Day 3, Practical 2, Hely's solution

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1 Simulation setting 1

Task 1.

```
library(data.table)
sim.fun <- function(n) {</pre>
    # baseline covariates
    X0.1 < - runif(n, -2, 2)
    X0.2 <- rnorm(n)</pre>
    X0.3 \leftarrow rbinom(n, 1, 0.2)
    # baseline treatment (randomized)
    A0 < - rbinom(n, 1, 0.5)
    # follow-up covariates
    X1.1 \leftarrow rbinom(n, 1, plogis(-0.7 + 0.3*X0.3 + 0.8*A0))
    X1.2 <- rbinom(n, 1, plogis(0.25 - 0.55*X0.3))</pre>
    # follow-up treatment
    A1 \leftarrow rbinom(n, 1, prob=plogis(0.9 - 5*(1-A0) - 4.7*X1.1 - 4.8*X1.2))
    Y <- rbinom(n, 1, prob=plogis(-0.9 - 0.2*A0 + 1.2*X1.1 - 0.1*A1 - 0.8*A1*(X1
    .1==0)))
    return(data.table(X0.1=X0.1, X0.2=X0.2, X0.3=X0.3,
               AO=AO,
               X1.1=X1.1, X1.2=X1.2,
               A1=A1,
               Y=Y)
}
```

Task 2:

```
sim.fun <- function(n, intervene=list()) {

    # baseline covariates
    X0.1 <- runif(n, -2, 2)
    X0.2 <- rnorm(n)
    X0.3 <- rbinom(n, 1, 0.2)</pre>
```

```
# baseline treatment (randomized)
                if ("A0" %in% names(intervene)) {
                AO <- intervene$AO
                } else {
                A0 < - rbinom(n, 1, 0.5)
                }
                # follow-up covariates
                X1.1 \leftarrow rbinom(n, 1, plogis(-0.7 + 0.3*X0.3 + 0.8*A0))
                X1.2 \leftarrow rbinom(n, 1, plogis(0.25 - 0.55*X0.3))
                # follow-up treatment
                if ("A1" %in% names(intervene)) {
                A1 <- intervene$A1(X1.1)
                } else {
                A1 <- rbinom(n, 1, prob=plogis(0.9 - 5*(1-A0) - 4.7*X1.1 - 4.8*X1.2))
                # outcome
                Y \leftarrow rbinom(n, 1, prob=plogis(-0.9 - 0.2*A0 + 1.2*X1.1 - 0.1*A1 - 0.8*A1*(X1 - 0.8
                 .1==0)))
                if (length(names(intervene))>0) {
                return(mean(Y))
                } else {
                return(data.table(X0.1=X0.1, X0.2=X0.2, X0.3=X0.3,
                                                         AO=AO,
                                                        X1.1=X1.1, X1.2=X1.2,
                                                        A1=A1,
                                                        Y=Y)
                }
}
```

Task 3.

```
set.seed(15)
(sim.data <- sim.fum(1000))</pre>
```

```
X0.2 X0.3 A0 X1.1 X1.2 A1 Y
  1: 0.40845618 -0.19620228
                           0 1
                                    1
  2: -1.21982429 0.59503302
                             1 1
                             0 1
  3: 1.86583493 -1.60888231
                                         0 0 0
                                    0
  4: 0.60362212 0.04123507
                             0 1
                                    0
                                         0 1 0
                                        1 0 0
  5: -0.53171243 -1.25139144
                             0 0
                             0 1
1996: -0.72294993 -0.43541869
                                    0
                                        0 0 1
1997: -0.13592934 -0.80204340
                             0 0
                                    0
                                        0 0 0
1998: -1.67193359 -1.57095686
                             0 0
                                    0
                                       1 0 0
1999: 0.68115948 -0.72905589
                             0 1
                                    0 0 1 0
2000: 0.04187752 0.10649384
                             0 0
                                    0
                                         0 0 1
```

```
##-- naive approach 1:
fit.glm1 <- glm(Y \sim A1 + X1.1 + X1.2 + A0 + X0.1 + X0.2 + X0.3)
       family=binomial, data=sim.data)
sim.data[, EY1:=predict(fit.glm1, type="response", newdata=copy(sim.data)[, ':='(A0=1,
     A1=1)])]
sim.data[, EY0:=predict(fit.glm1, type="response", newdata=copy(sim.data)[, ':='(A0=0,
     A1=0)])]
message("naive est1 = ", sim.data[, mean(EY1-EY0)])
##-- naive approach 2:
fit.glm2 <- glm(Y \sim A1 + A0 + X0.1 + X0.2 + X0.3,
       family=binomial, data=sim.data)
sim.data[, EY1:=predict(fit.glm2, type="response", newdata=copy(sim.data)[, ':='(A0=1,
     A1=1)])]
sim.data[, EY0:=predict(fit.glm2, type="response", newdata=copy(sim.data)[, ':='(A0=0,
     A1=0)])]
message("naive est2 = ", sim.data[, mean(EY1-EY0)])
```

naive est1 = -0.181898731224723naive est2 = -0.241852261901675

Task 4. See Figure 1.

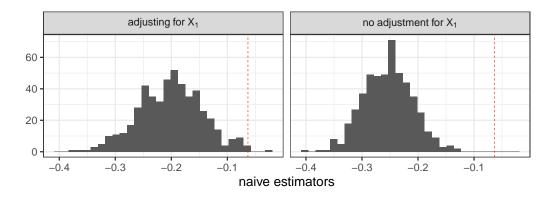


Figure 1

2 Simulation setting 2

Task 5.

```
library(data.table)
sim.fun2 <- function(n=1e6) {
    U <- rbinom(n, 1, prob=0.5)
    A <- rbinom(n, 1, prob=0.5)
    D2 <- D1 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.1*A))
    D2[D1==0] <- rbinom(n, 1, prob=plogis(2.1-3.9*U))[D1==0]
    return(data.table(A=A,D1=D1,D2=D2))
}</pre>
```

Task 6.

```
set.seed(100)
head(dat2 <- sim.fun2())</pre>
```

```
A D1 D2

1: 1 1 1

2: 1 0 1

3: 0 1 1

4: 0 1 1

5: 1 1 1

6: 1 1 1
```

6.1:

```
summary(fit.D1 \leftarrow glm(D1 \sim A, data=dat2, family=binomial))$coefficients
               Estimate Std. Error
                                        z value Pr(>|z|)
(Intercept) 0.3346503 0.002869386 116.6279
             -0.9172592 0.004113874 -222.9672
(We may note that the coefficient in front of A corresponds to:)
   p.A0 <- 0.5*(plogis(1.3-1.8)+plogis(1.3))
   p.A1 \leftarrow 0.5*(plogis(1.3-1.8-1.1)+plogis(1.3-1.1))
   log((p.A1/(1-p.A1)) / (p.A0/(1-p.A0)))
[1] -0.9098137
(And the intercept coefficient to:)
   qlogis(p.A0)
[1] 0.3297059
6.2:
   summary(fit.D2 <- glm(D2 \sim A, data=dat2[D1==0,], family=binomial)) \\ \\ scoefficients
                                          z value Pr(>|z|)
               Estimate Std. Error
(Intercept) -0.6884959 0.004631504 -148.65492
              0.2961096 0.005865281
                                        50.48515
6.3:
   dat2[, id:=1:.N]
   dat21 <- copy(dat2)[, k:=1]
   dat21$D <- dat21$D1
   dat22 <- dat2[dat2$D1==0,][, k:=2]</pre>
   dat22$D <- dat22$D2
   head(dat2.stacked <- rbind(dat21, dat22)[, t:=factor(k)])</pre>
   A D1 D2 id k D t
1: 0 1 1 1 1 1 1
2: 1 0 1 2 1 0 1
3: 1 1 1 3 1 1 1
4:0004101
5: 1 0 1 5 1 0 1
6:1 0 1 6 1 0 1
6.4:
   summary(fit.D <- glm(D~A+t, data=dat2.stacked, family=binomial))$coefficients
```

```
Estimate Std. Error z value Pr(>|z|)
(Intercept) 0.1322593 0.002595677 50.95367
            -0.5117185 0.003311977 -154.50546
                                                       0
Α
t2
            -0.3365993 0.003509668 -95.90632
                                                       Λ
   summary(fit.D <- glm(D~A*t, data=dat2.stacked, family=binomial))$coefficients
              Estimate Std. Error
                                     z value Pr(>|z|)
(Intercept) 0.3271974 0.002866385 114.1498
            -0.9103901 0.004112876 -221.3512
                                                      0
t2
                                                      0
            -1.0156933 0.005446742 -186.4772
A:t2
             1.2064997 0.007163607 168.4207
6.5:
   set.seed(10)
   U <- rbinom(n, 1, prob=0.5)</pre>
   A <- 1
   D2.1 <- D1.1 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.1*A))
   D2.1[D1.1==0] <- rbinom(n, 1, prob=plogis(2.1-3.9*U))[D1.1==0]
   D2.0 <- D1.0 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.0*A))
   D2.0[D1.0==0] <- rbinom(n, 1, prob=plogis(2.0-3.9*U))[D1.0==0]
```

Counterfactual risk difference: -0.106

```
set.seed(10)
U <- rbinom(n, 1, prob=0.5)
A <- 1
D2.1 <- D1.1 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.1*A))
D2.1[D1.1==0] <- rbinom(n, 1, prob=plogis(2.1-3.9*U))[D1.1==0]
A <- 0
D2.0 <- D1.0 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.0*A))
D2.0[D1.0==0] <- rbinom(n, 1, prob=plogis(2.0-3.9*U))[D1.0==0]
message(paste0("Counterfactual hazard ratio at 2nd time-point: ", mean(D2.1[D1.1==0])/
mean(D2.0[D1.0==0])))</pre>
```

Counterfactual hazard ratio at 2nd time-point: 1.21615895286972

message(paste0("Counterfactual risk difference: ", mean(D2.1-D2.0)))

2.1 Treatment switching

Task 7.

```
library(data.table)
sim.fun2 <- function(n=1e6) {
    U <- rbinom(n, 1, prob=0.5)
    A1 <- A <- rbinom(n, 1, prob=0.5)
    D2 <- D1 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.1*A))
    D2[D1==0] <- rbinom(n, 1, prob=plogis(2.1-3.9*U))[D1==0]
    A1[D1==0] <- rbinom(n, 1, prob=plogis(1.2+0.5*A))[D1==0]
    return(data.table(A=A,Aswitch=1*(A1!=A),D1=D1,D2=D2))
}</pre>
```

Task 8.

```
set.seed(100)
head(dat2 <- sim.fun2())</pre>
```

summary(glm(D2~A+Aswitch, family=binomial, data=dat2[A==1]))\$coefficients

```
Estimate Std. Error z value Pr(>|z|)
(Intercept) 0.5770818 0.003103637 185.93729 0
Aswitch -0.9519854 0.009647494 -98.67696 0
```

Task 9.

```
A1 <- A <- rep(1, n)
D2.11 <- D1.11 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.1*A))
D2.11[D1.11==0] <- rbinom(n, 1, prob=plogis(2.1-3.9*U))[D1.11==0]
A1[D1.11==0] <- 1

A1 <- A <- rep(1, n)
D2.10 <- D1.10 <- rbinom(n, 1, prob=plogis(1.3-1.8*U-1.1*A))
D2.10[D1.10==0] <- rbinom(n, 1, prob=plogis(2.1-3.9*U))[D1.10==0]
A1[D1.10==0] <- 0

mean(D2.11 - D2.10)
```

[1] -0.000303