Causal Modeling in R: Whole Game

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- Specify causal question (e.g. target trial)
- 2 Draw assumptions (causal diagram)
- 3 Model assumptions (e.g., propensity)
- 4 Diagnose model (e.g., balance)
- 5 Estimate causal effects (e.g., IPW)
- 6 Sensitivity analysis (more later!)

We'll focus on the broader ideas behind each step and what they look like all together; we don't expect you to fully digest each idea. We'll spend the rest of the workshop taking up each step in detail

Do people who quit smoking gain weight?

```
library(causaldata)
                nhefs_complete_uc <- nhefs_complete |>
                         filter(censored == 0)
                nhefs_complete_uc
# A tibble: 1,566 × 67
                                                          qsmk death yrdth modth dadth
                        segn
                                                                                                                                                                                                                                                   sbp
                                                                                                                                                                                                                                                                                       dbp sex
                  <dbl> 
      1
                              233
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      5
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                                                                             0
                             420
                                                                                                                                                                                                                      17
10
                                                                                                                                               86
                                                                                                                                                                                  10
                                                                                                                                                                                                                                                   184
                                                                                                                                                                                                                                                                                       106 0
```

5

Did those who quit smoking gain weight?

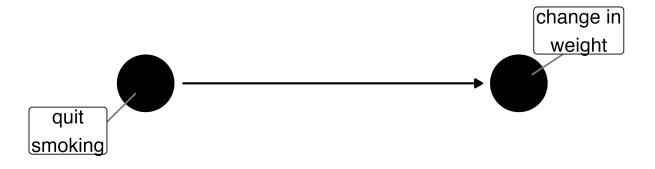
Did those who quit smoking gain weight?

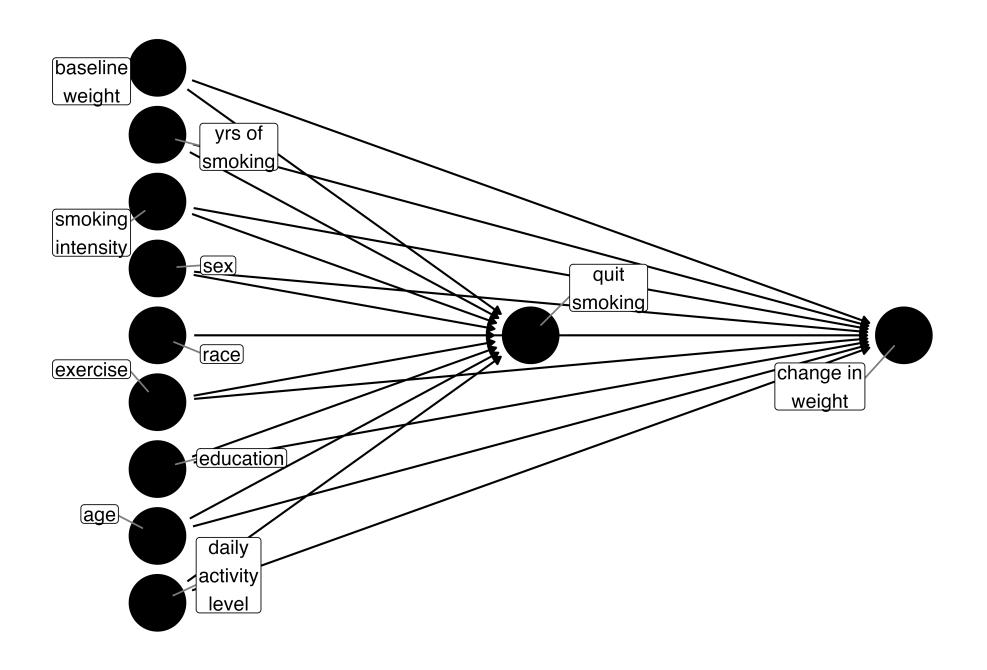
```
1 # ~2.5 KGs gained for quit vs. not quit
2 nhefs_complete_uc |>
3 group_by(qsmk) |>
4 summarize(
5 mean_weight_change = mean(wt82_71),
6 sd = sd(wt82_71),
7 .groups = "drop"
8 )

# A tibble: 2 × 3
```

```
# A tibble: 2 × 3
    qsmk mean_weight_change sd
    <dbl> <dbl> <dbl> 1.98 7.45
    2 1 4.53 8.75
```

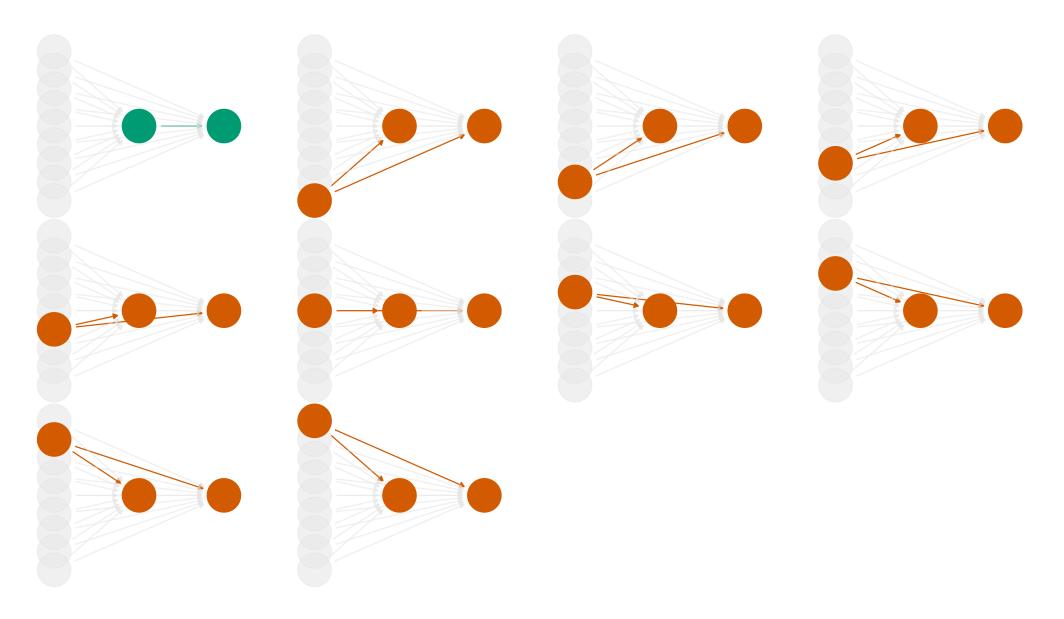
draw your assumptions



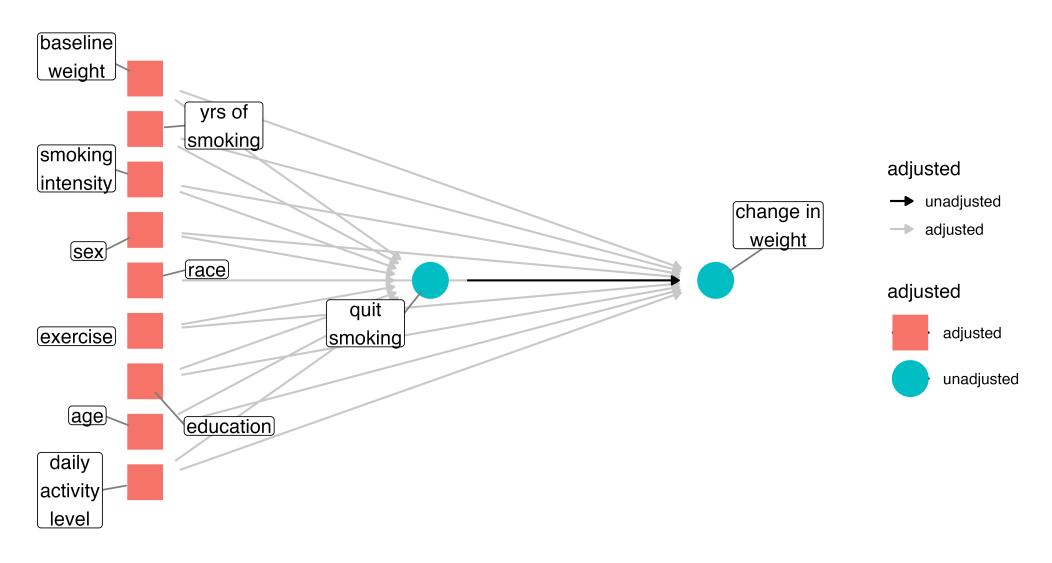


What do I need to control for?

true effect confounding effect



{active, age, education, exercise, race, sex, smokeintensity, smokeyrs, wt71}



Multivariable regression: what's the association?

```
lm(
     wt82 71 \sim qsmk + sex +
       race + age + I(age^2) + education +
       smokeintensity + I(smokeintensity^2) +
       smokeyrs + I(smokeyrs^2) + exercise + active +
       wt71 + I(wt71^2),
     data = nhefs_complete_uc
   ) |>
    tidy(conf.int = TRUE) |>
     filter(term == "qsmk")
10
```

model your assumptions

counterfactual: what if <u>everyone</u> quit smoking vs. what if <u>no one</u> quit smoking

Fit propensity score model

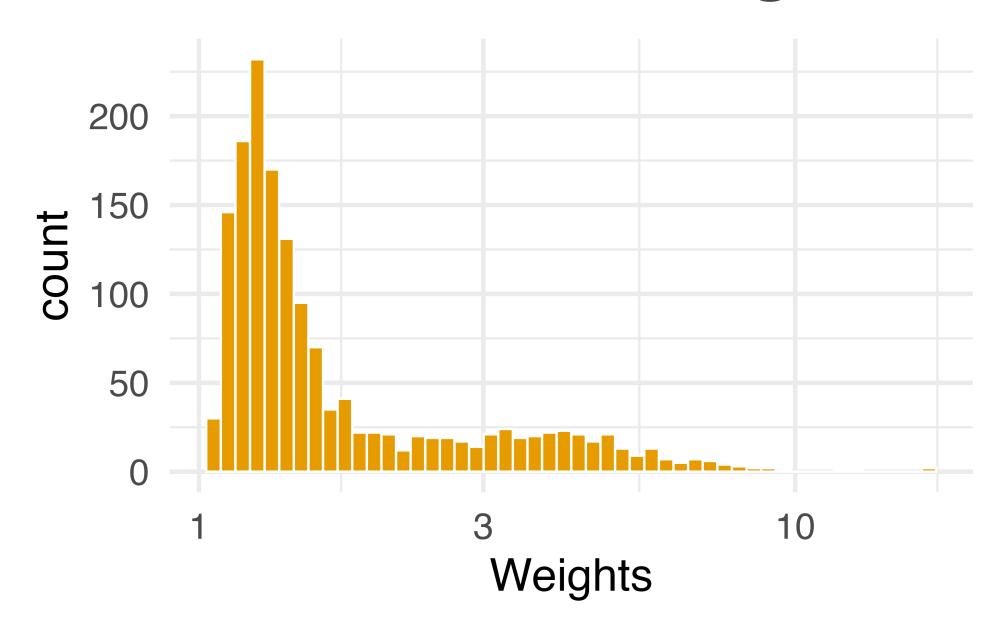
```
propensity_model <- glm(
    qsmk ~ sex +
        race + age + I(age^2) + education +
        smokeintensity + I(smokeintensity^2) +
        smokeyrs + I(smokeyrs^2) + exercise + active +
        wt71 + I(wt71^2),
    family = binomial(),
    data = nhefs_complete_uc
    )
}</pre>
```

Calculate inverse probability weights

```
1 library(propensity)
2 nhefs_complete_uc <- propensity_model |>
3  # predict whether quit smoking
4 augment(type.predict = "response", data = nhefs_complete_uc) |>
5  # calculate inverse probability
6 mutate(wts = wt_ate(.fitted, qsmk))
```

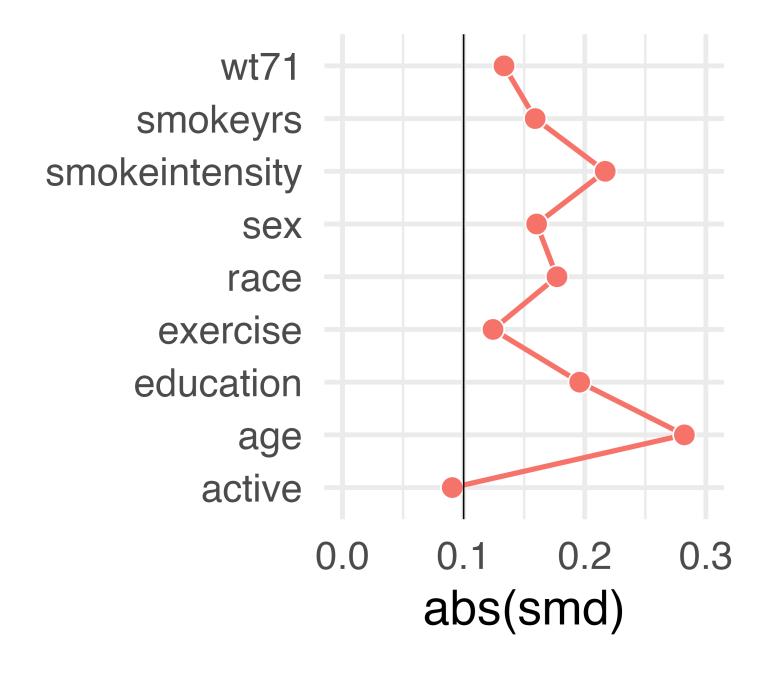
diagnose your model assumptions

What's the distribution of weights?



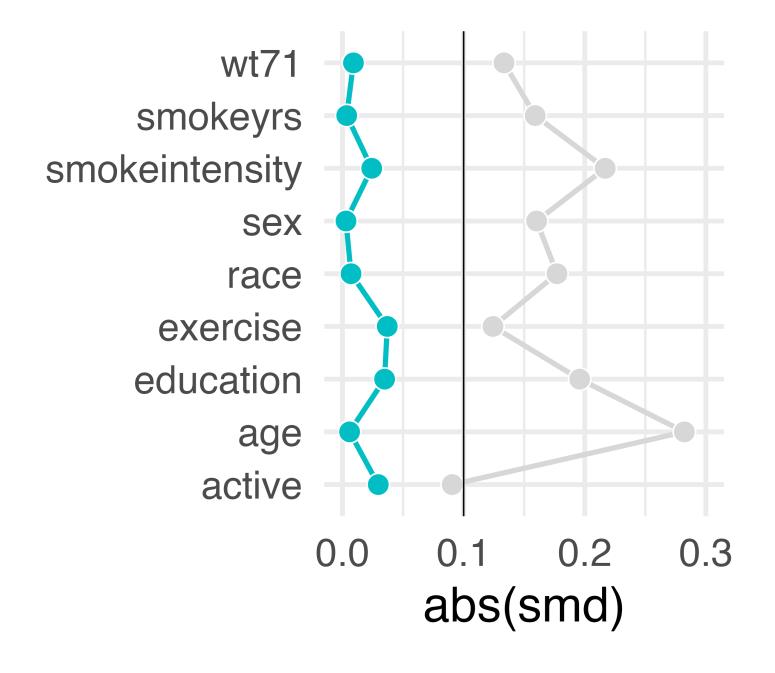
What are the weights doing to the sample?

What are the weights doing to the sample?



method

observed



method

- observed
- wts

estimate the causal effects

Estimate causal effect with IPW

```
ipw_model <- lm(
wt82_71 ~ qsmk,
data = nhefs_complete_uc,
weights = wts

)

ipw_estimate <- ipw_model |>
tidy(conf.int = TRUE) |>
filter(term == "qsmk")
```

Estimate causal effect with IPW

Let's fix our confidence intervals with robust SEs!

```
1  # also see robustbase, survey, gee, and others
2  library(estimatr)
3  ipw_model_robust <- lm_robust(
4   wt82_71 ~ qsmk,
5   data = nhefs_complete_uc,
6   weights = wts
7  )
8
9  ipw_estimate_robust <- ipw_model_robust |>
10  tidy(conf.int = TRUE) |>
11  filter(term == "qsmk")
```

Let's fix our confidence intervals with robust SEs!

Let's fix our confidence intervals with the bootstrap!

```
1  # fit ipw model for a single bootstrap sample
2  fit_ipw_not_quite_rightly <- function(.split, ...) {
3  # get bootstrapped data frame
4   .df <- as.data.frame(.split)
5
6  # fit ipw model
7  lm(wt82_71 ~ qsmk, data = .df, weights = wts) |>
    tidy()
9 }
```

```
1 fit ipw <- function(.split, ...) {</pre>
     # get bootstrapped data frame
 2
      .df <- as.data.frame(.split)</pre>
 4
 5
     # fit propensity score model
      propensity model <- glm(</pre>
 6
 7
        qsmk \sim sex +
 8
          race + age + I(age^2) + education +
9
          smokeintensity + I(smokeintensity^2) +
          smokeyrs + I(smokeyrs^2) + exercise + active +
10
11
        wt71 + I(wt71^2),
12
       family = binomial(),
13
        data = .df
14
15
16
     # calculate inverse probability weights
      .df <- propensity model |>
17
18
        augment(type.predict = "response", data = .df) |>
19
        mutate(wts = wt_ate(.fitted, qsmk))
20
21
     # fit correctly bootstrapped ipw model
22
     lm(wt82 71 ~ qsmk, data = .df, weights = wts) |>
23
       tidy()
24 }
```

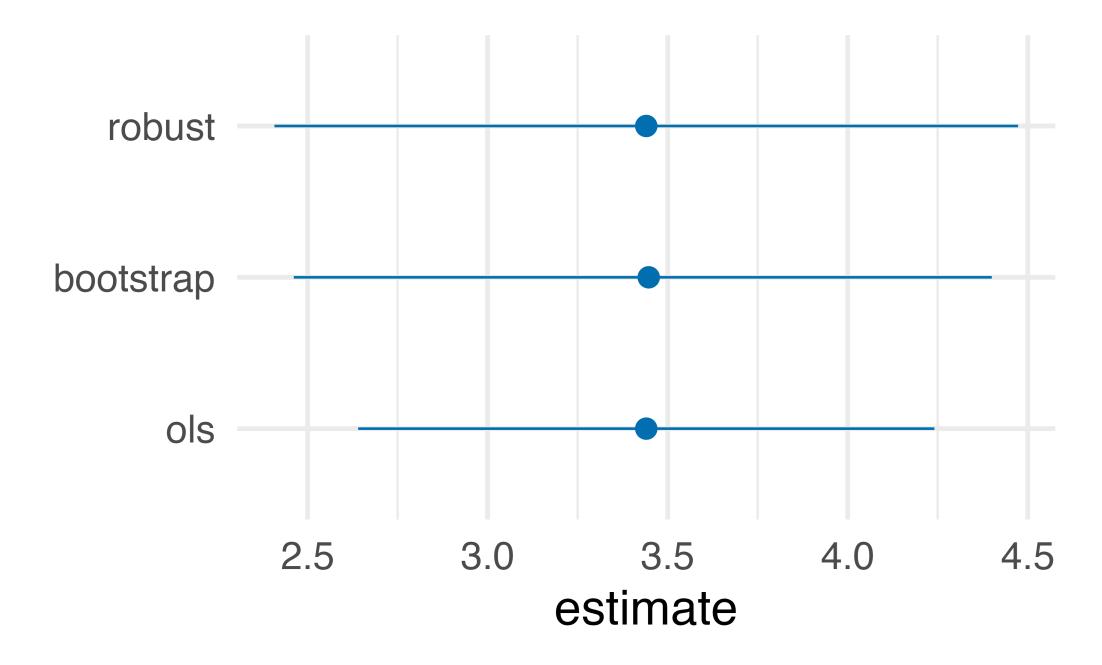
Using {rsample} to bootstrap our causal effect

```
1 # fit ipw model to bootstrapped samples
2 ipw_results <- bootstraps(nhefs_complete_uc, 1000, apparent = TRUE) |>
3 mutate(results = map(splits, fit_ipw))
```

Using {rsample} to bootstrap our causal effect

```
1 # get t-statistic-based CIs
2 boot_estimate <- int_t(ipw_results, results) |>
3   filter(term == "qsmk")
4
5 boot_estimate
```

Using {rsample} to bootstrap our causal effect



Our causal effect estimate: 3.5 kg (95% CI 2.4 kg, 4.4 kg)

Review the Quarto file... later!

Resources

Causal Inference: Comprehensive text on causal inference. Free online.

Bootstrap confidence intervals with {rsample}

R-causal: Our GitHub org with R packages and examples