SUMMER 2025



INTRODUCTION TO STATISTICAL MODELING

Center for Biomedical Research Support

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Access materials

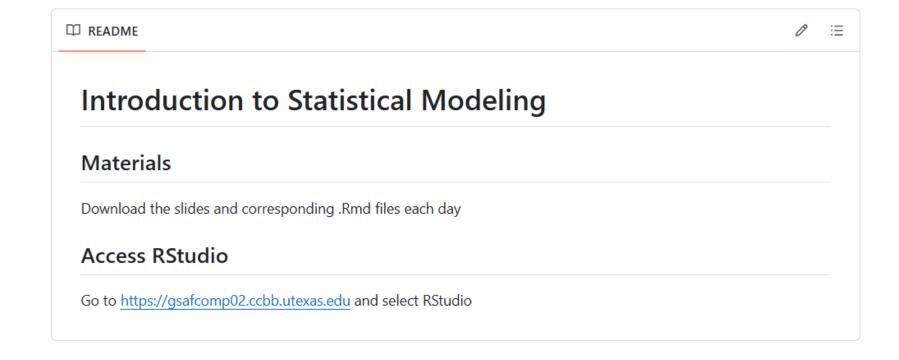




Layla Guyot laylaguyot

Statistics and Data Science enthusiast: teacher and researcher in education, focusing on bridging the gap between academia and industry.

https://github.com/laylaguyot/ CBRS Intro Statistical Modeling



Day 1 Exploring Data

- Study design and variables
- Descriptive statistics and visualizations
- Introduction to hypothesis testing

Day 2 Making Inferences

- Probability, random variables, and common probability distributions
- Sampling distributions and Central Limit Theorem
- Confidence intervals, t-tests, ANOVA, and Chi-square tests

Day 3 Linear Regression

- Simple Linear Regression
- Multiple Regression with different types of predictors
- Model assumptions, evaluation, and comparisons

Day 4 Logistic Regression

- Odds
- Logistic Regression
- Model evaluation with ROC curves or confusion matrix

Day 5 Model Building

- Underfitting, overfitting, and cross-validation
- Regularization with Lasso and Ridge
- Missing data

Summary of all tests:

Name of test	Variables involved	Hypotheses	Test statistic	df	Assumptions*	Effect size
One-sample t-test	numeric response	H_0 : $\mu = \mu_0$ H_A : $\mu \neq \mu_0$	$t = \frac{\bar{X} - \mu_0}{SE}$	n-1	✓ normal	$\bar{X} \pm t^* \cdot SE$
Independent t-test	numeric response binary predictor	$H_0: \mu_1 = \mu_2$ $H_A: \mu_1 \neq \mu_2$	$t = \frac{\bar{X}_1 - \bar{X}_2}{SE}$	$n_1 + n_2 - 1$	✓ normal✓ equal variance	$ar{X}_1 - ar{X}_2 \pm t^* \cdot SE$
ANOVA	numeric response categorical predictor	H_0 : $\mu_1 = \mu_2 = \cdots$ H_A : not all equal	$F = \frac{MS_{group}}{MS_{error}}$	$df_{group} = k - 1$ $df_{error} = n - k$	✓ normal✓ equal variance	Post Hoc Model fit <i>R</i> ²
Chi2 Goodness- of-Fit	categorical response	H_0 : distrib is H_A : distrib is not	$\chi^2 = \sum \frac{(obs - exp)^2}{exp}$	# cat — 1	✓ sample size	percentages
Chi2 Test of Independence	categorical response categorical predictor	H_0 : independent H_A : not independent	$\chi^2 = \sum \frac{(obs - exp)^2}{exp}$	$(\# cat_1 - 1)$ $\cdot (\# cat_2 - 1)$	✓ sample size	percentages

^{*}Random sample and Independent observations are common assumptions to all these tests

Checking in

Mentimeter

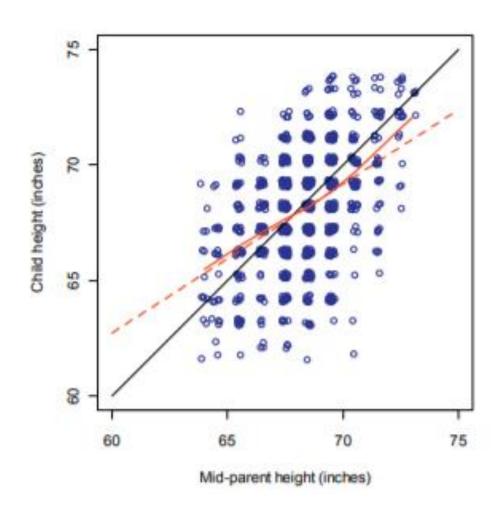
What is something you have learned (or re-discovered) during this workshop so far?



Linear Regression

Where does the term "regression" come from?

- Coined by Francis Galton in the late 1800s.
- Galton observed that tall parents tended to have shorter children (closer to the population average) and vice versa. He called this pattern "regression toward mediocrity", what we now call regression to the mean.
- Galton and Pearson used family height data to describe and model these trends.



Mhile foundational to modern statistics for the least squares linear model, Galton and Pearson's work is inseparable from their **promotion of eugenics**.

Population model:

$$Y = \beta_0 + \beta_1 X_1 + \varepsilon$$

• Estimated regression function:

$$\widehat{Y} = b_0 + b_1 X_1$$

• Residuals:

$$e_i = Y_i - \widehat{Y}_i$$

error/residual

What should we test about to check for a "significant" linear relationship?

How to determine the line to model our data?

Least Squares Regression Line

$$\widehat{Y} = b_0 + b_1 X$$
 with $b_1 = r \frac{s_Y}{s_X}$ $b_0 = \overline{Y} - b_1 \overline{X}$



Besides random sample, there are 4 assumptions for the SLR model:

Linearity: the mean response is a linear function of X_i

Independent observations: the errors, ϵ_i , are independent

Normality of residuals: the errors, ϵ_i , are normally distributed

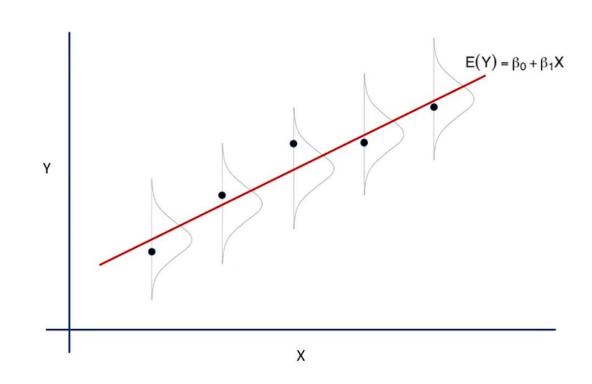
Equal variances: the errors, ϵ_i , have equal variances (σ^2)

Besides random sample, there are 4 assumptions for the SLR model:

Linearity

Independent

Normality

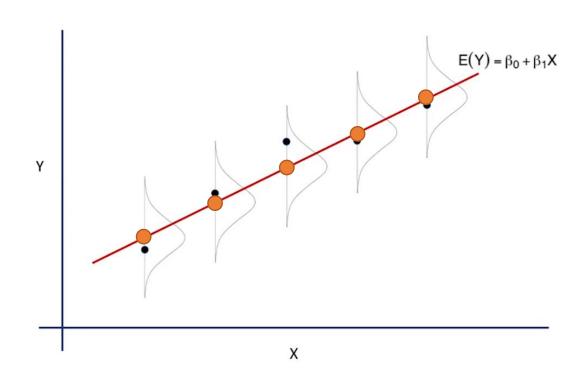


Besides random sample, there are 4 assumptions for the SLR model:

→ **L**inearity

Independent

Normality

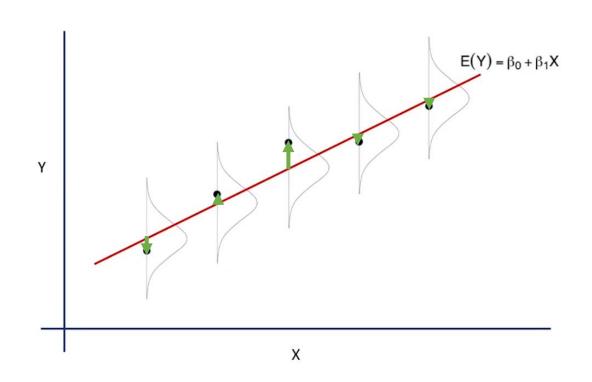


Besides random sample, there are 4 assumptions for the SLR model:

Linearity

→ Independent

Normality

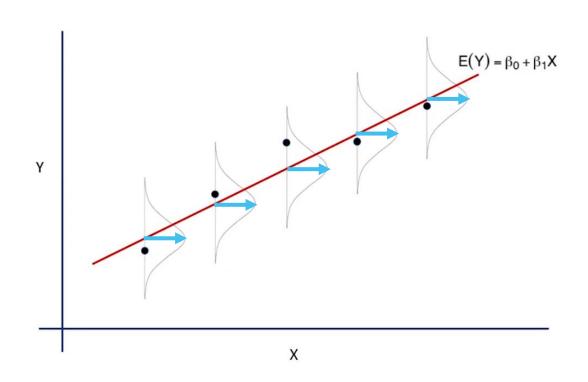


Besides random sample, there are 4 assumptions for the SLR model:

Linearity

Independent

→ Normality

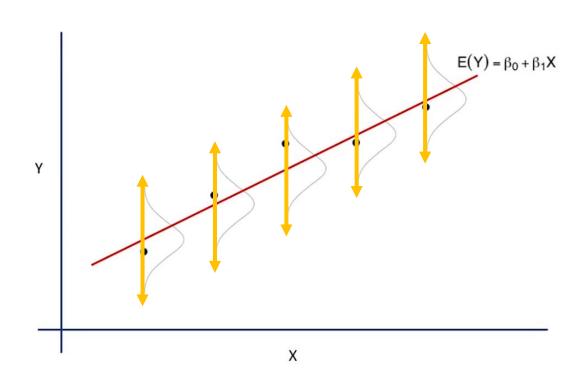


Besides random sample, there are 4 assumptions for the SLR model:

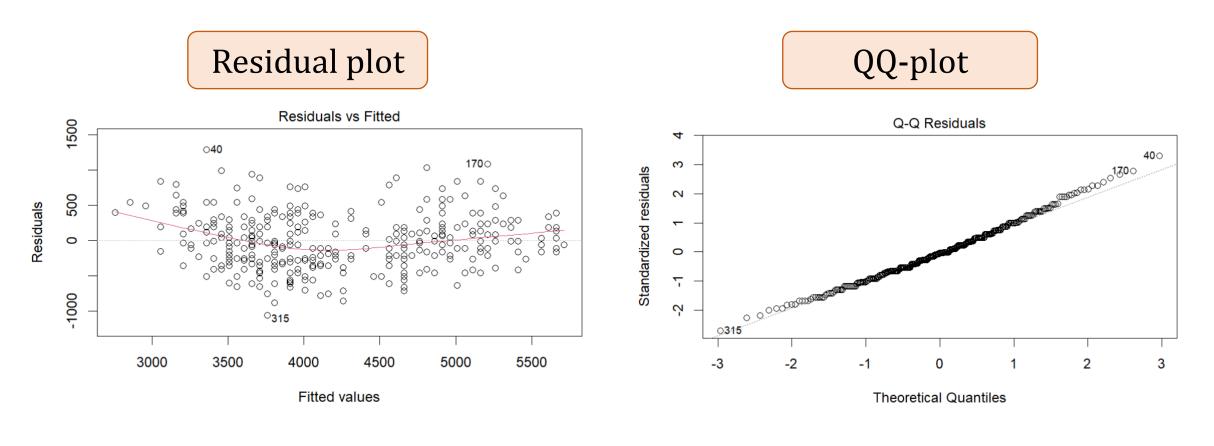
Linearity

Independent

Normality



Checking for assumptions:



Linearity, Equal variance

Normality

Checking for assumptions:

Residual plot

✓ Nonlinearity
Look for clear nonlinear patterns in
the residuals plot

✓ Unequal error variance Look for unequal distances from y=0

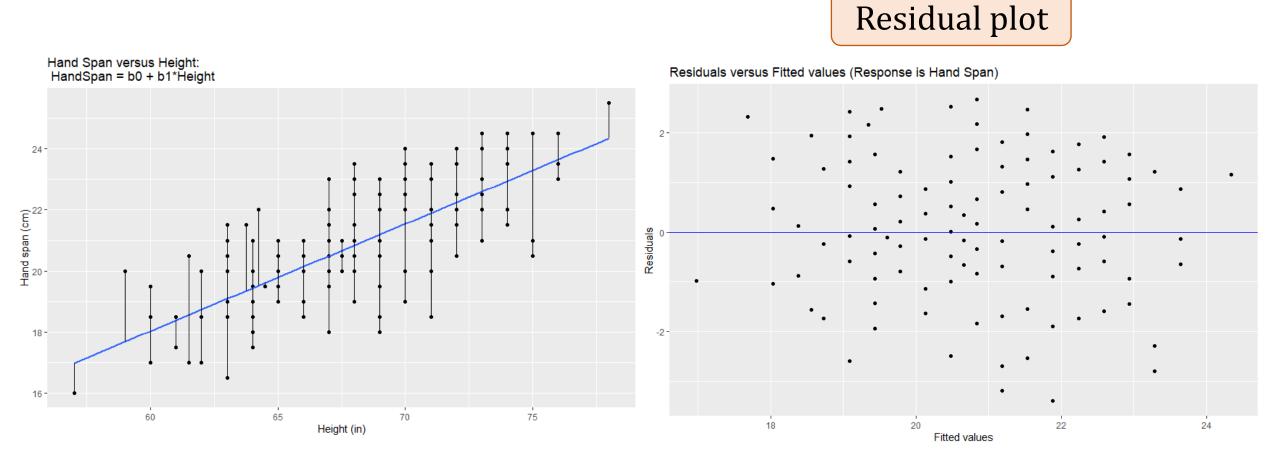
✓ Outliers

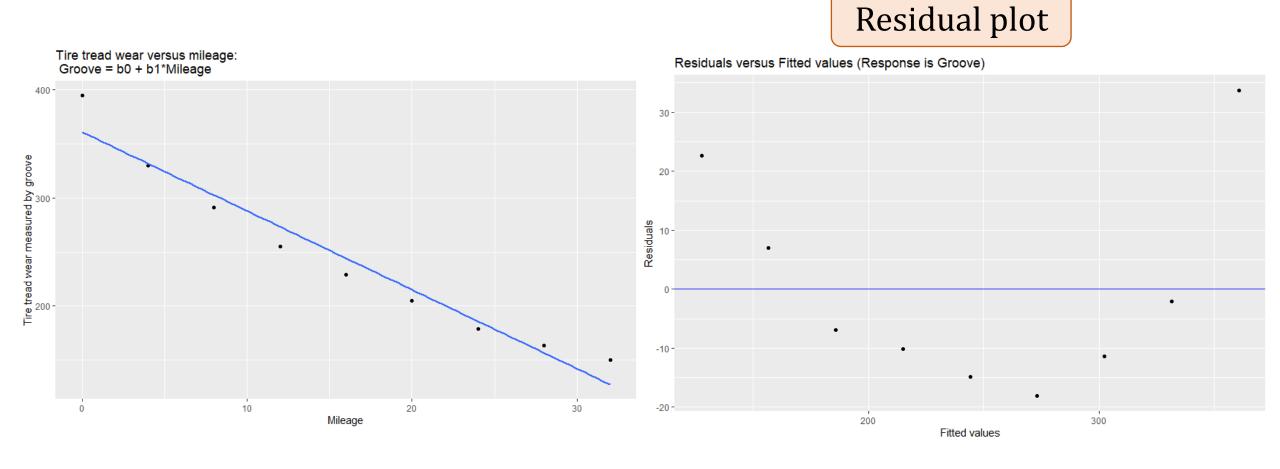
Look for extreme values in the residuals plot

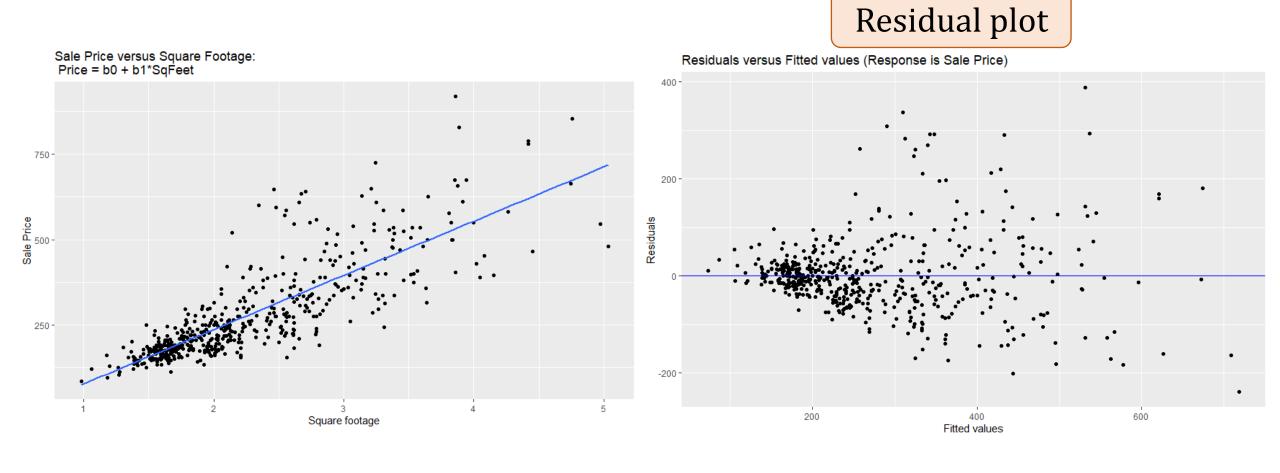
QQ-plot

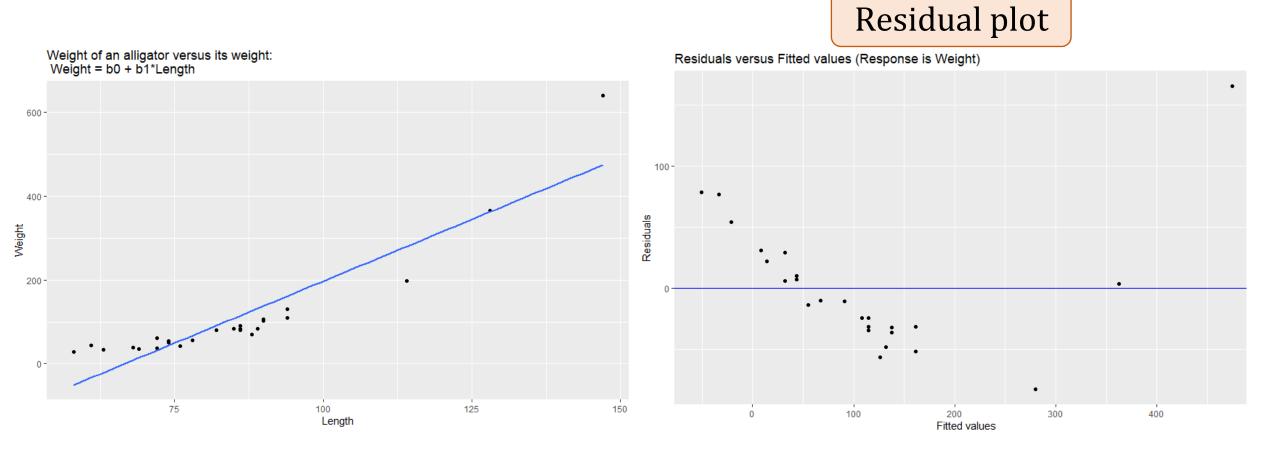
✓ Not normally Look for deviation from the normality:

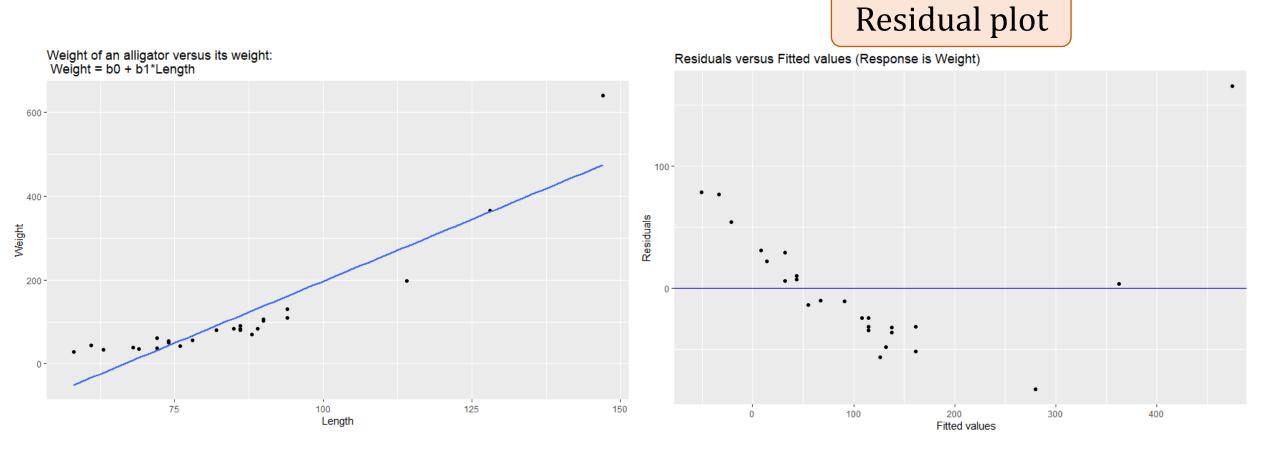
We compare the observed sample percentiles to the theoretical percentiles of the normal distribution. The normal probability plot should be approximately linear.

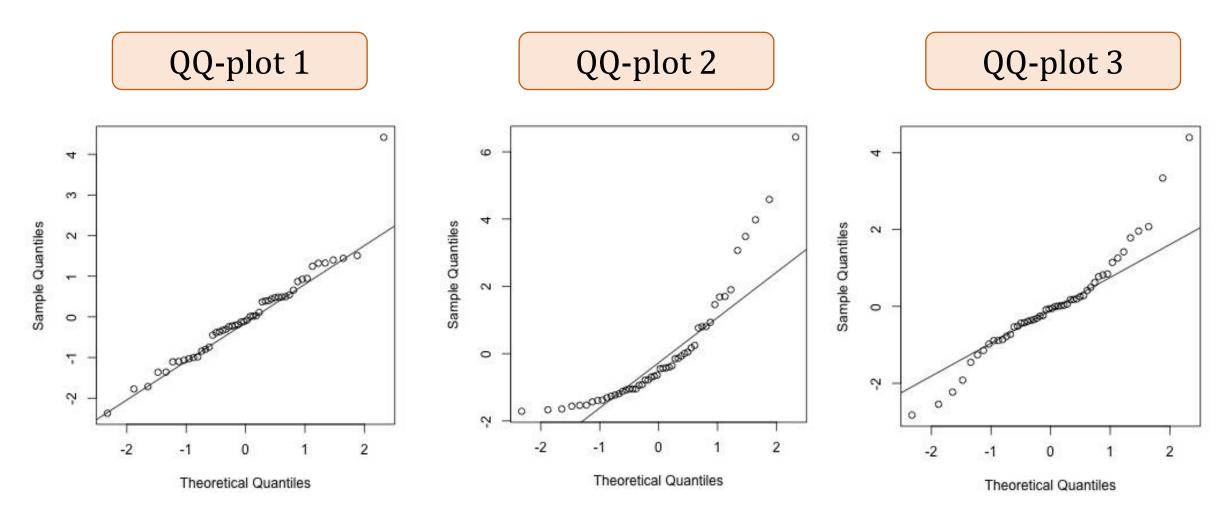


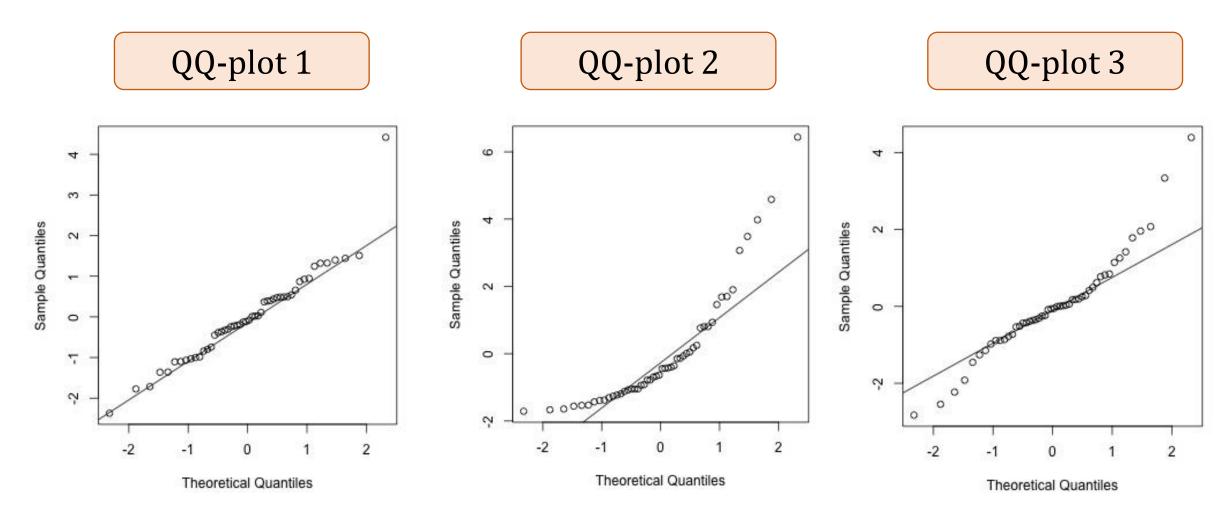














Comparing population slope to 0

1. State your hypotheses

$$H_0: \beta_1 = 0$$
$$H_A: \beta_1 \neq 0$$

$$H_A: \beta_1 \neq 0$$

Comparing population slope to 0

- 1. State your hypotheses
- 2. Calculate the test statistic *t* (based on sample data)

$$t = \frac{b_1}{SE_{b_1}}$$

We will explain this standard error later

Comparing population slope to 0

- 1. State your hypotheses
- 2. Calculate the test statistic *t* (based on sample data)
- 3. Compare test statistic to null distribution (calculate *p*-value)
- 4. Make a conclusion in context, reporting the appropriate statistics (t, df, p-value).

$$df = n - 2$$

When reporting results of a significant test, we should report a measure of the effect size with a **confidence interval** of the **population slope**:

$$b_1 \pm t_{df}^* \cdot SE_{b_1}$$

We will explain this standard error later

Inference for predicted values

Example:

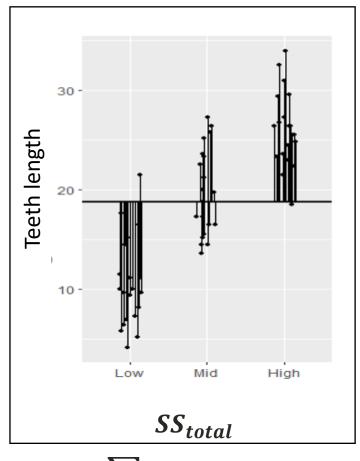
- 1. What is the mean hand span, μ , of <u>all</u> US adults?
 - Confidence interval for the mean response
- 2. What is the hand span, *Y*, of an **individual** US adult?
 - Prediction interval for a predicted response

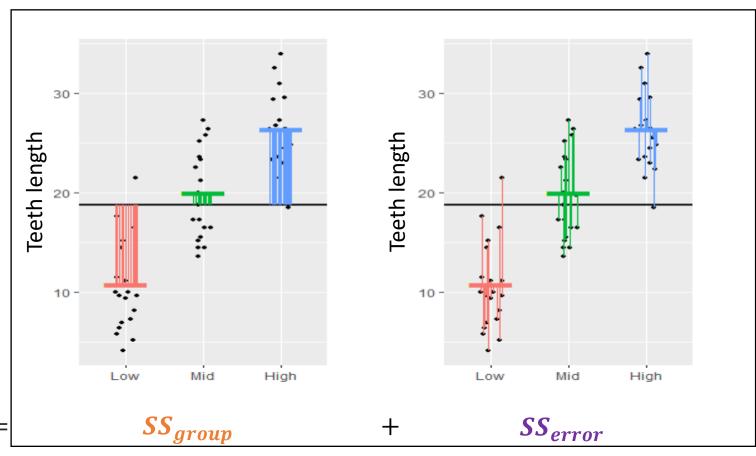
To evaluate the performance of a linear regression model, we can consider:

- R^2 : measures the percentage of variation in the response variable that can be explained by the predictor variable.
 - The higher the \mathbb{R}^2 , the better the model is.
- Mean Squared Error (MSE): measures the average squared residuals. The lower the MSE, the better the model is.
- Root Mean Squared Error (RMSE): measures how far apart the predicted values are from the observed values in a dataset, on average.
 - The lower the RMSE, the better the model is.



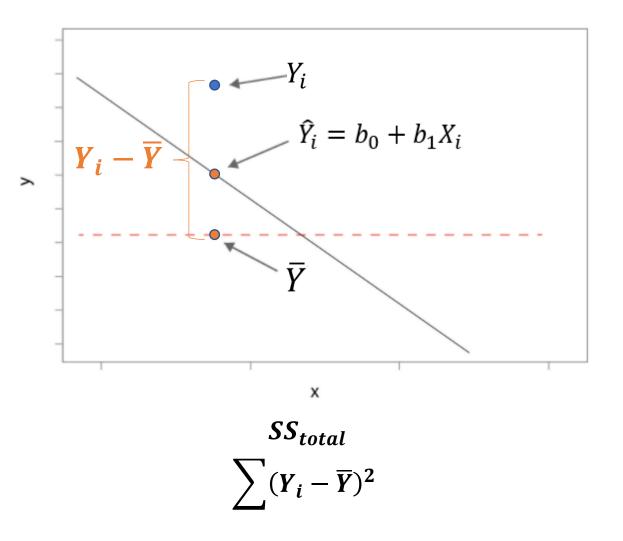
Decomposing variation in ANOVA



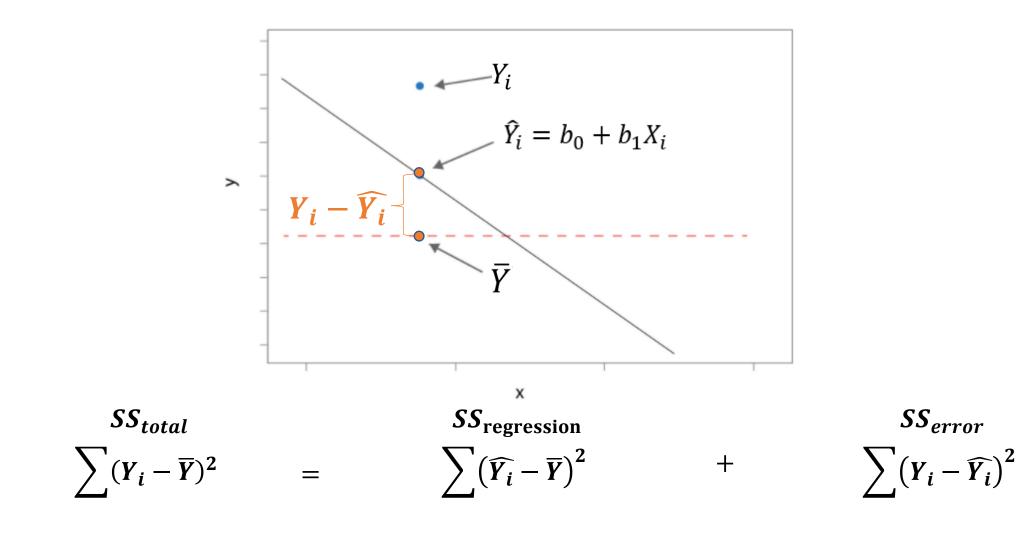


$$\sum (X_{ik} - \overline{X})^2 \qquad = \qquad \sum n_k (\overline{X}_k - \overline{X})^2 \qquad + \qquad \sum \sum (X_{ik} - \overline{X})^2$$

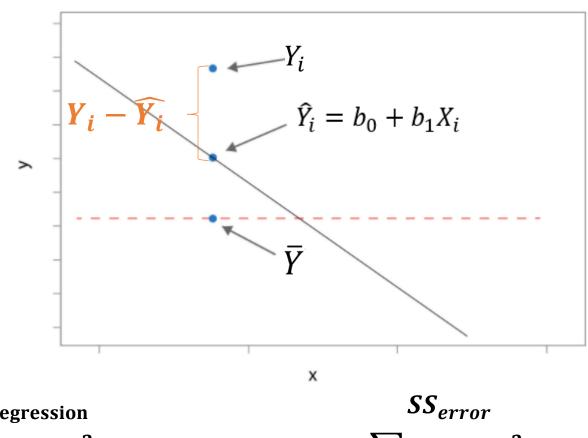
Decomposing variation in Simple Linear Regression



Decomposing variation in Simple Linear Regression



Decomposing variation in Simple Linear Regression



$$SS_{total} = \sum (Y_i - \overline{Y})^2 =$$

$$SS_{\text{regression}}$$

$$\sum (\widehat{Y_i} - \overline{Y})^2$$

$$\sum (Y_i - \widehat{Y}_i)^2$$

Connection with ANOVA

Inference with ANOVA

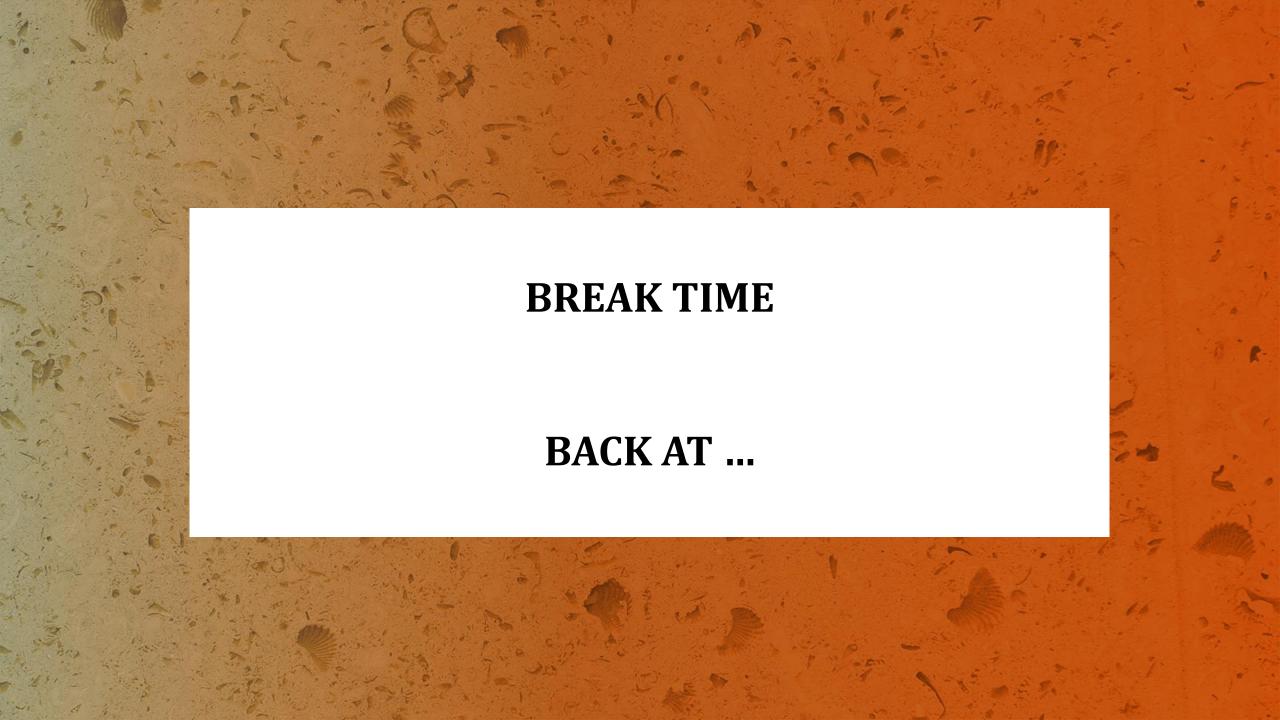
$$H_0$$
: $\beta_1 = 0$
 H_A : $\beta_1 \neq 0$

$$H_A$$
: $\beta_1 \neq 0$

$$F = rac{MS_{regression}}{MS_{error}}$$
 with $MS_{regression} = rac{SS_{regression}}{1}$

$$MS_{error} = \frac{SS_{error}}{n-2}$$





Population model:

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_p X_p + \varepsilon$$

• Estimated regression function:

$$\widehat{Y} = b_0 + b_1 X_1 + b_2 X_2 + \dots + b_p X_p$$

• Residuals:

$$e_i = Y_i - \widehat{Y}_i$$

Example: On a sample of 38 college students, the following variables were collected:

Y: Performance IQ scores (PIQ) of the revised Wechsler Adult Intelligence Scale

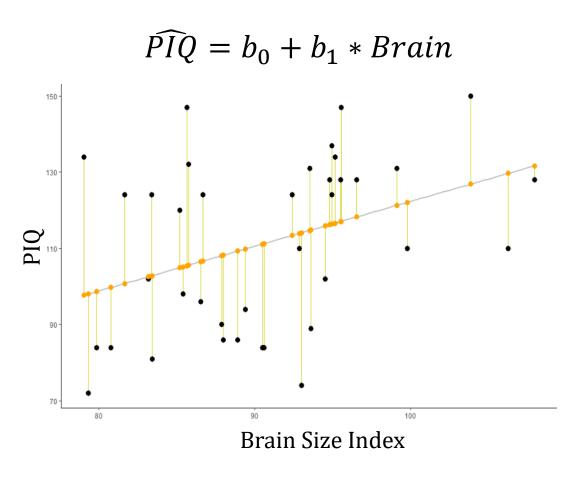
 X_1 : Brain size index based on MRI scans

 X_2 : Height (in inches)

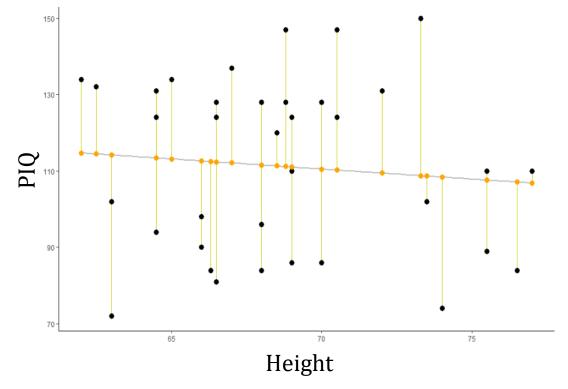
 X_3 : Weight (in pounds)

Can a student's intelligence be predicted by brain size and height?

Can a person's intelligence be predicted by brain size? by height?



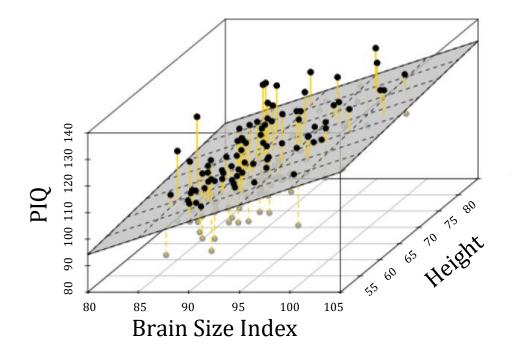
$$\widehat{PIQ} = b_0 + b_1 * Height$$



Can a person's intelligence be predicted by brain size and height?

$$\widehat{PIQ} = b_0 + b_1 * Brain + b_2 * Height$$

For 2 predictors (brain size index and height), the equation technically yields a plane:



Can a person's intelligence be predicted by brain size and height?

$$\widehat{PIQ} = b_0 + b_1 * Brain + b_2 * Height$$

Interpret the values of the slopes

As brain size index increases by 1, PIQ increases by 2.1 on average, while holding height constant.

As height increases by 1 inch, PIQ decreases by 2.7 on average, while holding brain size index constant.

```
> my_model <- lm(PIQ ~ Brain + Height, data = iqsize)
> summary(my_model)
Call:
lm(formula = PIQ ~ Brain + Height, data = igsize)
Residuals:
   Min
            1Q Median 3Q
-32.750 -12.090 -3.841 14.174 51.690
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 111.2757 55.8673 1.992 0.054243 .
Brain 2.0606 0.5466 3.770 0.000604 ***
Height -2.7299 0.9932 -2.749 0.009399 **
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Residual standard error: 19.51 on 35 degrees of freedom
Multiple R-squared: 0.2949, Adjusted R-squared: 0.2546
F-statistic: 7.321 on 2 and 35 DF, p-value: 0.002208
```

Comparing population slope to 0

for each slope

1. State your hypotheses

$$H_0: \beta_i = 0$$

$$H_A: \beta_i \neq 0$$

- 2. Calculate the test statistic $t = \frac{b_i}{SE_{b_i}}$
- 3. Compare test statistic to null distribution (calculate *p*-value)
- 3. Make a conclusion in context, reporting the appropriate statistics (t, df, p-value).

$$df = n - p - 1$$

Can a person's intelligence be predicted by brain size and height?

$$\widehat{PIQ} = b_0 + b_1 * Brain + b_2 * Height$$

Interpret the values of the slopes

Brain size index is a significant predictor of PIQ while holding height constant (t = 3.77, df = 35, p = 0.0006).

Height is a significant predictor of PIQ while holding brain size index constant (t = -2.75, df = 35, p = 0.009).

```
> my_model <- lm(PIQ ~ Brain + Height, data = igsize)
> summary(my_model)
Call:
lm(formula = PIQ ~ Brain + Height, data = iqsize)
Residuals:
           1Q Median 3Q
-32.750 -12.090 -3.841 14.174 51.690
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 111.2757 55.8673 1.992 0.054243 .
Brain 2.0606 0.5466 3.770 0.000604 ***
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Independent observations: the errors, ϵ_i , are independent

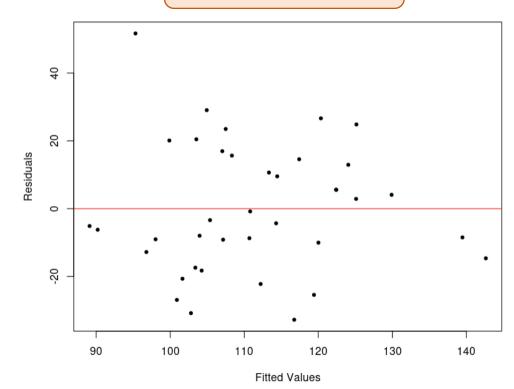
Normality of residuals: the errors, ϵ_i , are normally distributed

Equal variances: the errors, ϵ_i , have equal variances (σ^2)

Same assumptions than for the Simple Linear Regression model!

Are the assumptions met?

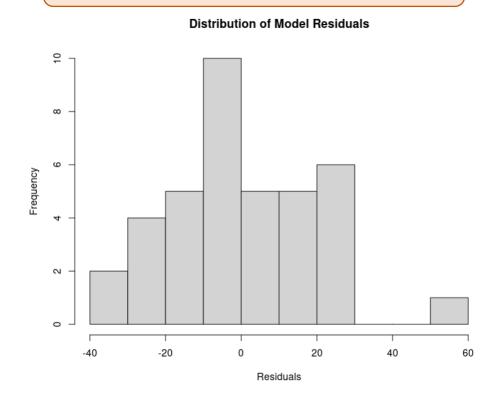




Linearity

Equal variance

Histogram of the residuals



Normality

Shapiro-Wilk normality test

data: my_model\$residuals
W = 0.976, p-value = 0.5764

Can a person's intelligence be predicted by brain size and height?

$$\widehat{PIQ} = b_0 + b_1 * Brain + b_2 * Height$$

Anything else we should check?

```
> my_model <- lm(PIQ ~ Brain + Height, data = igsize)
> summary(my_model)
Call:
lm(formula = PIQ ~ Brain + Height, data = iqsize)
Residuals:
   Min 10 Median 30
                                 Max
-32,750 -12,090 -3,841 14,174 51,690
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 111.2757 55.8673 1.992 0.054243 .
Brain 2.0606 0.5466 3.770 0.000604 ***
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```

Can a person's intelligence be predicted by brain size and height?

$$\widehat{PIQ} = b_0 + b_1 * Brain + b_2 * Height$$

What could be a potential issue if we had many predictors? What would happen to R²?

```
> my_model <- lm(PIQ ~ Brain + Height, data = iqsize)
> summary(my_model)
Call:
lm(formula = PIQ ~ Brain + Height, data = iqsize)
Residuals:
   Min 1Q Median 3Q
-32.750 -12.090 -3.841 14.174 51.690
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 111.2757 55.8673 1.992 0.054243 .
Brain 2.0606 0.5466 3.770 0.000604 ***
Height -2.7299 0.9932 -2.749 0.009399 **
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F-statistic: 7.321 on 2 and 35 DF, p-value: 0.002208
```

Can a person's intelligence be predicted by brain size and height?

Model with Brain size, Height

F-statistic: 7.321 on 2 and 35 DF, p-value: 0.002208

```
> my_model <- lm(PIQ ~ Brain + Height, data = iqsize)
> summary(my_model)
                                                       better?
Call:
lm(formula = PIQ ~ Brain + Height, data = igsize)
Residuals:
   Min
            10 Median
                           30
                                  Max
-32.750 -12.090 -3.841 14.174 51.690
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) 111.2757 55.8673 1.992 0.054243 .
         2.0606 0.5466 3.770 0.000604 ***
Brain
Height
            -2.7299 0.9932 -2.749 0.009399 **
Signif, codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Residual standard error: 19.51 on 35 degrees of freedom
Multiple R-squared: 0.2949, Adjusted R-squared: 0.2546
```



```
» my_model <- lm(PIQ ~ Brain + Height + Weight, data = igsize)</p>
Call:
lm(formula = PIQ ~ Brain + Height + Weight, data = iqsize)
Residuals:
          1Q Median
  Min
                        30 Max
-32.74 -12.09 -3.84 14.17 51.69
Coefficients:
             Estimate Std. Error t value Pr(>|t|)
(Intercept) 1.114e+02 6.297e+01 1.768 0.085979 .
Brain
            2.060e+00 5.634e-01 3.657 0.000856 ***
Height
        -2.732e+00 1.229e+00 -2.222 0.033034 *
Weight
            5.599e-04 1.971e-01 0.003 0.997750
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 19.79 on 34 degrees of freedom
Multiple R-squared: 0.2949, Adjusted R-squared: 0.2327
F-statistic: 4.741 on 3 and 34 DF, p-value: 0.007215
```



On a random sample of 32 births, Daniel (1999) collected these three variables:

Y: Birth weight of baby (Wgt) in grams

 X_1 : Length of gestation (Gest) in weeks

 X_2 : Smoking status of mother (Smoke = 1 if yes or Smoke = 0 if no)

Does smoking during pregnancy affect birth weight, while accounting for the length of gestation?

Does smoking during pregnancy affect birth weight, while accounting for the length of gestation?

```
\widehat{Wgt} = -2389.573 + 143.1 \, Gest - 244.544 \, Smoke
```

Interpret the values of the slopes

As the length of gestation increases by 1 week, birth weight increases by 143.1 grams on average, while holding smoking status constant.

Birth weight decreases by 244.544 grams on average for babies born from smoking mothers compared to nonsmoking, while holding length of gestation constant.

```
> reg <- lm(Wgt~Gest+Smoke,birthsmokers)
> summary(reg)
call:
lm(formula = Wgt ~ Gest + Smoke, data = birthsmokers)
Residuals:
    Min
             1Q Median
-223,693 -92,063 -9,365 79,663 197,507
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) -2389.573 349.206 -6.843 1.63e-07 ***
Gest
        143.100 9.128 15.677 1.07e-15 ***
Smoke -244,544
                       41.982 -5.825 2.58e-06 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Residual standard error: 115.5 on 29 degrees of freedom
Multiple R-squared: 0.8964, Adjusted R-squared: 0.8892
F-statistic: 125.4 on 2 and 29 DF, p-value: 5.289e-15
```

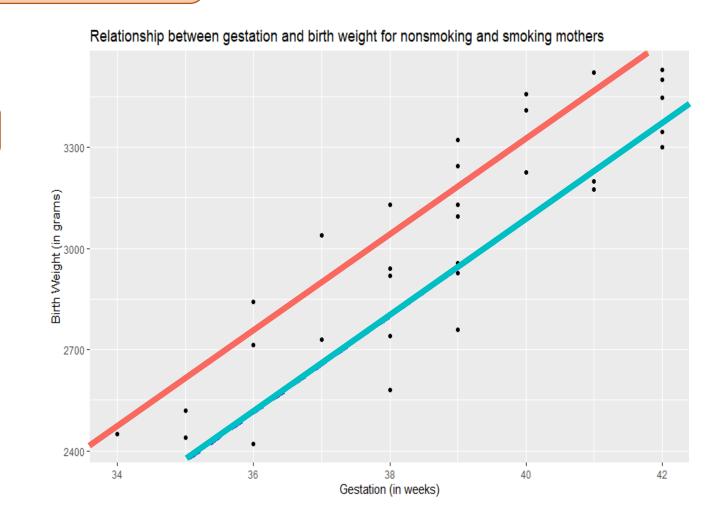
Does smoking during pregnancy affect birth weight, while accounting for the length of gestation?

$$\widehat{Wgt} = -2389.573 + 143.1 \, Gest - 244.544 \, Smoke$$

Interpret the values of the slopes

As the length of gestation increases by 1 week, birth weight increases by 143.1 grams on average, while holding smoking status constant.

Birth weight decreases by 244.544 grams on average for babies born from smoking mothers compared to nonsmoking, while holding length of gestation constant.



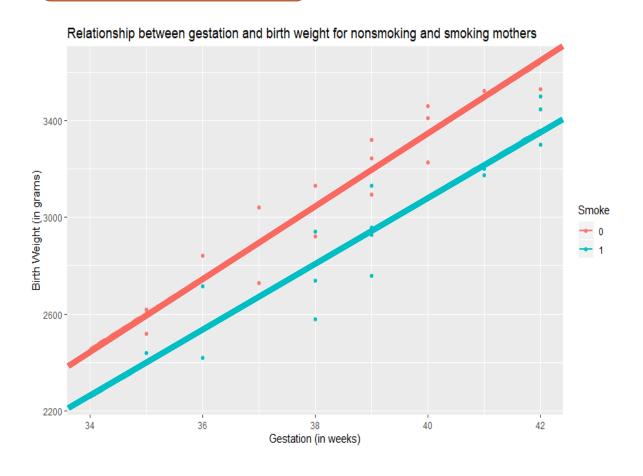
Interactions

- ➤ Does the effect of one predictor on the response depends on the levels of another predictor?
- ➤ Model with interaction effect for two predictors:

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_{1*2} (X_1 * X_2) + \varepsilon$$

Main effects: effects between each predictor and the response Interaction effect: a predictor might affect the response differently, depending on another predictor

Interactions



Does the effect of the gestation length on birth weight depend on smoking status?

OR

Does the effect of smoking on birth weight depend on the length of gestation?

Is there an interaction between the length of gestation and smoking status to predict birth weight?

```
\widehat{Wgt} = -2546.1 + 147.2 \ Gest + 71.6 \ Smoke - 8.2 \ Gest * Smoke
```

Interpret the slopes of the main effects.

As the length of gestation increases by 1 week, birth weight increases by 147.2 grams for <u>nonsmoking</u> mothers.

Birth weight is 71.6 grams more on average for babies born from smoking mothers compared to nonsmoking mothers for a length gestation of 0.

Does not make sense to interpret

```
> reg <- lm(Wgt~Gest*Smoke, data = birthsmokers)</p>
> summary(reg)
Call:
lm(formula = Wgt ~ Gest * Smoke, data = birthsmokers)
Residuals:
             1Q Median
    Min
                             30
                                     Max
-228.528 -89.560 0.273 83.629 184.529
Coefficients:
           Estimate Std. Error t value Pr(>|t|)
(Intercept) -2546.138
                      501,067 -5.081 2.22e-05
Gest
    147.207 13.120 11.220 7.15e-12 ***
Smoke 71.574 716.950 0.100
                                     0.921
Gest:Smoke -8.178
                    18.515 -0.442 0.662
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 117.2 on 28 degrees of freedom
Multiple R-squared: 0.8971, Adjusted R-squared: 0.8861
F-statistic: 81.37 on 3 and 28 DF, p-value: 6.144e-14
```

Interactions

> Centering numeric predictors to better interpret slopes:

 $Variable_c = Variable - mean(Variable)$

How do we interpret $Variable_c = 0$?

Is there an interaction between the length of gestation and smoking status to predict birth weight?

```
\widehat{Wgt} = 31144.3 + 147.2 \ Gest - 244.6 \ Smoke - 8.2 \ Gest * Smoke
```

Interpret the slopes of the main effects.

As the length of gestation increases by 1 week, birth weight increases by 147.2 grams for <u>nonsmoking</u> mothers.

Birth weight is 244.6 grams less on average for babies born from smoking mothers compared to nonsmoking mothers for an average gestation length.

```
> Gest_c <- birthsmokers$Gest-mean(birthsmokers$Gest)
> req <- lm(Wgt~Gest_c*Smoke, data = birthsmokers)</p>
> summary(reg)
Call:
lm(formula = Wgt ~ Gest_c * Smoke, data = birthsmokers)
Residuals:
    Min 1Q Median 3Q
                                     Max
-228.528 -89.560 0.273
                           83.629 184.529
Coefficients:
             Estimate Std. Error t value Pr(>|t|)
                         30.110 104.429 < 2e-16 ***
(Intercept)
             3144.329
Gest c
           147.207 13.120 11.220 7.15e-12 ***
Smoke: -244.563 42.577 -5.744 3.65e-06 ***
Gest_c:Smoke -8.178 18.515 -0.442
                                         0.662
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
Residual standard error: 117.2 on 28 degrees of freedom
Multiple R-squared: 0.8971, Adjusted R-squared: 0.8861
F-statistic: 81.37 on 3 and 28 DF, p-value: 6.144e-14
```

Is there an interaction between the length of gestation and smoking status to predict birth weight?

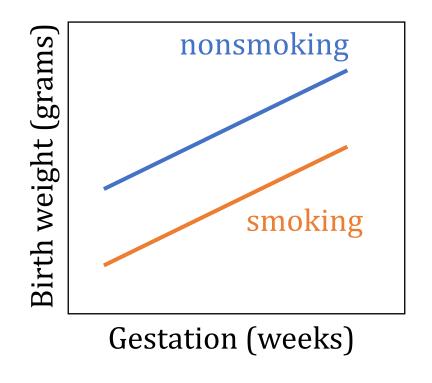
```
\widehat{Wgt} = 31144.3 + 147.2 \ Gest - 244.6 \ Smoke - 8.2 \ Gest * Smoke
```

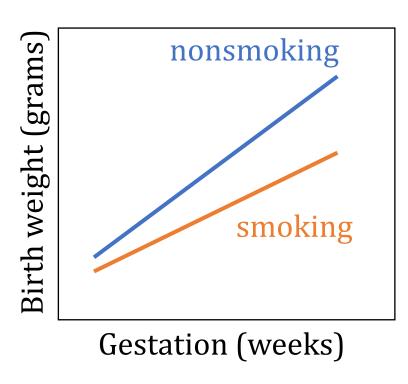
Interpret the slope for interaction effect.

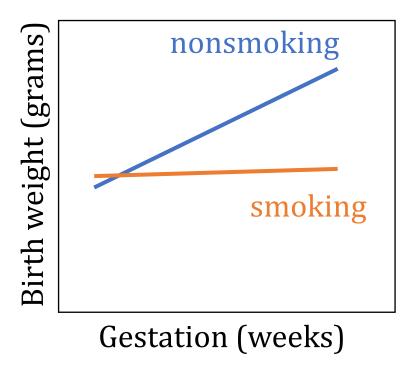
The effect of the length of gestation on the birth weight decreases by 8.178 grams on average for smoking mothers compared to nonsmoking mothers.

```
> Gest_c <- birthsmokers$Gest-mean(birthsmokers$Gest)
> reg <- lm(Wgt~Gest_c*Smoke, data = birthsmokers)
> summary(reg)
Call:
lm(formula = Wat ~ Gest c * Smoke. data = birthsmokers)
Residuals:
    Min 1Q Median 3Q
                                     Max
-228.528 -89.560 0.273 83.629 184.529
Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 3144.329 30.110 104.429 < 2e-16 ***
Gest c 147.207 13.120 11.220 7.15e-12 ***
Smoke: -244.563 42.577 -5.744 3.65e-06 ***
Gest_c:Smoke -8.178 18.515 -0.442
                                         0.662
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '1
Residual standard error: 117.2 on 28 degrees of freedom
Multiple R-squared: 0.8971, Adjusted R-squared: 0.8861
F-statistic: 81.37 on 3 and 28 DF, p-value: 6.144e-14
```

In which of these graphs do you see an interaction between gestation and smoking status?









A case study: Predicting course ratings

<u>Hamermesh & Parker (2004) Beauty in the classroom:</u> instructors' pulchritude and putative pedagogical productivity. <u>Economics of Education Review.</u>



Collected data at the University of Texas at Austin:

- Evaluations at the end of the semester for the academic years 2000-2002: 463 courses.
- Instructors at all levels, with pictures on their department's websites: 94 professors.
- Beauty judgements were made by 6 students who had not attended the classes and were not aware of the course evaluations: combined into a beauty score (out of 10).

What are the sampling units? What is the population?

Preliminary Analysis: Descriptive statistics

Table 1 page 3

Descriptive statistics, courses, instructors and evaluations

Variable	All
Course evaluation	4.022 (0.525)
Instructor evaluation	4.217 (0.540)
Number of students	55.18 (75.07)
Percent evaluating	74.43
Female	0.359
Minority	0.099
Non-native English	0.037
Tenure track	0.851
Lower division	0.339
One credit	0.029
Number of courses	463
Number of faculty	94

Look at the list of variables: Which one can be considered as the response? Which ones as potential predictors?

Note: Means with standard deviations in parentheses. All statistics except for those describing the number of students, the percent evaluating the instructor and the lower-upper division distinction are weighted by the number of students completing the course evaluation forms.

Preliminary Analysis: Descriptive statistics

Table 1 page 3 Descriptive statistics, courses, instructors and evaluations		
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Look at the descriptive statistics. Interpret the highlighted values.

Note: Means with standard deviations in parentheses. All statistics except for those describing the number of students, the percent evaluating the instructor and the lower-upper division distinction are weighted by the number of students completing the course evaluation forms.

Preliminary Analysis: Comparisons

Table 2 page 3

Beauty evaluations, individual and composite

	Average	Standard deviation	Standardized	
			Minimum	Maximum
Individual ratings:				
Male, upper division-1	4.43	2.18	-1.57	2.10
Male, upper division—2	4.87	1.65	-2.34	2.50
Female, upper division-1	5.18	2.05	-2.03	1.84
Female, upper division-2	5.39	2.10	-2.10	2.20
Male, lower division	3.53	1.70	-1.49	2.04
Female, lower division	4.14	1.88	-1.67	2.05
Composite standardized rating				
_	0	0.83	-1.54	1.88

Compare the highlighted values.

Preliminary Analysis: Comparisons

Table 2 page 3

Beauty evaluations, individual and composite

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Female, lower division	4.14	1.88	-1.67	2.05
Composite standardized rating				
	0	0.83	-1.54	1.88

If we wanted to compare the highlighted values with a test, what test shall we conduct?

Preliminary Analysis: Comparisons

Table 2 page 3
Beauty evaluations, individual and composite

	Average	Standard deviation	Standardized	
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Composite standardized rating				
-	0	0.83	-1.54	1.88

What did the researcher do when they "standardized" the beauty scores?

Predicting course ratings: Multiple Regression

Table 3 page 4
Weighted least-squares estimates of the determinants of class ratings

Variable	All
Composite standardized beauty	0.275 (0.059)
Female	-0.239 (0.085)
Minority	-0.249 (0.112)
Non-native English	-0.253 (0.134)
Tenure track	-0.136 (0.094)
Lower division	-0.046 (0.111)
One-credit course	0.687 (0.166)
R^2	.279
N courses	463
N faculty	94

Look at the estimates for each predictor. Which predictors have a positive effect on the class ratings (controlling for other variables)?

Note: Robust standard errors in parentheses here and in Table 4.

On page 4, the authors wrote: "The striking fact from the estimates [...] is the statistical significance of the composite standardized beauty measure".

Does Table 3 show which variables are significant?

Predicting course ratings



Based on the RStudio output, which predictors have a significant effect on class ratings?

```
Coefficients:
               Estimate Std. Error t value
                                              Pr(>|t|)
              4.22314 0.06512 64.856 < 0.000000000000000000
(Intercept)
genderfemale
           -0.23899 0.04586 -5.212
                                            0.000000284 ***
minorityyes -0.24894 0.07999 -3.112
                                               0.00197 **
nativeno
        -0.25271 0.11985 -2.109
                                               0.03553 *
tenureyes -0.13592 0.06247 -2.176
                                              0.03007 *
divisionlower -0.04589 0.04380 -1.048
                                               0.29523
creditssingle 0.68651
                        0.13685 5.016
                                           0.000000757 ***
            0 (**** 0.001 (*** 0.01 (** 0.05 (... 0.1 (... 1
Signif. codes:
Residual standard error: 2.719 on 455 degrees of freedom
Multiple R-squared: 0.2788, Adjusted R-squared: 0.2677
F-statistic: 25.13 on 7 and 455 DF, p-value: < 0.00000000000000022
```

Predicting course ratings



The authors acknowledged some issues that came up during their study:

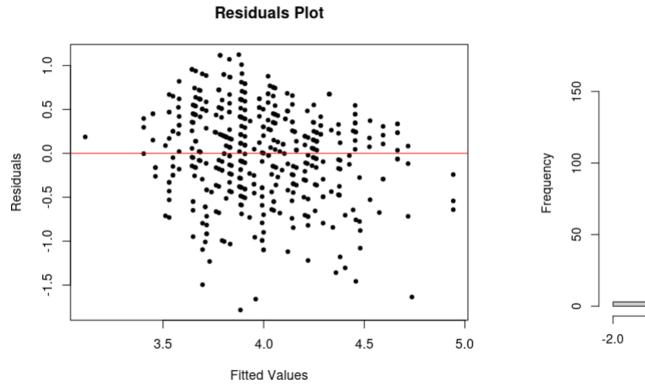
- "the most serious potential problem may result from a type of sample selectivity"
- "our measure of beauty may merely be a proxy"
- "whether higher instructional ratings mean that the faculty member is a better teacher"

Why is important for the authors to acknowledge these issues?

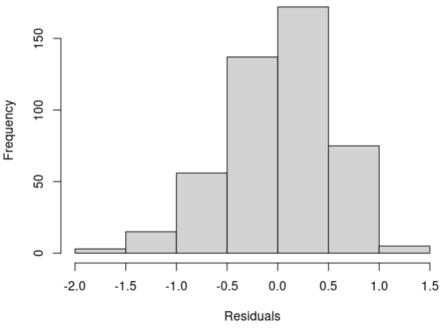
What other potential issues should they have also addressed?

Predicting course ratings





Distribution of Model Residuals



Check the assumptions

Predicting course ratings



Table 3 page 4

Weighted least-squares estimates of the determinants of class ratings

Variable	All
Composite standardized beauty	0.275 (0.059)
Female	-0.239 (0.085)
Minority	-0.249 (0.112)
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Tenure track	-0.136 (0.094)
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One-credit course	0.687 (0.166)
R^2	.279
N courses	463
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Note: Robust standard errors in parentheses here and in Table 4.

What's the model fit? What does it mean?

Next

Day 4 Logistic Regression

- Odds
- Logistic Regression
- Model evaluation with ROC curves or confusion matrix

Any questions? comments?

