Social Science Inquiry II

Week 6: Linear models, part II

Molly Offer-Westort

Department of Political Science, University of Chicago

Winter 2022

Loading packages for this class

> library(ggplot2)

Housekeeping

► Extra credit problem

▶ We have some data that are produced from a random sampling procedure, where they are sampled from the same population.

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- We've selected an estimating procedure, and produced a point estimate of some target estimand using our estimating procedure.
- ▶ We then produced an estimate of the standard error of our estimate.
- Now we would like to be able to say something what that means.

One way to do this is to use our estimated standard errors to give an interval of uncertainty around our point estimate.	

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▶ A valid confidence interval CI_n for a target parameter θ with coverage $1-\alpha$

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- $ightharpoonup Cl_n$ is a random interval. It is a function of the data we observe.
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- ▶ If you use valid confidence repeatedly in your work, 95% of the time, your confidence intervals will include the true value of the relevant θ .

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•	We could trivially define valid confidence intervals by including the
	entire support of the data. (Why wouldn't we want to do that?)

► The formula for the 95% confidence interval is:

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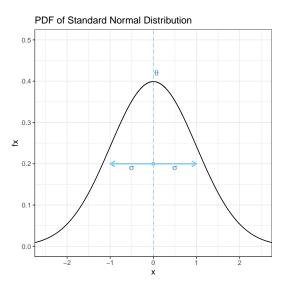
► The 1.96 value tells us how many standard errors away from the mean we need to include in our interval to get valid coverage.

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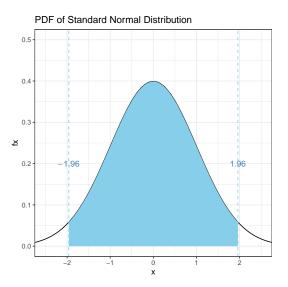
$$CI_n = (\hat{\theta}_n - 1.96 \times \hat{\text{se}}, \ \theta_n + 1.96 \times \hat{\text{se}})$$

- ► The 1.96 value tells us how many standard errors away from the mean we need to include in our interval to get valid coverage.
- ► This formula is based on a *normal approximation*, i.e., we assume the data is going to look like a normal distribution.

The normal distribution has a bell curve shape, with more density around the middle, and less density at more extreme values.



If we want to describe symmetric bounds around the mean that contain 95% of the distribution, this would be from the 2.5th percentile to the 97.5th percentile.



Returning to our example where we flip a coin twice, let X be the number of heads we observe. Our coin is *not* fair, and the probability of getting a heads is 0.8.

> X <- c(0, 1, 2)
> fx <- c(1/16, 3/8, 9/16)
> (Ex <- sum(X*fx))
[1] 1.5</pre>

Returning to our example where we flip a coin twice, let X be the number of heads we observe. Our coin is *not* fair, and the probability of getting a heads is 0.8.

Let's take a sample of size n = 100 from this distribution, and see what our confidence intervals look like

- > n <- 100
 > x_observed <- sample(X, prob = fx, replace = TRUE, size = n)
 > head(x observed)
- [1] 1 1 1 2 1 1

Our estimates of the mean and standard error of the mean.

```
> (theta_hat <- mean(x_observed))
[1] 1.5</pre>
```

> (se_hat <- sd(x_observed)/sqrt(n))</pre>

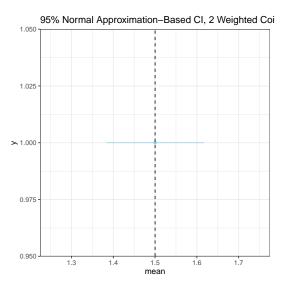
[1] 0.05945884

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- [1] 1.5
- > (se_hat <- sd(x_observed)/sqrt(n))</pre>
- [1] 0.05945884

Putting it together, the 95% normal approximation-based confidence interval.

- $> (CI95 \leftarrow c(theta_hat + c(-1,1)*1.96*se_hat))$
- [1] 1.383461 1.616539



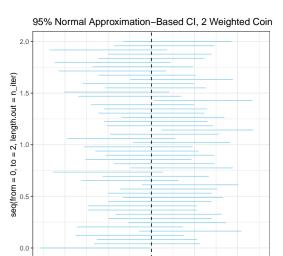
What if we did this many times?

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```
> n_iter <- 50
> x_mat <- replicate(n_iter, sample(X, prob = fx, replace = TRUE,
+ size = n))</pre>
```

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```
> n_iter <- 50
> x_mat <- replicate(n_iter, sample(X, prob = fx, replace = TRUE,
                                   size = n)
+
> CI 95f <- function(x){
+ theta hat <- mean(x)
+ se_hat <- sd(x)/sqrt(n)
+ CI hat <- theta hat +
     c('conf_lower' = -1, 'conf_upper' = 1)*1.96*se_hat
+ }
> sample_CIs <- as.data.frame(t(apply(x_mat, 2, CI_95f)))</pre>
> head(sample_CIs, 3)
  conf_lower conf_upper
   1.243391 1.496609
2 1.366997 1.613003
3 1.373477 1.606523
```



1.5

х

1.3

1.4

1.6

1.7

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Applied example

We can see this in action with respect to the paper by Devah Pager:

Pager, D. (2003). The mark of a criminal record. *American Journal of Sociology*, 108(5), 937-975.

► The study was an audit study, where pairs of white and pairs of black hypothetical job applicants applied to real jobs.

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- ► In each pair, one respondent listed a criminal record on job applications; the other did not. Otherwise, applicants were matched.
- ► The outcome is whether applicants got a callback.

```
> dfp <- data.frame(</pre>
    black = rep(c(0, 1), times = c(300, 400)),
    record = c(rep(c(0, 1), each = 150),
               rep(c(0, 1), each = 200)),
    call_back = c(
      # whites without criminal records
      rep(c(0, 1), times = c(99, 51)), # 150
      # whites with criminal records
      rep(c(0, 1), times = c(125, 25)), # 150;
      # - callbacks could be 25 or 26
      # blacks without criminal records
      rep(c(0, 1), times = c(172, 28)), # 200
      # blacks with criminal records
+
      rep(c(0, 1), times = c(190, 10)) # 200
+
```

>

```
> pager_agg <- aggregate(call_back~black + record, data = dfp, mean)</pre>
> pager_agg$race <- factor(pager_agg$black,
                            levels = c(1.0).
                            labels = c('Black', 'White'))
+
 pager_agg$criminal_record <- factor(pager_agg$record,</pre>
                                       levels = c(1.0).
                                       labels = c('Record', 'No Record'))
 pager_agg
  black record call back race criminal record
             0 0.3400000 White
                                       No Record
      0
2
               0.1400000 Black
                                       No Record
3
             1 0.1666667 White
                                          Record
             1 0.0500000 Black
                                          Record
```

American Journal of Sociology

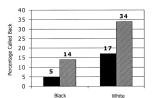


FIG. 6.—The effect of a criminal record for black and white job applicants. The main effects of race and criminal record are statically significant (P < .01). The interaction between the two is not significant in the full sample. Black bars represent criminal record; striped bars represent no criminal record.

Let's say our $\hat{\theta}$ here is the overall mean of call_back among black applicants.

```
> (theta_hat <- mean(dfp$call_back[which(dfp$black == 1)]) )
[1] 0.095</pre>
```

Let's say our $\hat{\theta}$ here is the overall mean of call_back among black applicants.

Our see is our estimate of the standard error of the mean,

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We get this by plugging in our unbiased sample variance estimate into the formula for the standard error of the mean.

```
> (se_hat <- sqrt(var(dfp$call_back[which(dfp$black == 1)])/
+ length(which(dfp$black == 1))))</pre>
```

- [1] 0.01467911
- >
- >

We can then get our 95% confidence intervals by plugging into the formula,

$$CI_n = (\hat{\theta}_n - 1.96 \times \hat{se}, \ \theta_n + 1.96 \times \hat{se})$$

 $> (CI \leftarrow c(theta_hat + c(-1,1)*1.96*se_hat))$

[1] 0.06622895 0.12377105

► As a special case of a linear model, when we regress *Y* on an indicator, we just get the sample mean of *Y*.

- ► As a special case of a linear model, when we regress *Y* on an indicator, we just get the sample mean of *Y*.
- ► In this case, estimating standard errors and confidence intervals follows the same procedures as for sample means.

```
> model <- lm(call_back ~1,
+ data = dfp[which(dfp$black == 1),])</pre>
```

```
> summary(model)
Call:
lm(formula = call_back ~ 1, data = dfp[which(dfp$black == 1),
   1)
Residuals:
  Min 10 Median 30 Max
-0.095 -0.095 -0.095 -0.095 0.905
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 0.09500 0.01468 6.472 2.84e-10 ***
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```

Residual standard error: 0.2936 on 399 degrees of freedom

> confint.default(model)

2.5 % 97.5 % (Intercept) 0.06622948 0.1237705

▶ We can think about parameters in a linear model in a similar way.

$$E[Y|X] = \beta_0 + \beta_1 X$$

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▶ The true population parameters are generally unknown.

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- We will think about our random sample being not just for one variable, but from the joint distribution of (Y, X).
- ▶ Then each $\hat{\beta}_k$ is also random, with its own sampling distribution.
- ▶ We can get a point estimate for each of the parameters, $\hat{\beta}_k$: the coefficients in our linear model.
- We also want to get an estimate of the standard errors of the estimates, $\sqrt{\hat{\mathrm{Var}}[\hat{\beta}_k]}$.

▶ Let's try this with the dfp data, where the outcome Y is call_back, regressed on black and record, interacted.

$$\hat{Y}_i = \hat{\beta}_0 + \hat{\beta}_1 \text{Black}_i + \hat{\beta}_2 \text{Record}_i + \hat{\beta}_3 \text{Black}_i \times \text{Record}_i$$

> model2 <- lm(call_back ~ black*record, data = dfp)</pre>

▶ How do we go about interpreting these coefficients and confidence intervals?

```
> summary(model2)
Call:
lm(formula = call_back ~ black * record, data = dfp)
Residuals:
   Min 10 Median 30 Max
-0.3400 -0.1667 -0.1400 -0.0500 0.9500
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 0.34000 0.02905 11.706 < 2e-16 ***
black -0.20000 0.03842 -5.205 2.55e-07 ***
record -0.17333 0.04108 -4.220 2.77e-05 ***
black:record 0.08333 0.05434 1.534 0.126
Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
Residual standard error: 0.3557 on 696 degrees of freedom
Multiple R-squared: 0.07713, Adjusted R-squared: 0.07315
F-statistic: 19.39 on 3 and 696 DF, p-value: 4.399e-12
```

► How do we go about interpreting these coefficients and confidence intervals?

```
> confint.default(model2)
```

```
2.5 % 97.5 % (Intercept) 0.28307277 0.39692723 black -0.27530765 -0.12469235 record -0.25384060 -0.09282607 black:record -0.02316776 0.18983443
```

References I