \leq 1 \quad \text{and} \quad 0 \leq \mathit{Y}_1 \leq 1 \implies 0 \leq \mathbb{E}(\mathit{Y}_0) \leq 1 \quad \text{and} \quad 0 \leq \mathbb{E}(\mathit{Y}_1) \leq 1$$

Shorthand

$$P_{11} \equiv \mathbb{P}(Y = 1|D = 1) = \mathbb{E}[Y|D = 1] = \mathbb{E}[Y_1|D = 1]$$

 $P_{10} \equiv \mathbb{P}(Y = 1|D = 0) = \mathbb{E}[Y|D = 0] = \mathbb{E}[Y_0|D = 0]$
 $p \equiv \mathbb{P}(D = 1) = \mathbb{E}(D).$

Assumption-Free Bounds: Improving on $-1 \le ATE \le 1$ Y and D Are Observed

 $ightharpoonup
ho \Longrightarrow P_{11} \equiv \mathbb{E}[Y_1|D=1], \ P_{10} \equiv \mathbb{E}[Y_0|D=0], \ ext{and} \ p \equiv \mathbb{E}(D) \ ext{are observed}$

Iterated Expectations

$$\mathbb{E}[Y_1] = \mathbb{E}_D[\mathbb{E}(Y_1|D)] = P_{11}p + \mathbb{E}[Y_1|D=0](1-p)$$

$$\mathbb{E}[Y_0] = \mathbb{E}_D[\mathbb{E}(Y_0|D)] = \mathbb{E}[Y_0|D=1]p + P_{10}(1-p).$$

Bound the Unobserved Quantities

 $ightharpoonup \mathbb{E}[Y_1|D=0]$ and $\mathbb{E}[Y_0|D=1]$ are between 0 and 1

$$pP_{11} \leq \mathbb{E}[Y_1] \leq pP_{11} + (1-p)$$

$$(1-p)P_{10} \leq \mathbb{E}[Y_0] \leq p + (1-p)P_{10}$$

Assumption-Free Bounds: Width Equals 1

Previous Slide

$$pP_{11} \le \mathbb{E}[Y_1] \le pP_{11} + (1-p)$$

 $(1-p)P_{10} \le \mathbb{E}[Y_0] \le p + (1-p)P_{10}$

Combine These

$$pP_{11} - (1-p)P_{10} - p \le \mathbb{E}[Y_1 - Y_0] \le pP_{11} - (1-p)P_{10} + (1-p).$$

Written More Compactly

$$q \leq \mathsf{ATE} \leq (q+1), \quad q \equiv [pP_{11} - (1-p)P_{10} - p]$$

▶ Half as wide as $-1 \le ATE \le 1$ but always includes zero

Add Assumptions, Tighten the Bounds (Details in Lecture Notes)

Monotone Treatment Selection (MTS)

▶ Suppose we know direction of self-selection into treatment, e.g. *positive*:

$$\mathbb{E}(Y_1|D=0) \leq \mathbb{E}(Y_1|D=1) \quad \text{and} \quad \mathbb{E}(Y_0|D=0) \leq \mathbb{E}(Y_0|D=1).$$

▶ Positive MTS gives an improved *upper bound* for the ATE:

$$q \le \mathsf{ATE} \le P_{11} - P_{10} \le (q+1), \quad q \equiv [pP_{11} - (1-p)P_{10} - p]$$

Monotone Treatment Response (MTR)

- ▶ Suppose we know the direction of the **causal effect**: e.g. *positive effect*: $Y_1 > Y_0$.
- ▶ Positive MTR gives an improved *lower* bound for the ATE, namely zero:

$$0 \leq \mathsf{ATE} \leq (q+1)$$

A Comparison of Bounds

- Preceding bounds are sharp under their respective assumptions. How tight are they?
- Example: suppose that 8% of Ivy League graduates earn a PhD versus 1.5% of the general public and that 0.2% of people attend an Ivy League institution.

$$(P_{11}=0.08,\,P_{10}=0.015,\,p=0.002) \implies q \equiv [pP_{11}-(1-p)P_{10}-p] \approx -0.017$$

No Asumptions: $[q,q+1] \approx [-0.017,0.983]$

Positive MTS: $[q,P_{11}-P_{10}] \approx [-0.017,0.065]$

Positive MTR: $[0,q+1] \approx [0,0.983]$

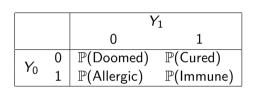
Positive MTS + MTR: $[0,P_{11}-P_{10}] = [0,0.065]$.

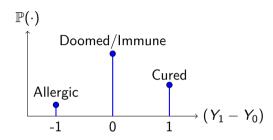
► Here positive MTR has little effect; positive MTS makes a dramatic difference!

Bounding the Distribution of Treatment Effects

- ightharpoonup Randomly assign $D \implies ATE$ point identified: no selection bias!
- \triangleright (Y_0, Y_1) never observed for same person; can't learn joint distribution.
- Anything that depends on this joint distribution is not point identified.
- ► Examples: $Var(Y_1 Y_0)$, $\mathbb{P}(Y_1 Y_0 > 0)$
- ▶ Can we partially identify the distribution of treatment effect $(Y_1 Y_0)$?
- ▶ Start with binary *Y* case; then consider the general case.

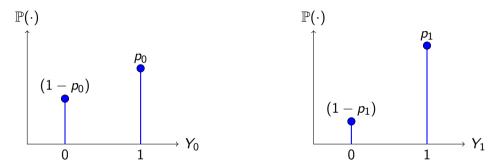
Unobserved: Joint Distribution of (Y_0, Y_1) , Distribution of $(Y_1 - Y_0)$





- Dangerous disease, and dangerous treatment.
- ▶ Treatment helps some people (the "Cured"), harms others (the "Allergic").
- ► Treatment has no effect on other people (the "Doomed" and "Immune")
- ► Are more people helped than harmed?

Observed: Marginal Distributions of Y_0 and Y_1



- ightharpoonup Assume (Y, D) come from a randomized, double-blind, placebo-controlled trial.
- \triangleright p_0 is the share of untreated who recover; p_1 is the share of treated who recover.
- ▶ The ATE is $p_1 p_0$
- ▶ Try to bound what we can't observe using what we can observe.

From Joint (Unobserved) to Marginals (Observed)

Recall:
$$p_0 \equiv \mathbb{P}(Y_0 = 1)$$
 and $p_1 \equiv \mathbb{P}(Y_1 = 1)$.

Shorthand: $\alpha \equiv \mathbb{P}(Allergic)$

Previous Slide

$$(1-p_0) = \mathbb{P}(\mathsf{Doomed}) + \mathbb{P}(\mathsf{Cured})$$
 $p_0 = \mathbb{P}(\mathsf{Allergic}) + \mathbb{P}(\mathsf{Immune})$
 $(1-p_1) = \mathbb{P}(\mathsf{Doomed}) + \mathbb{P}(\mathsf{Allergic})$
 $p_1 = \mathbb{P}(\mathsf{Cured}) + \mathbb{P}(\mathsf{Immune})$

Rearranging

$$\mathbb{P}(\mathsf{Immune}) = p_0 - \alpha$$
 $\mathbb{P}(\mathsf{Doomed}) = (1 - p_1) - \alpha$
 $\mathbb{P}(\mathsf{Cured}) = (p_1 - p_0) + \alpha$

lacktriangle Everything is written in terms of observables (p_0, p_1) and $\alpha!$

Bounding $\alpha \equiv \mathbb{P}(\mathsf{Allergic})$

Previous Slide

 $ightharpoonup \mathbb{P}(\mathsf{Immune}) = p_0 - \alpha, \ \mathbb{P}(\mathsf{Doomed}) = (1 - p_1) - \alpha, \ \mathbb{P}(\mathsf{Cured}) = (p_1 - p_0) + \alpha$

Probabilities are between 0 and 1

▶ Apply Immune, Doomed, and Cured to bound α :

$$0 \le (p_1 - p_0) + \alpha \le 1, \quad 0 \le (1 - p_1) - \alpha \le 1, \quad 0 \le p_0 - \alpha \le 1.$$

Simplify

▶ Rearrange the preceding, and combine with $0 \le \alpha \le 1$

$$\max\{-\mathsf{ATE},0\} \le \alpha \le \min\{p_0,(1-p_1)\}, \quad \mathsf{ATE} = (p_1-p_0).$$

(Pointwise) Sharp Bounds for Distribution of Treatment Effects

Previous Slide

- $ightharpoonup \mathbb{P}(\mathsf{Immune}) = p_0 \alpha, \ \mathbb{P}(\mathsf{Doomed}) = (1 p_1) \alpha, \ \mathbb{P}(\mathsf{Cured}) = (p_1 p_0) + \alpha$
- ► $\max\{-(p_1-p_0),0\} \le \alpha \le \{p_0,(1-p_1)\}$

Shorthand

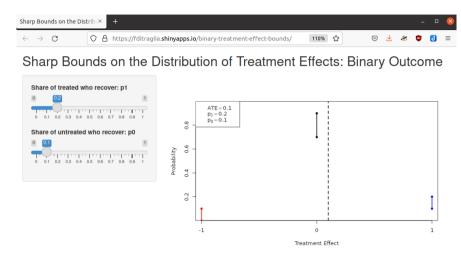
 $\underline{\alpha} \equiv \max\{-(p_1 - p_0), 0\}, \quad \overline{\alpha} \equiv \min\{p_0, (1 - p_1)\}$

Combine

 $ightharpoonup \operatorname{Recall}$ that $\alpha \equiv \mathbb{P}(\operatorname{Allergic}) = \mathbb{P}(Y_1 - Y_0 = -1)$

$$egin{aligned} \underline{lpha} & \leq \mathbb{P}(Y_1 - Y_0 = -1) \leq \overline{lpha} \ (1 - p_1) + p_0 - 2\overline{lpha} & \leq \mathbb{P}(Y_1 - Y_0 = 0) \leq (1 - p_1) + p_0 - 2\underline{lpha} \ (p_1 - p_0) + \underline{lpha} & \leq \mathbb{P}(Y_1 - Y_0 = 1) \leq (p_1 - p_0) + \overline{lpha} \end{aligned}$$

https://fditraglia.shinyapps.io/binary-treatment-effect-bounds/



The General Case: Fan & Park (2010)

- ightharpoonup Above we assumed that (Y_0, Y_1) were both binary.
- ▶ We asked which joint distributions were **not ruled out** based on the marginals.
- Pointwise sharp bounds for $\mathbb{P}(Y_1 Y_0 = -1)$, $\mathbb{P}(Y_1 Y_0 = 0)$ and $\mathbb{P}(Y_1 Y_0 = 1)$.
- Special case of a general result: Fan and Park (2010).
- Same basic idea, but math is harder when (Y_0, Y_1) may not be binary.
- This is a result you may actually use in practice!
- Explain their result without proving it.

Fan & Park (2010) Bounds

Observables

 $ightharpoonup F_0(y) \equiv \mathbb{P}(Y_0 \leq y) \text{ and } F_1(y) \equiv \mathbb{P}(Y_1 \leq y)$

Goal

▶ Sharp bounds for $F(\delta) \equiv \mathbb{P}(Y_1 - Y_0 \leq \delta)$

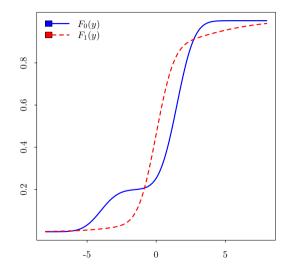
Notation

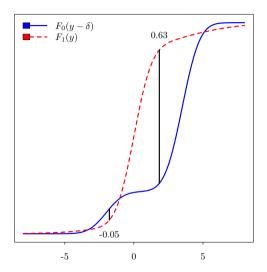
$$\underline{F}(\delta) \equiv \sup_{y} F_{1}(y) - F_{0}(y - \delta)$$
$$\overline{F}(\delta) \equiv 1 + \left[\inf_{y} F_{1}(y) - F_{0}(y - \delta)\right]$$

Theorem

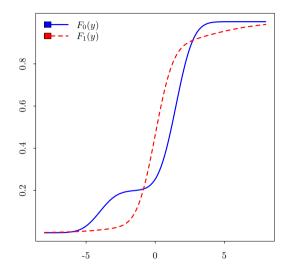
▶ For any δ , $0 \le \underline{F}(\delta) \le F(\delta) \le \overline{F}(\delta) \le 1$. These bounds are (pointwise) sharp.

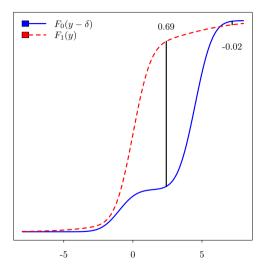
Left: $\delta = 0$, Right: $\delta = 2$



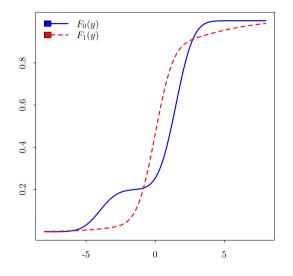


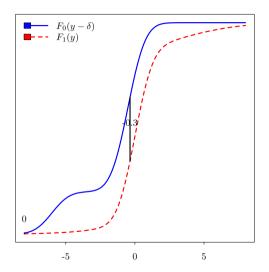
Left: $\delta = 0$, Right: $\delta = 3$





Left: $\delta = 0$, Right: $\delta = -2$





All the bounds!

