Fitting the outcome model

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Outcome Model

```
library(broom)

lm(outcome ~ exposure, data = df, weights = wts) %>%
  tidy()
```

✅ This will get us the point estimate
❌ This will get NOT us the correct confidence intervals
📦 {rsample}

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

```r
fit_ipw <- function(split, ...) {
  .df <- analysis(split)

  # fit propensity score model
  propensity_model <- glm(
    exposure ~ confounder_1 + confounder_2 + ...
    family = binomial(),
    data = .df
  )

  # calculate inverse probability weights
  .df <- propensity_model %>%
    augment(type.predict = "response", data = .df) %>%
    mutate(wts = 1 / ifelse(exposure == 0, 1 - .fitted, .fitted))

  # fit correctly bootsrapped ipw model
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2 Use {rsample} to bootstrap our causal effect

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library(rsample)

# fit ipw model to bootstrapped samples
ipw_results <- bootstraps(df, 1000, apparent = TRUE) %>%
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3 Pull out the causal effect

```
# get t-statistic-based CIs
boot_estimate <- int_t(ipw_results, results) %>%
  filter(term == "exposure")
```

Your Turn

1. **Create a function called** ipw_fit **that fits the propensity score model and the weighted outcome model for the effect between** qsmk **and** wt82_71

2. **Using the** bootstraps() **and** int_t() **functions to estimate the final effect.**

12:00