# Fitting the outcome model

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#### **Outcome Model**

```
1 library(broom)
2
3 lm(outcome ~ exposure, data = df, weight
4 tidy()
```

- This will get us the point estimate
- This will get NOT us the correct confidence intervals
- Let's bootstrap them with rsample

## 1. Create a function to run your analysis once on a sample of your data

```
1 fit ipw <- function(split, ...) {</pre>
     .df <- analysis(split)</pre>
 3
     # fit propensity score model
     propensity model <- glm(</pre>
       qsmk ~ sex +
         race + age + I(age^2) + education +
         smokeintensity + I(smokeintensity^2) +
         smokeyrs + I(smokeyrs^2) + exercise + active +
9
10
         wt71 + I(wt71^2),
     family = binomial(),
11
      data = .df
12
13
14
     # calculate inverse probability weights
15
     .df <- propensity model |>
16
       augment(type.predict = "response", data = .df) |>
17
       mutate(wts = wt ate(.fitted, qsmk, exposure type = "binary"))
18
19
     # fit correctly bootstrapped ipw model
20
     lm(wt82 71 ~ qsmk, data = .df, weights = wts) |>
21
22
       tidy()
```

```
library(rsample)

# fit ipw model to bootstrapped samples
bootstrapped_nhefs <- bootstraps(
    nhefs_complete_uc,
    times = 1000,
    apparent = TRUE

    )

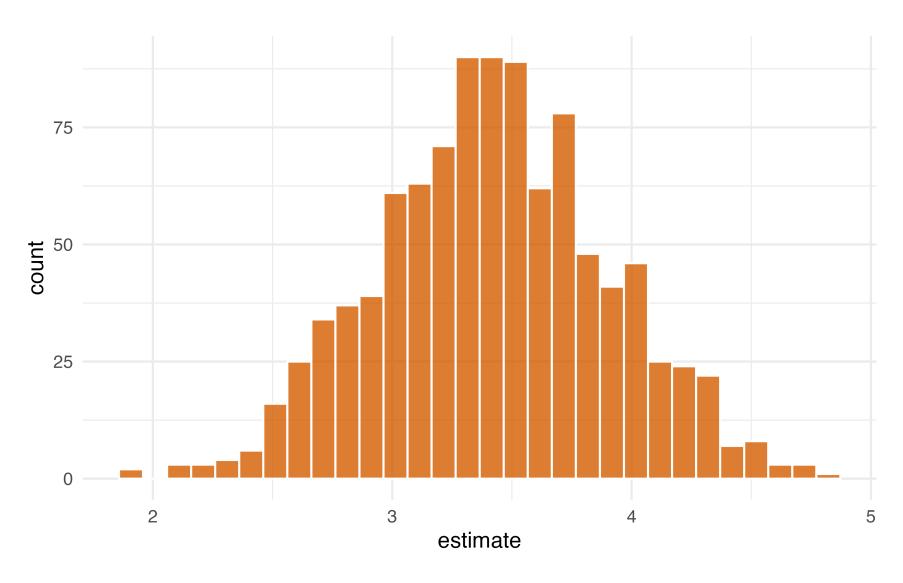
bootstrapped_nhefs</pre>
```

```
# Bootstrap sampling with apparent sample
# A tibble: 1,001 × 2
  splits
                      id
  st>
                      <chr>
 1 <split [1566/595]> Bootstrap0001
 2 <split [1566/588]> Bootstrap0002
 3 <split [1566/577]> Bootstrap0003
 4 <split [1566/592]> Bootstrap0004
 5 <split [1566/573]> Bootstrap0005
 6 <split [1566/577]> Bootstrap0006
 7 <split [1566/579]> Bootstrap0007
 8 <split [1566/577]> Bootstrap0008
 9 <split [1566/559]> Bootstrap0009
10 /-- 1:L [1[((/[00]) D--L-L----0010
```

```
ipw_results <- bootstrapped_nhefs |>
mutate(boot_fits = map(splits, fit_ipw))

ipw_results
```

```
# Bootstrap sampling with apparent sample
# A tibble: 1,001 × 3
                                   boot fits
  splits
                     id
  st>
                     <chr> <chr>
 1 <split [1566/587]> Bootstrap0001 <tibble [2 × 5]>
 2 <split [1566/555]> Bootstrap0002 <tibble [2 × 5]>
 3 <split [1566/590]> Bootstrap0003 <tibble [2 × 5]>
 4 <split [1566/599]> Bootstrap0004 <tibble [2 × 5]>
 5 <split [1566/580]> Bootstrap0005 <tibble [2 × 5]>
 6 <split [1566/574]> Bootstrap0006 <tibble [2 × 5]>
 7 <split [1566/572]> Bootstrap0007 <tibble [2 × 5]>
 8 <split [1566/569]> Bootstrap0008 <tibble [2 × 5]>
 9 <split [1566/562]> Bootstrap0009 <tibble [2 × 5]>
10 /---1!L [16///F01]\ D--L-L----0010 /L!LL1- [0 // F1\
```



#### 3. Pull out the causal effect

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

#### **Your Turn**

Create a function called ipw\_fit that fits the propensity score model and the weighted outcome model for the effect between park\_extra\_magic\_morning and wait\_minutes\_posted\_avg Using the bootstraps() and int\_t() functions to estimate the final effect.