

R Lab 2 - Identifiability & the Simple Substitution Estimator

Introduction to Causal Inference

Goals:

1. Review the steps 1-5 of the roadmap: (1) specify the causal model, (2) specify the causal question, (3) specify the observed data and its link to the causal model, (4) assess identifiability and (5) specify a statistical estimand and statistical model.
2. Obtain the value the statistical estimand closed form.
3. Obtain the value the statistical estimand with simulations.
4. Introduce and implement the simple substitution estimator based on the G-Computation formula.
5. Use simulations to evaluate the properties of estimators.

Next lab:

Code discrete SuperLearner to select the estimator with the lowest cross-validated risk. Use R **SuperLearner** package to build the best convex combination of candidate algorithms and to evaluate the performance of SuperLearner.

Reminder:

This is not an R class. However, software is an important bridge between the statistical concepts and implementation.

1 Background Story

“[The Hunger Games] is written in the voice of sixteen-year-old Katniss Everdeen, who lives in a post-apocalyptic world in the country of Panem where the countries of North America once existed. The Capitol, a highly advanced metropolis, holds hegemony over the rest of the nation. The Hunger Games are an annual event in which one boy and one girl aged 12 to 18 from each of the 12 districts surrounding the Capitol are selected by lottery [as ‘tributes’] to compete in a televised battle in which only one person can survive.” - Source: Wikipedia “The Hunger Games”

Some of the tributes have trained extensively for this tournament. The life experiences of other tributes have engendered certain abilities/advantages (e.g. strength, tree climbing, markmanship). Prior to the tournament, a committee of judges assigns a score to each the tribute indicating his/her likelihood of winning. Once the tournament starts, forming alliances and sponsorship can aid in survival. A lone victor returns to their district and is showered with wealth and other resources.

Suppose we are interested in the effect of forming an alliance on the probability of surviving through the first 24 hours. We have randomly sampled one tribute from each year of the games. Let $W1$ denote the tribute’s gender with $W1 = 1$ being male and $W1 = 0$ female. Let $W2$ denote the score from the judges. Let A be an indicator of whether an alliance is formed or not, and Y be an indicator of survival through the first 24 hours. Finally, let $W3$ be an indicator of whether the tribute receives aid from sponsors during the tournament. Our goal is to evaluate the effect of forming an alliance on the probability of surviving through the first 24 hours.

This study can be translated into the following directed acyclic graph:

1. Translate the DAG into the corresponding structural causal model \mathcal{M}^F .

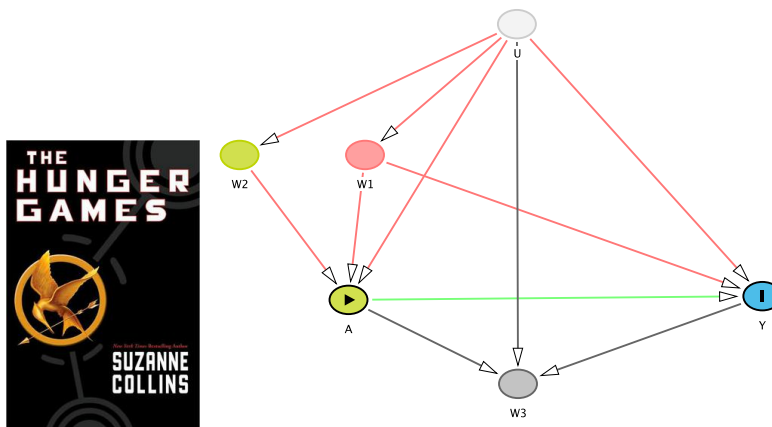


Figure 1: Directed Acyclic Graph for the Hunger Games study.

2. Are there any exclusion restrictions?
3. Are there any restrictions on the distribution of the exogenous variables P_U ? In other words, are there any independence assumptions?
4. Specify the causal question and parameter.
5. Suppose the observed data consist of n independent, identically distributed (i.i.d.) draws of the random variable $O = (W1, W2, A, Y, W3) \sim P_0$. Specify the link between the SCM and the observed data. Does the SCM place any restrictions on the statistical model \mathcal{M} ?
6. Using the backdoor criteria, assess identifiability of $\Psi^F(P_{U,X})$. If not identified, under what assumptions would it be?
7. Specify the target parameter of the observed data distribution $\Psi(P_0)$.

2 A specific data generating process

The above SCM is compatible with many possible data generating processes. Recall \mathcal{M}^F is a causal model for the set of possible distributions $P_{U,X}$ for (U, X) . Now, consider the a specific data generating process, where each of the exogenous nodes U_{X_i} is drawn independently from the following distributions.

$$\begin{aligned}
 U_{W1} &\sim \text{Uniform}(0, 1) \\
 U_{W2} &\sim \text{Normal}(\mu = 1, \sigma^2 = 2^2) \\
 U_A &\sim \text{Uniform}(0, 1) \\
 U_Y &\sim \text{Uniform}(0, 1) \\
 U_{W3} &\sim \text{Uniform}(0, 1)
 \end{aligned}$$

Given the U 's, the endogenous variables are deterministically generated as:

$$\begin{aligned}
 W1 &= \mathbb{I}[U_{W1} < 0.45] \\
 W2 &= 0.75 * U_{W2} \\
 A &= \mathbb{I}[U_A < \text{expit}(-1 + 2.6 * W1 + 0.9 * W2)] \\
 Y &= \mathbb{I}[U_Y < \text{expit}(-2 + A + 0.7 * W1)] \\
 W3 &= \mathbb{I}[U_{W3} < \text{expit}(-1 + 1.3 * A + 2.9 * Y)]
 \end{aligned}$$

The *expit* function is the inverse of the logistic function:

$$\begin{aligned}\text{logit}(x) &= \log\left(\frac{x}{1-x}\right) \\ \text{expit}(x) &= \frac{1}{1 + e^{-x}}\end{aligned}$$

1. **Evaluate $\Psi(P_0)$ in closed form.** Hint: plug in the necessary functions into the G-Computation formula.
2. **Interpret $\Psi(P_0)$.**

3 Translate this data generating process into simulations

1. **First set the seed to 252.**
2. **Set the number of draws $n = 5000$.**
3. **Sample n i.i.d. observations of random variable $O = (W1, W2, A, Y, W3) \sim P_0$.** In other words, simulate the background factors U and evaluate the structural equations F . The *expit* function in R is `plogis`.
4. **Create a data frame (`data.frame`), named `Obs`, to hold these values.** The rows are the n repetitions of the experiment and the columns are the random variables. In other words, the rows are the n tributes and the columns are their characteristics. **Use the `head` and `summary` functions to get a better understanding of the data generating experiment.**

4 Simple substitution estimator based on the G-Computation formula

In the Section 2, we used our knowledge of the true distribution of the observed data P_0 to obtain the value of the target parameter. Specifically, we plugged in the true conditional mean $E_0(Y|A, W)$ and the marginal distribution $P_0(W)$ into the G-computation formula:

$$\Psi(P_0) = E_0[E_0(Y|A = 1, W) - E_0(Y|A = 0, W)]$$

where W represents the covariates that satisfy the backdoor criteria for the effect of A on Y . In our example, $W1$ satisfies the backdoor criteria under the working model \mathcal{M}^{F*} .

In reality, we usually do not know the true distribution of the observed data P_0 . Instead, we only have a sample of n i.i.d. observations of O from P_0 . An intuitive estimator of the statistical estimand $\Psi(P_0)$ is the simple substitution estimator based on the G-Computation formula. Briefly, the algorithm estimates the relevant parts of the observed data distribution P_0 and plugs them into the parameter mapping Ψ :

- (A.) Estimate the conditional mean $E_0(Y|A, W)$ using the observed data as input.
- (B.) Estimate the marginal distribution of baseline covariates $P_0(W)$ using the observed data as input.
- (C.) Substitute these estimates into the target parameter mapping:

$$\hat{\Psi}(P_n) = \sum_{w1} [\hat{E}(Y|A = 1, W = w) - \hat{E}(Y|A = 0, W = w)] \hat{P}(W = w)$$

where P_n denotes the empirical distribution, which puts weight $1/n$ on each copy O_i , $i = 1, \dots, n$.

Formally, an estimator $\hat{\Psi}$ is a mapping from the set of possible empirical distributions P_n to the parameter space (\mathbb{R}) . In other words, $\hat{\Psi}$ is a function with input as the observed data (a realization of P_n) and output a value in the parameter space (e.g. a number). The estimator should respect the statistical model \mathcal{M} , which is non-parametric. In other words, we should not make any unfounded assumptions about the observed data distribution P_0 . In lecture, we will go over this estimator and its properties in detail.

4.1 Implementation with the NPMLE

1. **Estimate the conditional mean function with the non-parametric maximum likelihood estimator (NPMLE). Create strata of each possible value of $(A, W1)$ and take the empirical mean of Y in each strata.** This is equivalent to fitting a saturated regression model.

Hint: The following code creates a vector of the outcomes among unexposed ($A = 0$) females ($W1 = 0$). The NPMLE for the conditional probability of survival for this subgroup is the empirical mean of the resulting vector:

```
> # outcomes among unexposed females
> Y.a0w0<- Y[W1==0 & A==0]
> meanY.a0w0 <- mean(Y.a0w0)
> meanY.a0w0

[1] 0.1207116
```

In words, the observed probability of survival among females, who did not make an alliance, is $\hat{E}(Y|A = 0, W1 = 0) = 12.07\%$.

2. **Estimate the marginal distribution $P_0(W1 = w1)$ with the sample proportion:**

$$\hat{P}(W1 = w1) = \frac{1}{n} \sum_{i=1}^n \mathbb{I}(W1_i = w1)$$

Again, this non-parametric estimator does not place any restrictions on the statistical model.

3. **Substitute these estimates into the parameter mapping:**

$$\begin{aligned} \hat{\Psi}(P_n) = & [\hat{E}(Y|A = 1, W1 = 1) - \hat{E}(Y|A = 0, W1 = 1)] \hat{P}(W1 = 1) \\ & + [\hat{E}(Y|A = 1, W1 = 0) - \hat{E}(Y|A = 0, W1 = 0)] \hat{P}(W1 = 0) \end{aligned}$$

4.2 Implementation with parametric regression

In the previous subsection, we estimated the conditional risk $E_0(Y|A, W1)$ with the empirical mean outcome Y in strata of A and $W1$. This is equivalent to fitting a saturated parametric model for the conditional mean:

$$E_0(Y|A, W1) = P_0(Y = 1|A, W1) = \text{expit}(\beta_0 + \beta_1 A + \beta_2 W1 + \beta_3 A W1)$$

1. **To gain familiarity with R and the simple substitution estimator, use the `glm` function to fit the conditional mean function $E_0(Y|A, W1)$ with logistic regression. Be sure to specify the arguments `family='binomial'` and `data=0bs`.**

Hint: To get interaction terms, try the formula $Y \sim A * W1$.

2. **Copy the data set `0bs` into two new data frames `txt` and `control`. Then set `A=1` for all units in `txt` and `A=0` for all units in `control`.**

Hint: Columns of a data frame can be accessed with the `$` operator.

3. **Now use the `predict` function to get the expected outcomes for each individual under the intervention $\hat{E}(Y|A = 1, W1)$. Be sure to specify the arguments `newdata=txt` and the `type='response'`.**
4. **Now use the `predict` function to get the expected outcomes for each individual under the control $\hat{E}(Y|A = 0, W1)$. Be sure to specify the arguments `newdata=control` and the `type='response'`.**

5. **Evaluate the statistical parameter by substituting the predicted mean outcomes under the treatment and under the control into the G-Computation formula.** The sample proportion is a non-parametric maximum likelihood estimator of the marginal distribution of $W1$. So we can just take the empirical mean of the difference in the predicted outcomes for each subject:

$$\hat{\Psi}(P_n) = \frac{1}{n} \sum_{i=1}^n \left[\hat{E}(Y_i|A=1, W1_i) - \hat{E}(Y_i|A=0, W1_i) \right]$$

5 Estimate the bias, variance and mean squared error (MSE) of the substitution estimator.

Simulations are useful for evaluating the properties of estimators. We will focus on estimating the bias, variance and mean squared error of the simple substitution estimator. Specifically, for $R = 500$ iterations, we will sample $n = 200$ i.i.d. observations from P_0 , implement the simple substitution estimator based on the G-Computation formula, and save the resulting estimate ψ_n .

1. **Set R to 500 and n to 200.**
2. **Create a vector estimates of length $R = 500$ to hold the estimated values ψ_n obtained at each iteration.**
Hint: Use the `rep` function to create a vector of missing values NA.
3. **Inside a for loop from 1 to $R = 500$, sample n i.i.d. observations of random variable $O = (W1, W2, A, Y, W3)$; implement the simple substitution estimator using the saturated regression model (adjusting for A and $W1$), and save the resulting estimate ψ_n as an entry in the vector estimates.**

Hint: A simple example of a `for` loop is given below. More information on the syntax can be found with `?for`.

```
> # this code creates an empty vector "temp" of length 10
> # in the for loop, the empty values are replaced by 2*index
> temp<- rep(NA, 10)
> for(i in 1:10) {
+   temp[i]<- 2*i
+ }
> temp
```

```
[1]  2  4  6  8 10 12 14 16 18 20
```

4. **What is the average value of the estimates of $R = 500$ trials?**
5. **Estimate the bias of the estimator.** What is the average deviation of the estimate and the truth $\Psi(P_0)$? Hint: use the `mean` function.

$$\text{Bias}(\hat{\Psi}(P_n)) = E_0(\hat{\Psi}(P_n) - \Psi(P_0))$$

6. **Estimate the variance of the estimator.** How much do the estimates vary across samples? Hint: use the `var` function.

$$\text{Variance}(\hat{\Psi}(P_n)) = E_0 \left(\left(\hat{\Psi}(P_n) - E_0[\hat{\Psi}(P_n)] \right)^2 \right)$$

7. **Estimate the mean squared error of the estimator.** On average, how far are the estimates from the truth?

$$\begin{aligned} MSE(\hat{\Psi}(P_n)) &= E_0 \left(\left(\hat{\Psi}(P_n) - \Psi(P_0) \right)^2 \right) \\ &= Bias^2 + Variance \end{aligned}$$

6 More practice

Suppose the Capitol (people in charge of the Hunger Games) demand that you estimate the conditional mean outcome, given the intervention and all the covariates, according to following parametric regression model:

$$E_0(Y|A, W1, W2, W3) = \text{expit}(\beta_0 + \beta_1 A + \beta_2 W1 + \beta_2 W2 + \beta_3 W3)$$

In other words, they believe that conditional probability of survival through the first 24 hours is a linear (on the logit scale) function of the intervention (alliance), all the pre-exposure covariates ($W1, W2$) and a post-exposure covariate ($W3$). This “knowledge” changes our SCM \mathcal{M}^F , because it restricts the set of allowed functions f_Y . This “knowledge” also changes our statistical model \mathcal{M} , because it restricts the allowed conditional distributions for Y given $(A, W1, W2, W3)$.

1. **Does the backdoor criteria hold conditional on $W1, W2$ and $W3$ (assuming independence of the errors)?**
2. **For $R = 500$ iterations, repeat the above process of sampling $n = 200$ observations, fitting the conditional mean outcome with a logistic model (adjusting now for $A, W1, W2$ and $W3$), obtaining the predicted values under $A = 1$ and $A = 0$, and substituting the estimates into the target parameter mapping.**
3. **Compare the bias, variance and mean squared error of the substitution estimators when using a saturated model (equivalent to the NPMLE) and a misspecified parametric model to estimate the conditional mean $E_0(Y|A, W)$.**