Agenda

- 1. Interpreting coefficients
- 2. Comparing models
- 3. Interpreting t- and p-values

Interpreting coefficients Our general recipe for interpretation of a regression coefficient is, "For a 1-unit increase in **X**, we would expect to see a [blah]-unit increase in **Y**." Ideally, you should substitute in something more specific for each of the bolded areas. The units of your response and explanatory variables, the quantities the variables represent, whether the coefficient on the explanatory variable is negative or positive, etc.

- 1. How does the recipe change for a categorical variable?
- 2. How does the recipe change for multiple regression?

```
require(openintro)
## Warning in library(package, lib.loc = lib.loc, character.only = TRUE, logical.return
= TRUE, : there is no package called 'openintro'
m1 <- lm(math~read+write+socst, data=hsb2)
## Error in is.data.frame(data): object 'hsb2' not found
summary(m1)
##
## Call:
## lm(formula = Price ~ Service, data = NYC)
## Residuals:
      Min 1Q Median 3Q
##
## -17.6646 -4.7540 -0.2093 4.3368 26.2460
##
## Coefficients:
    Estimate Std. Error t value Pr(>|t|)
## (Intercept) -11.9778 5.1093 -2.344 0.0202 *
## Service 2.8184 0.2618 10.764 <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
## Residual standard error: 7.153 on 166 degrees of freedom
## Multiple R-squared: 0.4111, Adjusted R-squared: 0.4075
## F-statistic: 115.9 on 1 and 166 DF, p-value: < 2.2e-16
```

- 1. How would you interpret the coefficient on read?
- 2. How would you interpret the coefficient on write?
- 3. How would you interpret the coefficient on socst? Does it make sense?

```
m2 <- lm(math~read+write+socst+gender, data=hsb2)</pre>
## Error in is.data.frame(data): object 'hsb2' not found
summary(m2)
##
## Call:
## lm(formula = log10(hwy) ~ displ, data = vehicles)
## Residuals:
                1Q Median
## -0.37785 -0.04395 -0.00016 0.04626 0.29171
##
## Coefficients:
               Estimate Std. Error t value Pr(>|t|)
## (Intercept) 1.5515065 0.0010471 1482 <2e-16 ***
## displ -0.0578847 0.0002894
                                      -200
                                             <2e-16 ***
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.07191 on 33383 degrees of freedom
## (57 observations deleted due to missingness)
## Multiple R-squared: 0.5451, Adjusted R-squared: 0.545
## F-statistic: 4e+04 on 1 and 33383 DF, p-value: < 2.2e-16
```

1. How would you interpret the coefficient on gendermale?

```
m3 <- lm(math~read+write+ses, data=hsb2)
## Error in is.data.frame(data): object 'hsb2' not found
summary(m3)
##
## Call:
## lm(formula = Price ~ Food + Service, data = NYC)
## Residuals:
##
      Min 1Q Median
                               3Q
                                        Max
## -16.1333 -4.7053 0.4169 3.5992 27.0728
##
## Coefficients:
##
             Estimate Std. Error t value Pr(>|t|)
## (Intercept) -21.1586 5.6651 -3.735 0.000258 ***
## Food
              1.4954
                        0.4462 3.351 0.000997 ***
                         0.4185 4.072 7.22e-05 ***
## Service
               1.7041
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6.942 on 165 degrees of freedom
## Multiple R-squared: 0.4486, Adjusted R-squared: 0.4419
## F-statistic: 67.12 on 2 and 165 DF, p-value: < 2.2e-16
```

- 1. How would you interpret the coefficient on sesmiddle?
- 2. How would you interpret the coefficient on seshigh?

Comparing Models We can't use multiple R^2 to compare models, because it will increase no matter what additional variables we add. For example, compare the previous model with the following:

```
m4 <- lm(math~read+write+ses+rnorm(200), data=hsb2)
## Error in is.data.frame(data): object 'hsb2' not found
summary(m4)
## Error in summary(m4): object 'm4' not found</pre>
```

But, R_{adj}^2 allows us to make comparisons. With that in mind, which is the best model we have seen so far?

Assessing conditions Assessing the LINE conditions is the same for multiple regression as it was for simple linear regression. However, the residual vs. fitted plot becomes even more useful.

```
par(mfrow=c(2,2))
plot(m3, which=1)
plot(m3, which=2)
plot(m3, which=3)
hist(m3$residuals)
```

