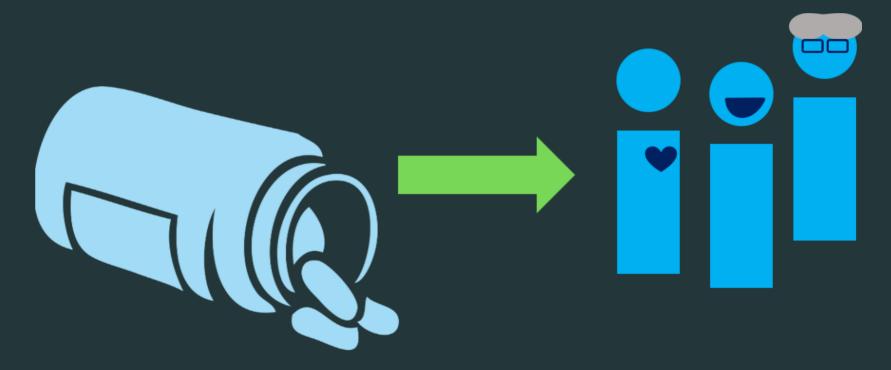
Lucy D'Agostino McGowan

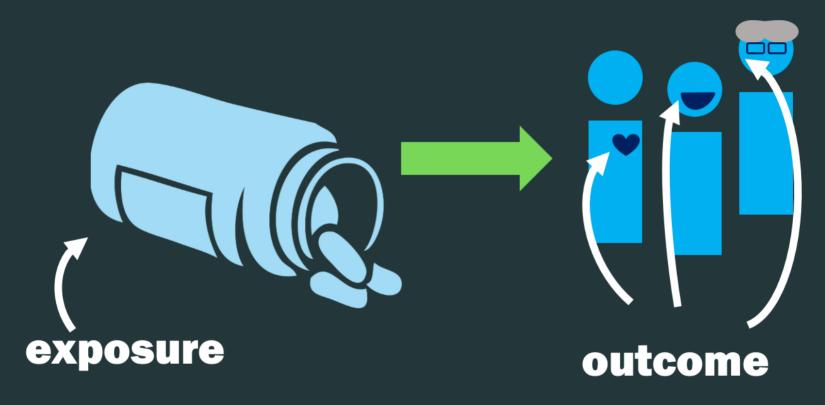
**Wake Forest University** 

2021-09-01 (updated: 2021-10-18)

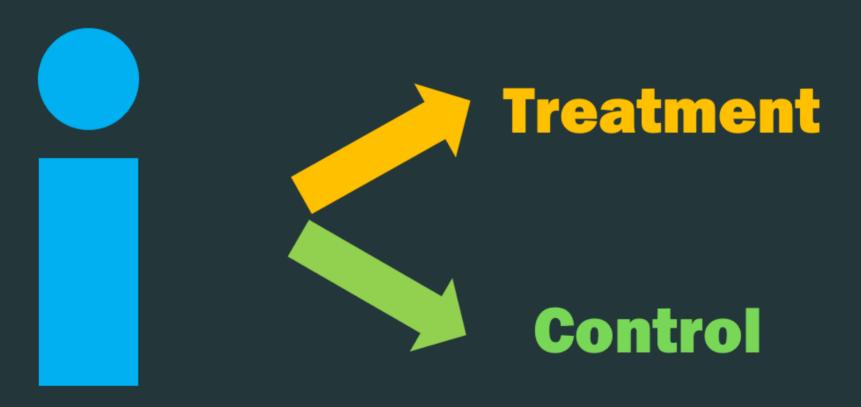
**Goal:** To answer a research question



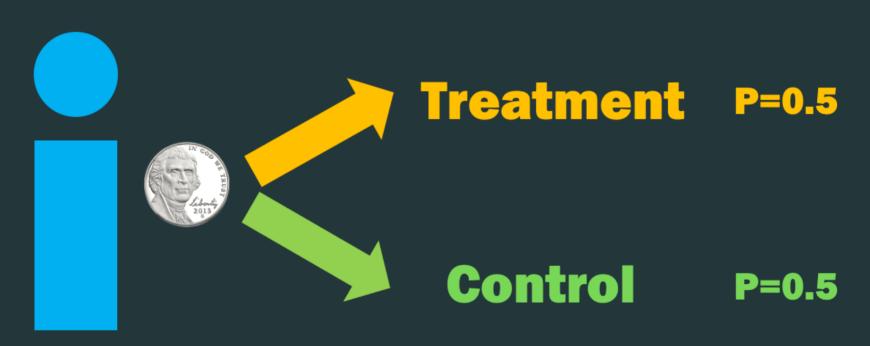
**Goal:** To answer a research question



#### **Randomized Controlled Trial**



#### **Randomized Controlled Trial**

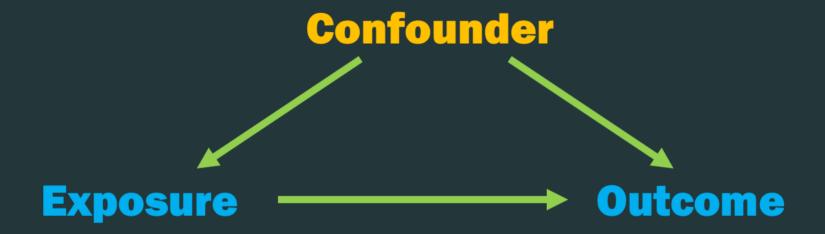




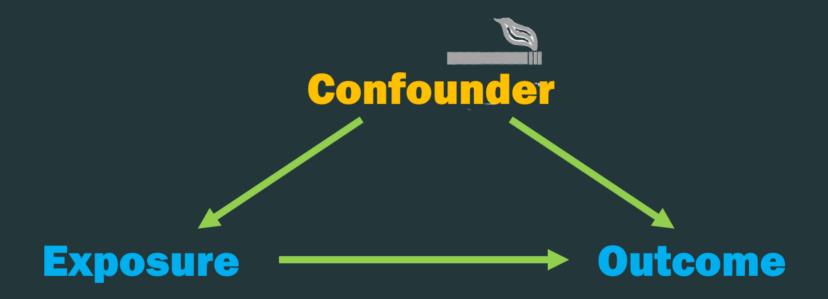




# Confounding



# Confounding



Rosenbaum and Rubin showed in observational studies, conditioning on **propensity scores** can lead to unbiased estimates of the exposure effect

- 1 There are no unmeasured confounders
- Every subject has a nonzero probability of receiving either exposure

# Fit a logistic regression predicting exposure using known covariates

$$Pr(exposure = 1) = \frac{1}{1 + \exp(-X\beta)}$$

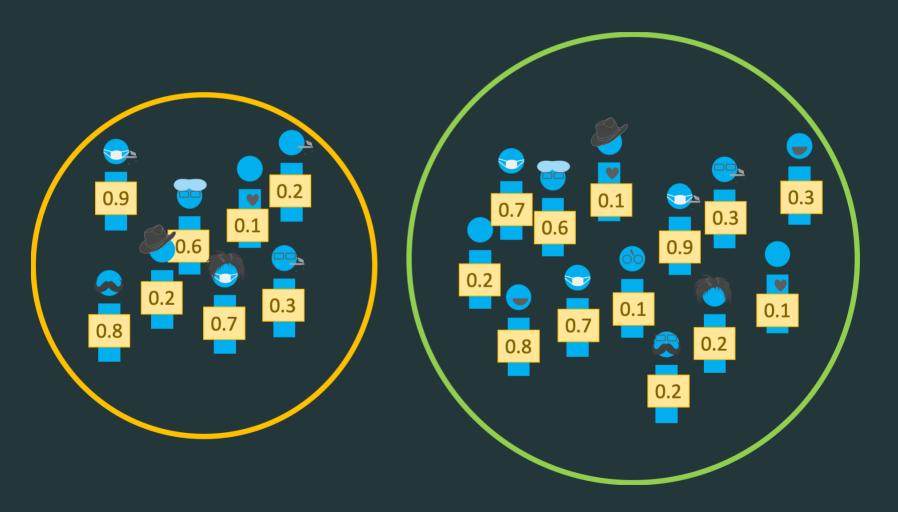
Each individuals' predicted values are the propensity scores

```
library(tidyverse)
library(broom)
```

```
glm(exposure ~ confounder_1 + confounder_2 + confounder_3 + ...,
    data = df,
    family = binomial())
```

```
glm(exposure ~ confounder_1 + confounder_2 + confounder_3 + ...,
    data = df,
    family = binomial()) %>%
    augment(type.predict = "response", data = df)
```

```
glm(exposure ~ confounder_1 + confounder_2 + confounder_3 + ...,
    data = df,
    family = binomial()) %>%
    augment(type.predict = "response", data = df)
```



#### Your turn

- 1 Using the confounders identified in the previous DAG, fit a propensity score model for qsmk
- 2 Stretch: Create two histograms, one of the propensity scores for those that quit smoking and one for those that do not