# p8122\_HW1

yc4384\_Yangyang\_Chen

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# Question 1a

```
individuals <- data.frame(
    Individual = 1:8,
    Y0 = c(0, 1, 0, 1, 1, 0, 1, 0),
    Y1 = c(0, 0, 1, 0, 0, 1, 0, 0)
)

# calculate treatment effects
individuals <- individuals %>%
    mutate(TE = Y1 - Y0) # add a col TE

individuals %>%
    kable()
```

Individual	Y0	Y1	$\overline{\text{TE}}$
1	0	0	0
2	1	0	-1
3	0	1	1
4	1	0	-1
5	1	0	-1
6	0	1	1
7	1	0	-1
8	0	0	0

## Question 1b

```
ACE <- individuals %>%
  summarise(ACE = mean(TE)) %>%
  pull(ACE)

cat("The average causal effect (ACE) is:", ACE, "\n")
```

## The average causal effect (ACE) is: -0.25

## Question 1c

## Table: Observed Outcomes under Assigned Treatment

```
individuals %>%
select(Individual, Y0, Y1, Assigned_Treatment, Observed) %>%
kable()
```

Individual	Y0	Y1	Assigned_Treatment	Observed
1	0	0	1	0
2	1	0	0	1
3	0	1	1	1
4	1	0	1	0
5	1	0	0	1
6	0	1	0	0
7	1	0	0	1
8	0	0	1	0

```
mean_treatment <- individuals %>%
  filter(Assigned_Treatment == 1) %>%
  summarise(mean_treatment = mean(Observed)) %>%
  pull(mean_treatment)

mean_control <- individuals %>%
  filter(Assigned_Treatment == 0) %>%
  summarise(mean_control = mean(Observed)) %>%
  pull(mean_control)

association <- mean_treatment - mean_control
  cat("The association between treatment and outcome under specific treatment assignment is:", association</pre>
```

## The association between treatment and outcome under specific treatment assignment is: -0.5

#### Interpretation:

The association between treatment and outcome under the specific treatment assignment is -0.5, which indicates individuals in the treatment group had worse health status (mean = 0.25) compared to the control group (mean = 0.75).

### Question 1d

## Table: Observed Outcomes under Random Assignment

```
individuals %>%
  select(Individual, Y0, Y1, Random_Assigned_Treatment, Random_Assigned_Observed) %>%
  kable()
```

_					
	Individual	Y0	Y1	$Random\_Assigned\_Treatment$	$Random\_Assigned\_Observed$
	1	0	0	0	0
	2	1	0	1	0
	3	0	1	0	0
	4	1	0	0	1
	5	1	0	1	0
	6	0	1	0	0
	7	1	0	1	0
	8	0	0	1	0

```
# Calculate the association for random assignment
mean_treatment_random <- individuals %>%
    filter(Random_Assigned_Treatment == 1) %>%
    summarise(mean_treatment_random = mean(Random_Assigned_Observed)) %>%
    pull(mean_treatment_random)

mean_control_random <- individuals %>%
    filter(Random_Assigned_Treatment == 0) %>%
    summarise(mean_control_random = mean(Random_Assigned_Observed)) %>%
    pull(mean_control_random)

association_random <- mean_treatment_random - mean_control_random
cat("The association between treatment and outcome under random assignment is:", association_random, "\)</pre>
```

## The association between treatment and outcome under random assignment is: -0.25

```
# Compare with part 1b result (ACE)
cat("The ACE calculated in part 1b is:", ACE, "\n")
```

```
## The ACE calculated in part 1b is: -0.25
```

```
cat("Comparison: Association under random assignment:", association random, "vs ACE:", ACE, "\n")
```

## Comparison: Association under random assignment: -0.25 vs ACE: -0.25