

## THE AVERAGE TREATED EFFECT ON THE TREATED

The *average treatment effect on the treated* (ATT) is

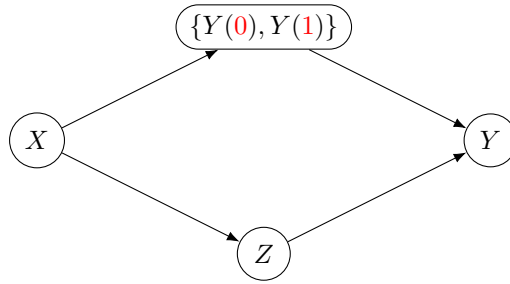
$$\mathbb{E}[Y(\mathbf{1}) - Y(\mathbf{0}) \mid Z = 1]$$

that is, the ATT aims to identify the causal effect on intervening to change  $Z = \mathbf{0}$  to  $Z = \mathbf{1}$  but *only* in the subpopulation of individuals who are *observed* to receive treatment. Note that

$$\mathbb{E}[Y(\mathbf{1}) - Y(\mathbf{0})] = \mathbb{E}[Y(\mathbf{1}) - Y(\mathbf{0}) \mid Z = 0] \Pr[Z = 0] + \mathbb{E}[Y(\mathbf{1}) - Y(\mathbf{0}) \mid Z = 1] \Pr[Z = 1]$$

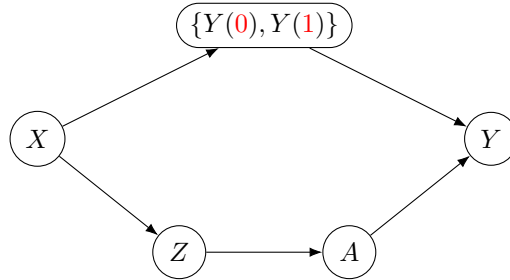
In this calculation, we imagine

- the observational distribution  $f_{X,Y,Z}^{\mathcal{O}}$  generating the observed data  $\{(x_i, y_i, z_i), i = 1, \dots, n\}$
- in the subgroup observed to have  $Z = 1$ , we then consider a second (hypothetical) experimental intervention to change  $Z$  to  $\mathbf{z}$  which over-rides the original  $Z$  if  $\mathbf{z} = \mathbf{0}$ ,
- we then consider comparison of outcomes between the two hypothetical subgroups indexed by  $\mathbf{z}$ .



$$f_X(x) f_{Z|X}(z|x) f_{Y(\mathbf{0}), Y(\mathbf{1})|X}(y_0, y_1|x) f_{Y|Z, Y(\mathbf{0}), Y(\mathbf{1})}(y|z, y_0, y_1)$$

For the ATT, we can represent the quantity of interest using a modified DAG that proposes a second hypothetical binary treatment,  $A$ . We allow  $Z$  to cause  $A$ , and then allow  $Z$  to act as a selection mechanism, but ensure that we have the conditional independence of  $X$  and  $A$  given  $Z$ ,  $X \perp\!\!\!\perp A \mid Z$ ,



$$f_X(x) f_{Z|X}(z|x) f_{Y(\mathbf{0}), Y(\mathbf{1})|X}(y_0, y_1|x) f_{A|Z}(a|z) f_{Y|A, Y(\mathbf{0}), Y(\mathbf{1})}(y|a, y_0, y_1)$$

This is the new ‘experimental’ distribution  $\mathcal{E}$ .

$$f_{Y|A, Y(\mathbf{0}), Y(\mathbf{1})}^{\mathcal{E}}(y|a, y_0, y_1) = \begin{cases} 1 & y = (1-a)y_0 + ay_1 \\ 0 & \text{otherwise} \end{cases}.$$

As  $A$  is binary, the model  $f_{A|Z}(a|z)$  must take the form

$$f_{A|Z}^{\mathcal{E}}(a|z) = p_z^a (1 - p_z)^{1-a} \quad a, z \in \{0, 1\}$$

for  $0 \leq p_z \leq 1$  for  $z = 0, 1$ . We can then express the ATT via the new DAG as

$$\mathbb{E}_{Y|A, Z}^{\mathcal{E}}[Y|A = \mathbf{1}, Z = 1] - \mathbb{E}_{Y|A, Z}^{\mathcal{E}}[Y|A = \mathbf{0}, Z = 1].$$

We have that from the DAG that

$$f_{Y|A,X,Z}^\varepsilon(y|a, x, z) \equiv f_{Y|A,X}^\varepsilon(y|a, x)$$

and hence as before

$$f_{Y|A,X}^\varepsilon(y|a, x) = \int f_{Y|A,Y(0),Y(1)}^\varepsilon(y|a, y_0, y_1) f_{Y(0),Y(1)|A,X}^\varepsilon(y_0, y_1|a, x) dy_0 dy_1 = f_{Y(a)|X}^\varepsilon(y|x).$$

Also from the DAG,  $X \perp\!\!\!\perp A \mid Z$ , so for all  $a, x, z$

$$f_{X|A,Z}^\varepsilon(x|a, z) \equiv f_{X|Z}^\varepsilon(x|z).$$

For  $\mathbf{a} = 0, 1$ , we therefore have

$$\mathbb{E}_{Y|A,Z}^\varepsilon[Y \mid A = \mathbf{a}, Z = z] = \iint y f_{Y|A,X}^\varepsilon(y \mid \mathbf{a}, x) f_{X|Z}^\varepsilon(x \mid z) dy dx.$$

Note that in this integral there is a potential *incompatibility* in the conditioning between

$$f_{Y|A,X}^\varepsilon(y \mid \mathbf{a}, x) \quad \text{and} \quad f_{X|Z}^\varepsilon(x \mid z)$$

when we try to write the integral in terms of the data generating mechanism. As before, choosing the form of the outcome conditional model

$$f_{Y|A,X}^\varepsilon(y \mid \mathbf{a}, x)$$

is to be avoided if possible due to the danger of mis-specification. If the choice is made correctly, then it can form the basis of a regression estimator.

We seek to resolve the incompatibility using the importance sampling trick, and write the expectation with respect to the observational model

$$f_{Y|X,Z}^\circ(y|x, z) f_{Z|X}^\circ(z|x) f_X^\circ(x).$$

First note that

$$f_{X|Z}^\varepsilon(x \mid z) = \frac{f_{Z|X}^\varepsilon(z \mid x) f_X^\varepsilon(x)}{f_Z^\varepsilon(z)}$$

so the integral can be rewritten

$$\frac{1}{f_Z^\varepsilon(z)} \iint y f_{Y|A,X}^\varepsilon(y \mid \mathbf{a}, x) f_{Z|X}^\varepsilon(z \mid x) f_X^\varepsilon(x) dy dx.$$

For  $\mathbf{z} = 0, 1$ , we can re-write the integrand using the importance sampling trick as

$$y f_{Y|A,X}^\varepsilon(y \mid \mathbf{a}, x) \frac{f_{Z|X}^\varepsilon(z \mid x)}{f_{Z|X}^\varepsilon(\mathbf{a} \mid x)} f_{Z|X}^\varepsilon(\mathbf{a} \mid x) f_X^\varepsilon(x)$$

which can be rearranged to

$$\left\{ y \frac{f_{Z|X}^\varepsilon(z \mid x)}{f_{Z|X}^\varepsilon(\mathbf{a} \mid x)} \right\} f_{Y|A,X}^\varepsilon(y \mid \mathbf{a}, x) f_{Z|X}^\varepsilon(\mathbf{a} \mid x) f_X^\varepsilon(x),$$

Comparing the observational and experimental DAGs, we see that for all  $x$  and  $z$

$$f_{Z|X}^\varepsilon(z \mid x) \equiv f_{Z|X}^\circ(z \mid x) \quad f_X^\varepsilon(x) \equiv f_X^\circ(x) \quad f_Z^\varepsilon(z) \equiv f_Z^\circ(z).$$

Also, we have for any  $t$  and  $y$  that

$$f_{Y|A,X}^\varepsilon(y \mid t, x) \equiv f_{Y|X,Z}^\circ(y \mid x, t).$$

Therefore we have

$$f_{Y|A,X}^\varepsilon(y \mid \mathbf{a}, x) f_{Z|X}^\varepsilon(\mathbf{a} \mid x) f_X^\varepsilon(x) \equiv f_{Y|X,Z}^\circ(y \mid x, \mathbf{a}) f_{Z|X}^\circ(\mathbf{a} \mid x) f_X^\circ(x).$$

Thus

$$\begin{aligned}
\mathbb{E}_{Y|A,Z}^\varepsilon[Y | A = \mathbf{a}, Z = z] &= \frac{1}{f_Z^\circ(z)} \iint \left\{ y \frac{f_{Z|X}^\circ(z | x)}{f_{Z|X}^\circ(\mathbf{a} | x)} \right\} f_{X,Y,Z}^\circ(x, y, \mathbf{a}) dy dx \\
&= \frac{1}{f_Z^\circ(z)} \iiint \left\{ \mathbb{1}_{\{\mathbf{a}\}}(t) y \frac{f_{Z|X}^\circ(z | x)}{f_{Z|X}^\circ(t | x)} \right\} f_{X,Y,Z}^\circ(x, y, t) dy dx dt \\
&= \frac{1}{f_Z^\circ(z)} \mathbb{E}_{X,Y,Z}^\circ \left[ \mathbb{1}_{\{\mathbf{a}\}}(Z) Y \frac{f_{Z|X}^\circ(z | X)}{f_{Z|X}^\circ(Z | X)} \right].
\end{aligned}$$

For the ATT, we are interested only in  $z = 1$ . The moment-based estimator is therefore

$$\widehat{\mathbb{E}}_{Y|A,Z}^\varepsilon[Y | A = \mathbf{a}, Z = 1] = \frac{\sum_{i=1}^n \mathbb{1}_{\{\mathbf{a}\}}(Z_i) w_1(X_i, Z_i) Y_i}{\sum_{i=1}^n \mathbb{1}_{\{1\}}(Z_i)} \quad \text{where} \quad w_z(X_i, Z_i) = \frac{f_{Z|X}^\circ(z | X_i)}{f_{Z|X}^\circ(Z_i | X_i)}$$

- When  $\mathbf{a} = 1$ ,

$$\mathbb{1}_{\{\mathbf{a}\}}(Z_i) w_1(X_i, Z_i) = \mathbb{1}_{\{1\}}(Z_i) = Z_i \quad \text{w.p. 1}$$

as the weight is identically 1, so therefore

$$\widehat{\mathbb{E}}_{Y|A,Z}^\varepsilon[Y | A = 1, Z = 1] = \frac{\sum_{i=1}^n Z_i Y_i}{\sum_{i=1}^n Z_i}$$

that is, the mean in the treated group.

- When  $\mathbf{a} = 0$ ,

$$\mathbb{1}_{\{\mathbf{a}\}}(Z_i) w_1(X_i, Z_i) = \mathbb{1}_{\{0\}}(Z_i) \frac{f_{Z|X}^\circ(1 | X_i)}{f_{Z|X}^\circ(Z_i | X_i)} = (1 - Z_i) \frac{f_{Z|X}^\circ(1 | X_i)}{f_{Z|X}^\circ(0 | X_i)}.$$

Therefore

$$\widehat{\mathbb{E}}_{Y|A,Z}^\varepsilon[Y | A = 0, Z = 1] = \frac{\sum_{i=1}^n (1 - Z_i) w(X_i) Y_i}{\sum_{i=1}^n Z_i}$$

where

$$w(X_i) = \frac{f_{Z|X}^\circ(1 | X_i)}{f_{Z|X}^\circ(0 | X_i)} = \frac{e(X_i)}{1 - e(X_i)}$$

That is, this estimator is a weighted sum of contributions from the *untreated* individuals.

Thus the estimator for the ATT is

$$\widehat{\mathbb{E}}_{Y|A,Z}^\varepsilon[Y | A = 1, Z = 1] - \widehat{\mathbb{E}}_{Y|A,Z}^\varepsilon[Y | A = 0, Z = 1] = \frac{\sum_{i=1}^n (Z_i - (1 - Z_i) w(X_i)) Y_i}{\sum_{i=1}^n Z_i}. \quad (1)$$

Under the standard assumptions, this estimator is consistent for the ATT and asymptotically normally distributed if

$$e(x) = f_{Z|X}^\circ(1 | X_i)$$

is correctly specified; that is, the estimator is *singly robust*.

**Example:** In this simulation study, we have

- $X \sim Uniform(0, 10)$
- $Z|X = x \sim Bernoulli(e(x))$ , with

$$e(x) = \frac{\exp\{-3 + 0.2x\}}{1 + \exp\{-3 + 0.2x\}}$$

- $Y|X = x, Z = z \sim Normal(\mu(x, z), 1)$  with

$$\mu(x, z) = x - \frac{1}{2}x \log x + \frac{5x}{1+x} + z(2 + 0.25x)$$

Thus the ATT is

$$2 + 0.25\mu_X(1)$$

where

$$\mu_X(1) = \int x f_{X|Z}^{\circ}(x|1) dx = \frac{\iint \mathbb{1}_{\{1\}}(z) x f_{X,Z}^{\circ}(x, z) dx dz}{\iint \mathbb{1}_{\{1\}}(z) f_{X,Z}^{\circ}(x, z) dx dz}$$

which can be estimated by the sample mean in the subset of observations with  $Z = 1$ .

```
set.seed(2984)
a1<-c(-3,0.2)
psi<-c(2,0.25)
h.func<-function(xv) {return(xv-0.5*xv*log(xv)+5*xv/(1+xv))}
#Large sample Monte Carlo calculation of true ATT
N <- 1000000
X<-runif(N,0,10)
pi.vec<-1/(1+exp(-cbind(1,X) %*% a1))
Z<-rbinom(N,1,pi.vec)
att <- psi[1] + psi[2]*mean(X[Z==1])
att

+ [1] 3.591837

Y<-h.func(X)+Z*(psi[1]+psi[2]*X)+rnorm(N)
```

Under correct specification, a regression estimator may be used

```
fit0<-lm(Y~1+offset(h.func(X))+Z+Z:X)
psi.hat<-coef(fit0)
psi.hat[1]+psi.hat[2]*mean(X[Z==1])

+          Z
+ 3.591741
```

However, under mis-specification, the incorrect answer is obtained:

```
fit1<-lm(Y~X+Z+Z:X)
psi.hat<-coef(fit1)
psi.hat[2]+psi.hat[3]*mean(X[Z==1])

+          X
+ 18.08968
```

The estimator in (1) is computed as follows, and gives the correct answer:

```
eX<-fitted(glm(Z~X,family=binomial))
w.hat<-Z + (1-Z)*eX/(1-eX)
att.hat<-sum((Z-(1-Z)*w.hat)*Y)/sum(Z)
att.hat

+ [1] 3.59099
```

**Double robustness:** To achieve *double robustness*, we augment the estimand for  $\mathbf{a} = \mathbf{0}$  as follows:

$$\mathbb{E}[Y(\mathbf{0}) \mid Z = 1] = \mathbb{E}[Y(\mathbf{0}) - \mu(X, \mathbf{0}) \mid Z = 1] + \mathbb{E}[\mu(X, \mathbf{0}) \mid Z = 1]$$

where

$$\mu(x, z) = \mathbb{E}[Y \mid X = x, Z = z]$$

is the modelled conditional mean for  $Y$ . To estimate the first term, we use

$$\widehat{\mathbb{E}}[Y(\mathbf{0}) - \mu(X, \mathbf{0}) \mid Z = 1] = \frac{\sum_{i=1}^n (1 - Z_i) w(X_i) (Y_i - \mu(X_i, \mathbf{0}))}{\sum_{i=1}^n Z_i}$$

as in the singly robust case. For the second term, we have

$$\widehat{\mathbb{E}}[\mu(X, \mathbf{0}) \mid Z = 1] = \frac{\sum_{i=1}^n Z_i \mu(X_i, \mathbf{0})}{\sum_{i=1}^n Z_i}$$

Therefore

$$\widehat{\mathbb{E}}[Y(\mathbf{0}) \mid Z = 1] = \frac{\sum_{i=1}^n (1 - Z_i) w(X_i) (Y_i - \mu(X_i, \mathbf{0})) + Z_i \mu(X_i, \mathbf{0})}{\sum_{i=1}^n Z_i}$$

which yields the augmented ATT estimator

$$\frac{\sum_{i=1}^n (Z_i - (1 - Z_i) w(X_i)) (Y_i - \mu(X_i, \mathbf{0}))}{\sum_{i=1}^n Z_i}.$$

```
#DR estimator with mu(x,z) = beta0 + Z (psi[0]+psi[1]*X)
w.mod <- lm(Y~Z+Z:X,weights=w.hat)
att.dr1<-coef(w.mod)[2]+coef(w.mod)[3]*mean(X[Z==1])
att.dr1

+          Z
+ 3.590976

#DR estimator with mu(x,z) = beta0 + beta1 X + Z (psi[0]+psi[1]*X)
w.mod2 <- lm(Y~Z*X,weights=w.hat)
att.dr2<-coef(w.mod2)[2]+coef(w.mod2)[4]*mean(X[Z==1])
att.dr2

+          Z
+ 3.590978
```

**Monte Carlo study:** we carry out a Monte Carlo study with 10000 replicates with  $n = 1000$ :

```
nreps<-10000
ests.mat<-matrix(0,nrow=nreps,ncol=4)
n<-1000
for(irep in 1:nreps) {
  X<-runif(n,0,10)
  eX0<-1/(1+exp(-cbind(1,X) %*% a1))
  Z<-rbinom(n,1,eX0)
```

```

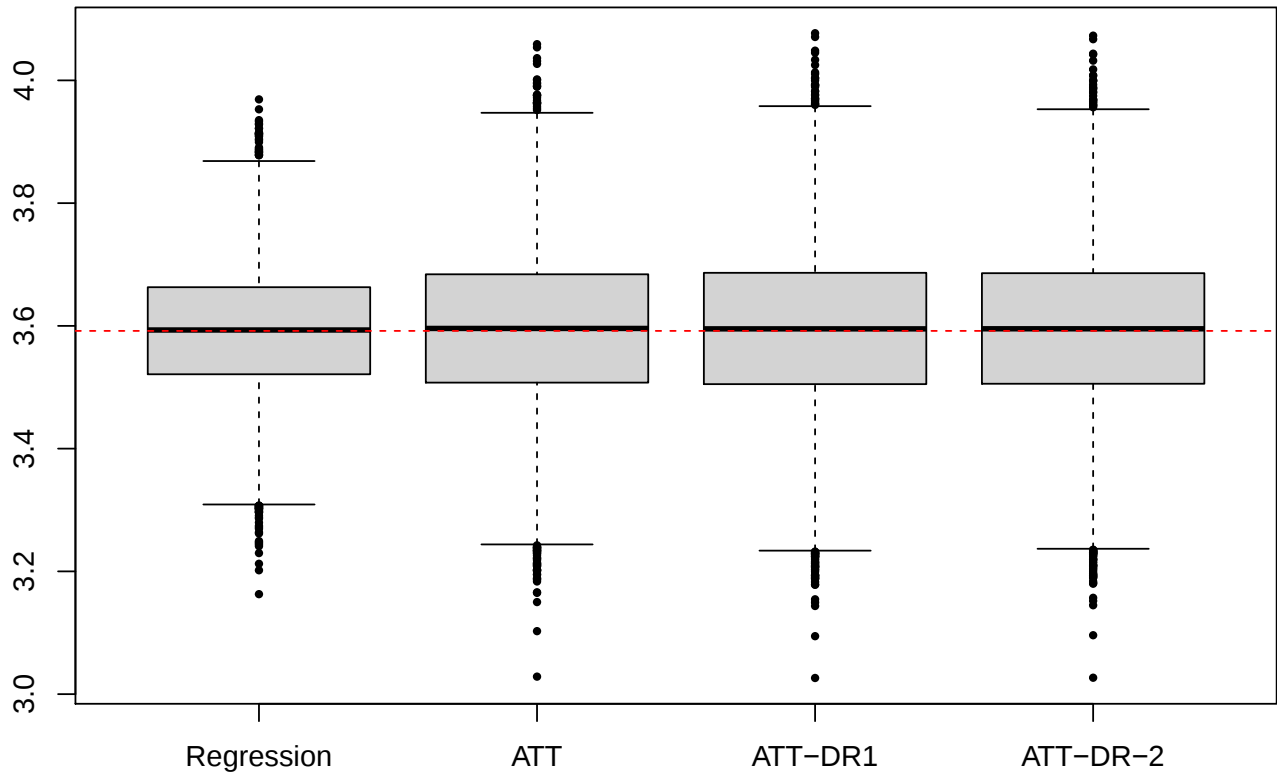
Y<-psi[1]*Z+psi[2]*Z*X+h.func(X)+rnorm(n)
fit0<-lm(Y~-1+offset(h.func(X))+Z+Z:X)
psi.hat<-coef(fit0)
ests.mat[irep,1]<-psi.hat[1]+psi.hat[2]*mean(X[Z==1]) #Regression estimator
eX<-fitted(glm(Z~X,family=binomial))
w.hat<-Z + (1-Z)*eX/(1-eX)
ests.mat[irep,2]<-sum((Z-(1-Z)*w.hat)*Y)/sum(Z) #Singly robust ATT estimator
w.mod <- lm(Y~Z+Z:X,weights=w.hat)
ests.mat[irep,3]<-coef(w.mod)[2]+coef(w.mod)[3]*mean(X[Z==1]) #Doubly robust ATT estimator 1
w.mod2 <- lm(Y~Z*X,weights=w.hat)
ests.mat[irep,4]<-coef(w.mod2)[2]+coef(w.mod2)[4]*mean(X[Z==1]) #Doubly robust ATT estimator 2
}

```

```

nv<-c('Regression','ATT','ATT-DR1','ATT-DR-2')
par(mar=c(3,3,2,0))
boxplot(ests.mat,names=nv,pch=19,cex=0.5)
abline(h=att,col='red',lty=2)

```



```

apply(ests.mat,2,var)*n #Variance
+ [1] 10.98275 17.02629 17.96629 17.74381

```