

Lesson 14: MLR Model Diagnostics

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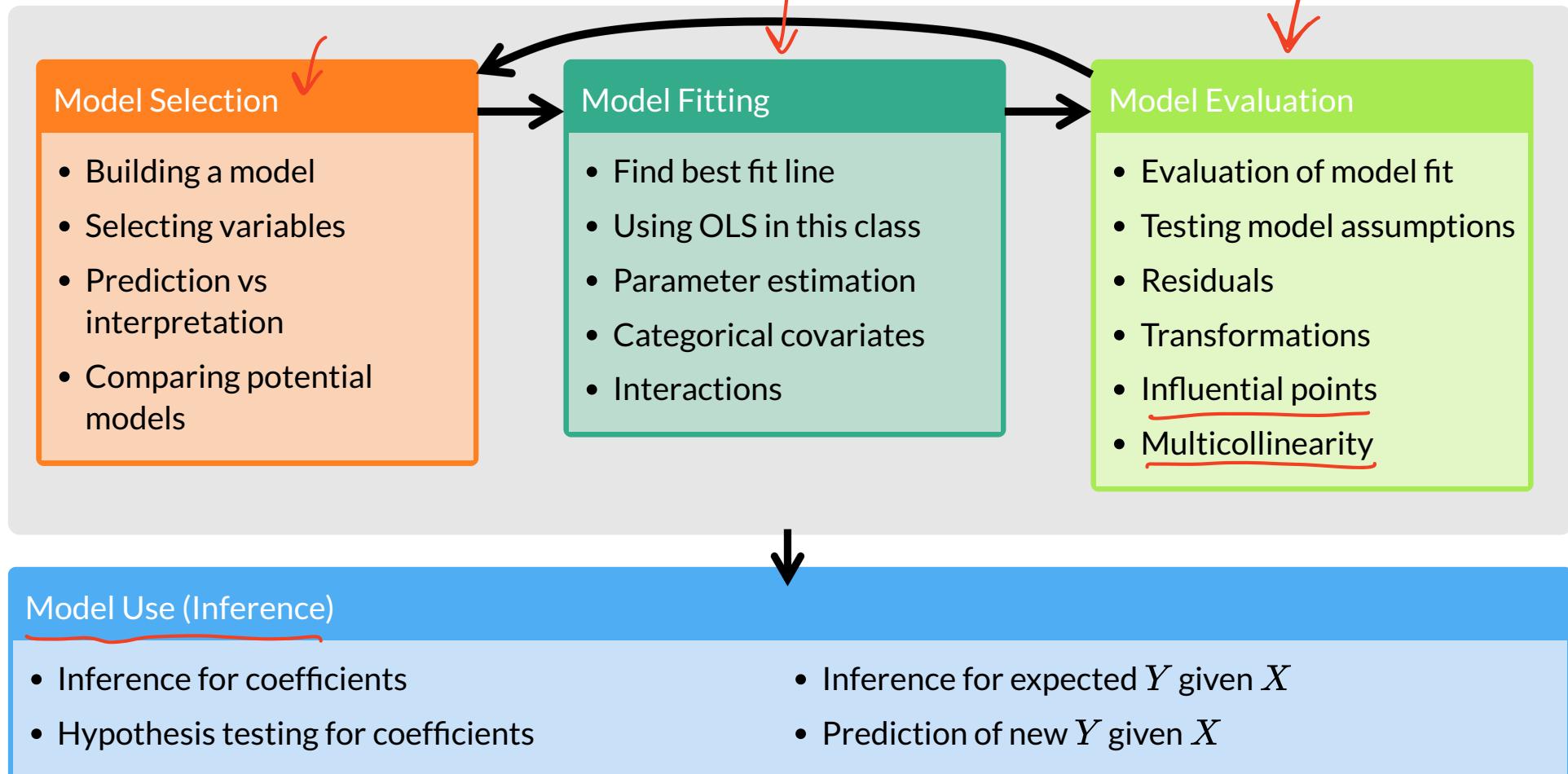
2024-03-13

Learning Objectives

- 1. Apply tools from SLR (Lesson 6: SLR Diagnostics) in MLR to evaluate LINE assumptions, including residual plots and QQ-plots
- 2. Apply tools involving standardized residuals, leverage, and Cook's distance from SLR (Lesson 7: SLR Diagnostics 2) in MLR to flag potentially influential points
- 3. Use Variance Inflation Factor (VIF) and its general form to detect and correct multicollinearity

Regression analysis process

lm()



Let's remind ourselves of the final model

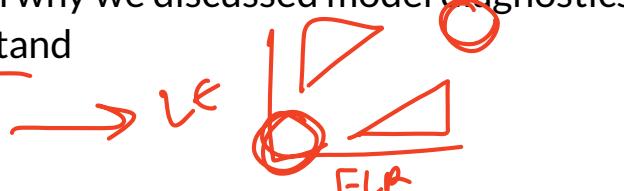
- Our **final model** contains
 - Female Literacy Rate `FLR`
 - CO2 Emissions in quartiles `C02_q`
 - Income levels in groups assigned by Gapminder `income_levels1`
 - World regions `four_regions`
 - Membership of global and economic groups `members_oecd_g77`
 - Food Supply `FoodSupplykcPPD`
 - Clean Water Supply `WaterSupplePct`

► Display regression table for final model

term	estimate	std.error	statistic	p.value
(Intercept)	39.877	4.889	8.157	0.000
FemaleLiteracyRate	-0.073	0.047	-1.555	0.125
CO2_q(0.806,2.54]	1.099	1.914	0.574	0.568
CO2_q(2.54,4.66]	-0.292	2.419	-0.121	0.904
CO2_q(4.66,35.2]	-0.595	2.524	-0.236	0.814
income_levels1Lower middle income	5.441	2.343	2.322	0.024
income_levels1Upper middle income	6.111	2.954	2.069	0.043
income_levels1High income	7.959	3.277	2.429	0.018
four_regionsAmericas	9.003	2.050	4.391	0.000
four_regionsAsia	5.260	1.637	3.213	0.002
four_regionsEurope	6.855	2.871	2.387	0.020
WaterSourcePrct	0.166	0.066	2.496	0.015
FoodSupplykcPPD	0.004	0.002	1.825	0.073
members_oecd_g77oecd	1.119	2.674	0.418	0.677
members_oecd_g77others	1.047	2.511	0.417	0.678

It's a lot to visualize

- Part of the reason why we discussed model diagnostics in SLR was so that we could have accompanying visuals to help us understand



- With 7 variables in our final model, it is hard to visualize outliers and influential points

- I highly encourage you revisit Lesson 6 and 7 (SLR Diagnostics) to help understand these notes

Remember our friend `augment()`?

- Run `final_model` through `augment()` (`final_model` is input)
 - So we assigned `final_model` as the output of the `lm()` function
- Will give us values about each observation in the context of the fitted regression model
 - cook's distance (`.cooksdist`), fitted value (`.fitted`, \hat{Y}_i), leverage (`.hat`), residual (`.resid`), standardized residuals (`.std.resid`)

```
1 aug = augment(final_model)
2 head(aug) %>% relocate(.fitted, .resid, .std.resid, .hat, .cooksdist, .after = LifeExpectancyYrs)
```

A tibble: 6 × 14

	LifeExpectancyYrs	.fitted	.resid	.std.resid	.hat	.cooksdist	FemaleLiteracyRate
1	56.7	61.5	-4.78	-1.43	0.327	0.0663	13
2	76.7	75.3	1.38	0.387	0.227	0.00293	95.7
3	60.9	58.6	2.30	0.684	0.320	0.0147	58.6
4	76.9	74.7	2.21	0.620	0.238	0.00799	99.4
5	76	76.9	-0.879	-0.233	0.145	0.000614	97.9
6	73.8	74.6	-0.796	-0.214	0.168	0.000618	99.5

i 7 more variables: CO2_q <fct>, income_levels1 <fct>, four_regions <fct>,

RDocumentation on the `augment()` function.

Learning Objectives

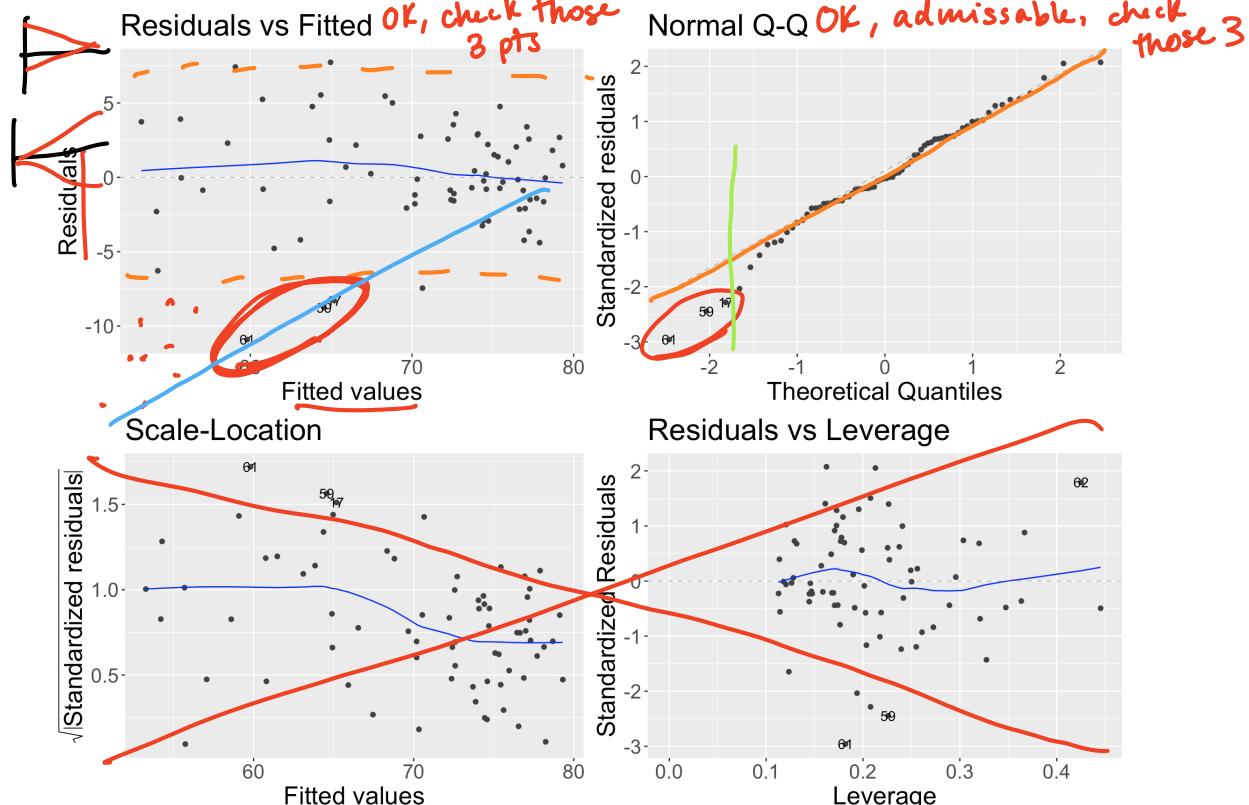
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Summary of the assumptions and their diagnostic tool

Assumption	What needs to hold?	Diagnostic tool
✓ Linearity	<ul style="list-style-type: none">Relationship between each X and Y is linear	<ul style="list-style-type: none">Scatterplot of Y vs. X
✓ Independence	<ul style="list-style-type: none">Observations are independent from each other	<ul style="list-style-type: none">Study design
Normality	<ul style="list-style-type: none"><u>Residuals</u> (and thus $Y X_1, X_2, \dots, X_p$) are <u>normally distributed</u>	<ul style="list-style-type: none">QQ plot of residualsDistribution of residuals
	<ul style="list-style-type: none"><u>Variance of residuals</u> (and thus $Y X_1, X_2, \dots, X_p$) is same across <u>fitted values</u> (homoscedasticity)	<ul style="list-style-type: none">Residual plot

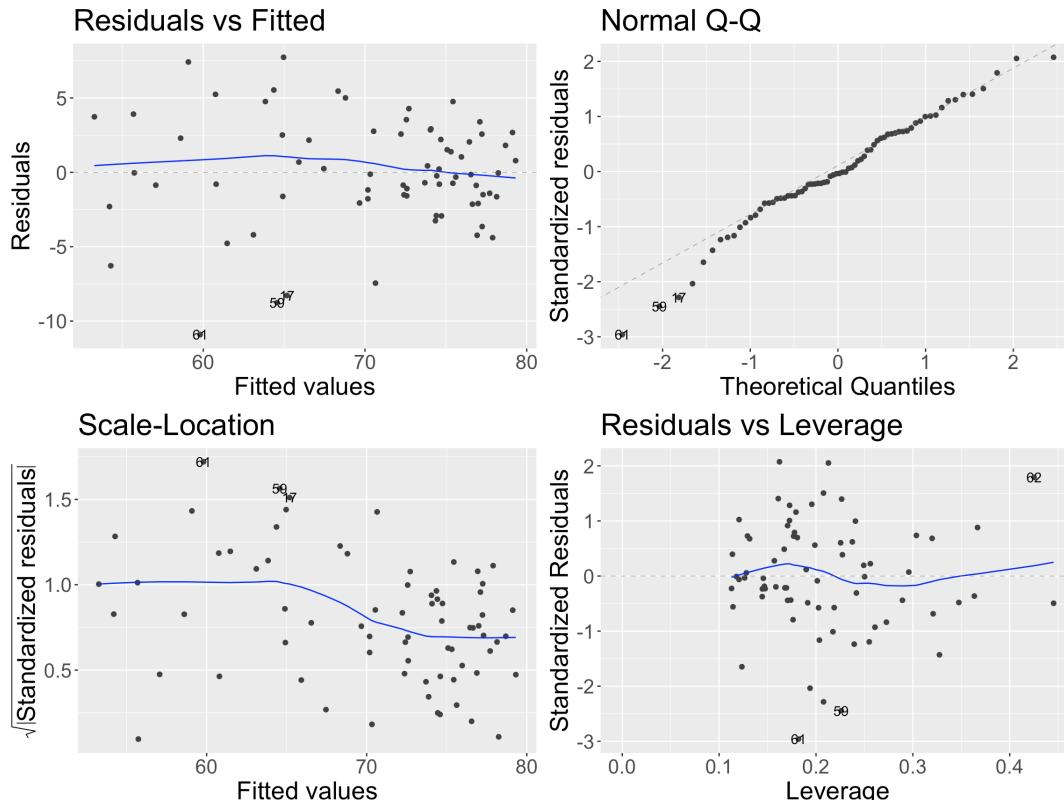
autoplot() to examine equality of variance and Normality

```
1 library(ggfortify)  
2 autoplot(final_model) + theme(text=element_text)
```



autoplot() to examine equality of variance and Normality

```
1 library(ggfortify)  
2 autoplot(final_model) + theme(text=element_text)
```



Looks like 3 obs are flagged:

- 17: Cote d'Ivoire
- 59: South Africa
- 61: Kingdom of Eswatini (formerly Swaziland in 2011)

Without them, QQ-plot and residual plot
look good

- Points on QQ-plot are close to identity line
- Residuals have pretty consistent spread across fitted values

But don't take them out!!!

- Instead, discuss what may be missing in our regression model that is not capturing the characteristics of these countries

Poll Everywhere Question 1

X

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Take a look at Lesson 7 notes with influential points in SLR. What do you think will happen with the influential points comparing SLR to our final model with MLR?

- Less influential points because more covariates might explain observations ✓  87%
- Same influential points because more covariates might not explain observations ✓  0%
- More influential points because more covariates explain less observations that were previously explained in SLR  7%
- Idk, can't we just find out? ✓  7%

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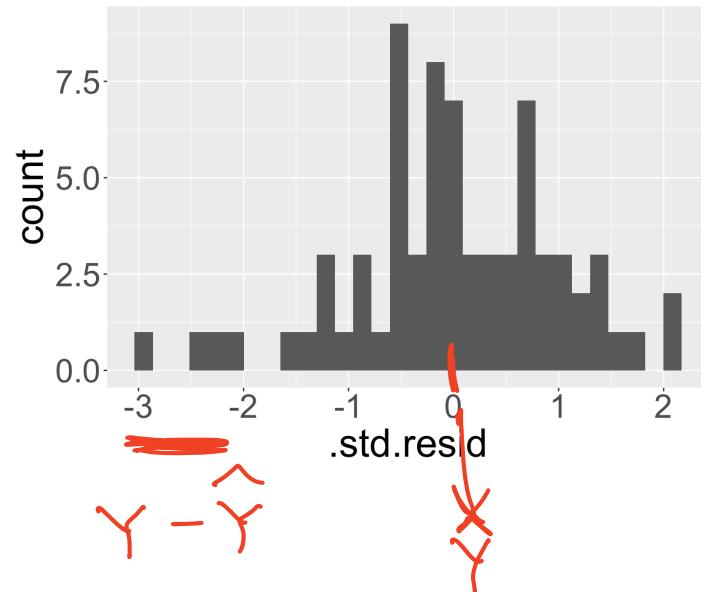
Identifying outliers

Internally standardized residual

$$r_i = \frac{\hat{\epsilon}_i}{\sqrt{\hat{\sigma}^2(1 - h_{ii})}}$$

- We flag an observation if the standardized residual is “large”
 - Different sources will define “large” differently
 - PennState site uses $|r_i| > 3$
 - `autoplot()` shows the 3 observations with the highest standardized residuals
 - Other sources use $|r_i| > 2$, which is a little more conservative

```
1 ggplot(data = aug) +  
2   geom_histogram(aes(x = .std.resid))
```



Countries that are outliers ($|r_i| > 2$)

- We can identify the countries that are outliers

```
1 aug %>% relocate(.std.resid, .after = country) %>%
2   filter(abs(.std.resid) > 2) %>% arrange(desc(abs(.std.resid)))
```

A tibble: 6 × 15

country	.std.resid	LifeExpectancyYrs	FemaleLiteracyRate	CO2_q	income_levels1
<chr>	<dbl>	<dbl>	<dbl>	<fct>	<fct>
1 Swaziland	-2.96	48.9	87.3	(0.8...	Lower middle ...
2 South Af...	-2.45	55.8	92.2	(4.6...	Upper middle ...
3 Cote d'I...	-2.28	56.9	47.6	[0.0...	Lower middle ...
4 Cape Ver...	2.07	72.7	80.3	(0.8...	Lower middle ...
5 Sudan	2.05	66.5	63.2	[0.0...	Lower middle ...
6 Vanuatu	-2.04	63.2	81.5	[0.0...	Lower middle ...

i 9 more variables: four_regions <fct>, WaterSourcePrct <dbl>,

Leverage h_i

- Values of leverage are: $0 \leq h_i \leq 1$
- We flag an observation if the leverage is “high”
 - Only good for SLR: Some textbooks use $h_i > 4/n$ where n = sample size $p=1$ coef
 - Only good for SLR: Some people suggest $h_i > 6/n$ $p=1$ coef
 - Works for MLR: $h_i > 3p/n$ where p = number of regression coefficients

```
1 aug = aug %>% relocate(.hat, .after = FemaleLiteracyRate)
2 aug %>% arrange(desc(.hat))
```

A tibble: 72 × 15

country	LifeExpectancyYrs	FemaleLiteracyRate	.hat	CO2_q	income_levels1
<chr>	<dbl>	<dbl>	<dbl>	<dbl>	<fct>
1 Mexico	75.8	92.3	0.445	(2.5...	Upper middle ...
2 Tajikistan	69.9	99.6	0.425	[0.0...	Lower middle ...
3 Bosnia and H...	76.9	96.7	0.367	(4.6...	Upper middle ...
4 Uzbekistan	69	99.2	0.363	(2.5...	Lower middle ...
5 Bangladesh	71	53.4	0.347	[0.0...	Lower middle ...
6 Afghanistan	56.7	13	0.327	[0.0...	Low income
7 Zimbabwe	51.9	80.1	0.321	[0.0...	Low income
-	-	-	-	-	-

Countries with high leverage ($h_i > 3p/n$)

- We can look at the countries that have high leverage: there are NONE

```
1 n = nrow(gapm2); p = length(final_model$coefficients) - 1
2 aug %>%
3   filter(.hat > 3*p/n) %>%
4   arrange(desc(.hat))

# A tibble: 0 × 15
# i 15 variables: country <chr>, LifeExpectancyYrs <dbl>,
#   FemaleLiteracyRate <dbl>, .hat <dbl>, CO2_q <fct>, income_levels1 <fct>,
#   four_regions <fct>, WaterSourcePrct <dbl>, FoodSupplykcPPD <dbl>,
#   members_oecd_g77 <chr>, .fitted <dbl>, .resid <dbl>, .sigma <dbl>,
#   .cooksrd <dbl>, .std.resid <dbl>
```

Identifying points with high Cook's distance

The Cook's distance for the i^{th} observation is

$$d_i = \frac{h_i}{2(1 - h_i)} \cdot r_i^2$$

where h_i is the leverage and r_i is the studentized residual

- No countries with high Cook's distance

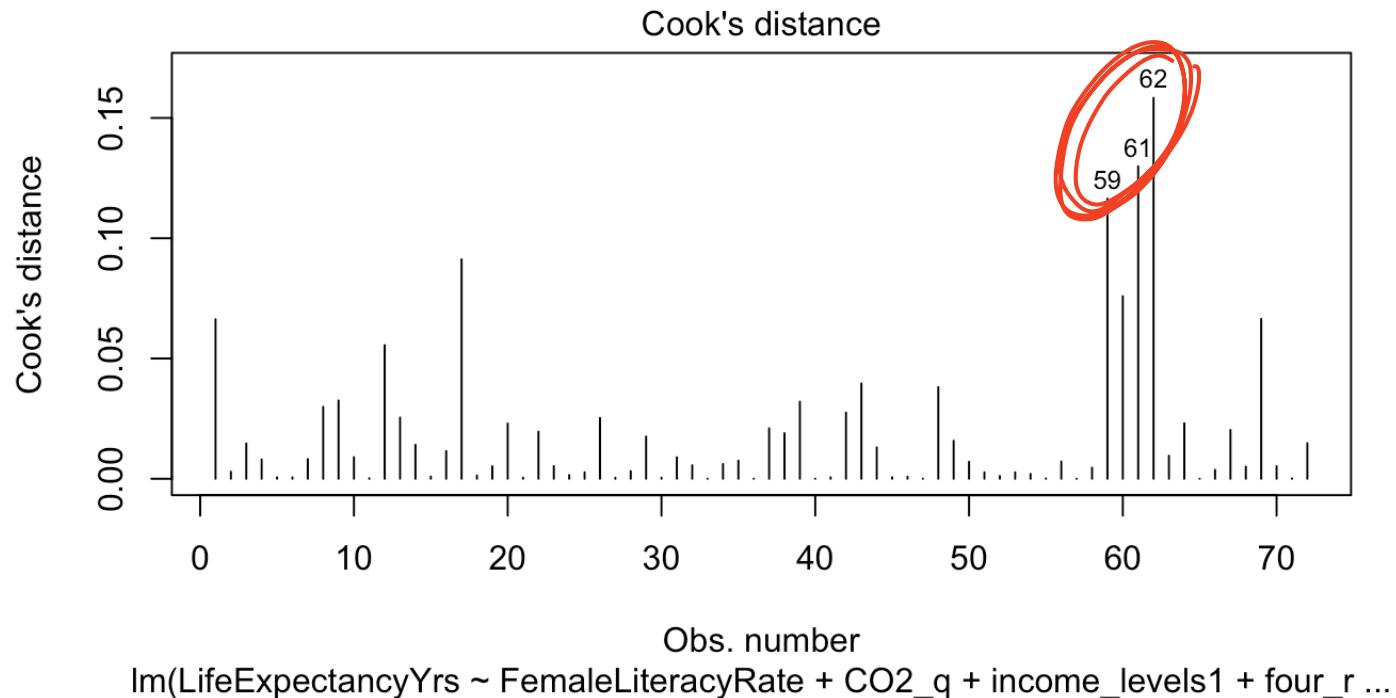
```
1 aug = aug %>% relocate(.cooksdi, .after = country)
2 aug %>% arrange(desc(.cooksdi)) %>% filter(.cooksdi > 1)

# A tibble: 0 × 15
# i 15 variables: country <chr>, .cooksdi <dbl>, LifeExpectancyYrs <dbl>,
#   FemaleLiteracyRate <dbl>, .hat <dbl>, CO2_q <fct>, income_levels1 <fct>,
#   four_regions <fct>, WaterSourcePrct <dbl>, FoodSupplykcPPD <dbl>,
#   members_oecd_g77 <chr>, .fitted <dbl>, .resid <dbl>, .sigma <dbl>,
#   .std.resid <dbl>
```

- Another rule for Cook's distance that is not strict:
 - Investigate observations that have $d_i > 1$
- Cook's distance values are already in the augment tibble: `.cooksdi`

Plotting Cook's Distance

```
1 # plot(model) shows figures similar to autoplot()  
2 # adds on Cook's distance though  
3 plot(final_model, which = 4)
```



How do we deal with influential points?

- If an observation is influential, we can check data errors:
 - Was there a data entry or collection problem?
 - If you have reason to believe that the observation does not hold within the population (or gives you cause to redefine your population)
- If an observation is influential, we can check our model:
 - Did you leave out any important predictors?
 - Should you consider adding some interaction terms?
 - Is there any nonlinearity that needs to be modeled?
- Basically, deleting an observation should be justified outside of the numbers!
 - If it's an honest data point, then it's giving us important information!
- Means we will need to discuss the limitations of our model
 - For example: Think about measurements that might help explain life expectancy that are NOT in our model
- A really well thought out explanation from StackExchange

Poll Everywhere Question 2

When we have detected problems in our model...

- We have talked about influential points
- We have talked about identifying issues with our LINE assumptions

What are our options once we have identified issues in our linear regression model?

- Are we missing a crucial measure in our dataset?
- Try a transformation if there is an issue with linearity or normality
 - Addressed in model selection
- Try a weighted least squares approach if unequal variance (oof, not enough time for us to get to)
- Try a robust estimation procedure if we have a lot of outlier issues (outside scope of class)

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What is multicollinearity? (adapted from parts of STAT 501 page)

So far, we've been ignoring something very important: multicollinearity

Multicollinearity

Two or more covariates in a multivariable regression model are *highly correlated*

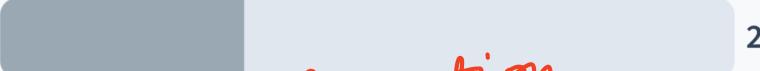
- Types of multicollinearity
 - **Structural multicollinearity**
 - Mathematical artifact caused by creating new covariates from other covariates
 - For example: If we have age, and decide to transform age to include age-squared
 - Then we have age and age-squared in the model: age-squared is perfectly predicted by age! *(centering!)*
 - **Data-based multicollinearity**
 - Result of a poorly designed experiment, reliance on purely observational data, or the inability to manipulate the system on which the data are collected.

Poll Everywhere Question 3

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In our example with life expectancy and the final model we decided on, which type of multicollinearity are we dealing with?

Structural multicollinearity  25%

how structuring transformation

Data-based multicollinearity  75%

Why is multicollinearity a problem?

In linear regression...

- Estimated regression coefficient of any one variable depends on other predictors included in the model
 - Not necessarily bad, but a big change might be an issue
- Hypothesis tests for any coefficient may yield different conclusions depending on other predictors included in the model
- Marginal contribution of any one predictor variable in reducing the error sum of squares depends on other predictors included in the model

When there is multicollinearity in our model:

- Precision of the estimated regression coefficients or correlated covariates decreases a lot
 - Basically, standard error increases and confidence intervals get wider, which means we're not as confident in our estimate anymore
 - Because highly correlated covariates are not adding much more information, but are constraining our model more

Did you notice anything about all the consequences of multicollinearity?

- All consequences relate to estimating a regression coefficient **precisely**
 - Recall that precision is linked to analysis **goals of association and interpretability**
 - See Lesson 12: Model Selection
- Multicollinearity is *not really an issue* when our **goal is prediction**
 - Highly correlated covariates/predictors will not hurt our prediction of an outcome

How do we detect multicollinearity?

- Variance inflation factors (VIF): quantifies how much the variance of the estimated coefficient for covariate k increases
 - Increases: from SLR with only covariate k to MLR with all other covariates
- General rule of thumb
 - $4 < VIF < 10$: Warrant investigation (but most people aren't investigating this...)
 - $VIF > 10$: Requires correction
 - Influencing regression coefficient estimates

VIF

$$VIF = \frac{1}{1 - R_k^2}$$

R_k^2 is the R^2 -value obtained by regressing the k^{th} covariate/predictor on the remaining predictors

$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_3 + \varepsilon$$

$VIF_1 : X_1 = \beta_0 + \beta_1 X_2 + \beta_2 X_3 + \varepsilon$

Diagram illustrating the decomposition of the regression model:

- The original model is $Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_3 + \varepsilon$.
- The first VIF calculation focuses on the coefficient of X_1 . It shows X_1 as a function of the remaining variables: $X_1 = \beta_0 + \beta_1 X_2 + \beta_2 X_3 + \varepsilon$.
- Arrows indicate dependencies:
 - An arrow labeled A^2 points from X_1 to X_2 , indicating a high correlation between X_1 and X_2 .
 - An arrow labeled A^2 points from X_1 to X_3 , indicating a high correlation between X_1 and X_3 .
 - An arrow labeled A points from X_2 to X_3 , indicating a high correlation between X_2 and X_3 .

Let's apply it to our final model

- Naive way to calculate this:

```
1 library(rms)
```

```
2 rms::vif(final_model) → every coefficient
```

FemaleLiteracyRate	CO2_q(0.806, 2.54]
4.863139	2.979224
CO2_q(2.54, 4.66]	CO2_q(4.66, 35.2]
4.758904	5.180216
income_levels1Lower middle income	income_levels1Upper middle income
5.290718	8.406927
income_levels1High income	four_regionsAmericas
7.293148	2.531966
four_regionsAsia	four_regionsEurope
2.096398	7.771994

- All $VIF < 10$
- Problem: multi-level covariates (CO2 Emissions and income level) have different VIF's even though they should be considered one variable

Let's apply it to our final model *correctly* (1/2)

- Calculate the GVIF and, more importantly, the $GVIF^{1/(2\cdot df)}$
- GVIF is the R^2 -value for regressing a covariate's group indicators on the remaining covariates
 - Captures the correlation between covariates better
- $GVIF^{1/(2\cdot df)}$ helps standardize GVIF based on how many levels each categorical covariate has
 - I'll refer to this as df-corrected GVIF or standardized GVIF
 - If continuous covariate, $GVIF^{1/(2\cdot df)} = \sqrt{GVIF}$

```
1 library(car)
2 car::vif(final_model)
```

car :: vif()
 pkg :: fn()

	GVIF	Df	$GVIF^{1/(2\cdot Df)}$
FemaleLiteracyRate	4.863139	1	2.205253
CO2_q	8.223951	3	1.420736
income_levels1	11.045885	3	1.492336
four_regions	13.935918	3	1.551277
WaterSourcePrct	4.824266	1	2.196421
FoodSupplykcPPD	3.499250	1	1.870628
members_oecd_g77	7.430919	2	1.651052

Let's apply it to our final model *correctly* (2/2)

- If continuous covariate, $\underline{GVIF^{1/(2\cdot df)}} = \underline{\sqrt{GVIF}} = \sqrt{VIF}$
- So we can square $\underline{GVIF^{1/(2\cdot df)}}$ and set VIF rules
- OR: we can correct any $\underline{GVIF^{1/(2\cdot df)} > \sqrt{10} = 3.162}$: *multicollinearity issues!*

```
1 car::vif(final_model)
```

	GVIF	Df	$GVIF^{(1/(2*Df))}$
FemaleLiteracyRate	4.863139	1	2.205253
CO2_q	8.223951	3	1.420736
income_levels1	11.045885	3	1.492336
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< 3.162

- All of these covariates are okay! No multicollinearity to correct in this dataset!

But what if we do need to make corrections for multicollinearity?

- We have been dealing with **data-based multicollinearity** in our example
- If we had issues with multicollinearity, then what are our options?
 - Remove the variable(s) with large VIF
 - Use expert knowledge in the field to decide
- If one variable has a large VIF, then there is usually another one or more variables with large VIFs
 - Basically, all the covariates that are correlated will have large VIFs
- Example: our two largest VIFs were for world region and income levels
 - Hypothetical: their $GVIF^{1/(2\cdot df)} > 3.162$
 - Remove one of them
 - I'm no expert, but from more of a data equity lens, there's a lot of generalizations made about world regions
 - I think relying on the income level of a country might give us more information as well

What about structural multicollinearity?

- Structural multicollinearity
 - Mathematical artifact caused by creating new covariates from other covariates
- For example: If we have age, and decide to transform age to include age-squared
 - Then we have age and age-squared in the model: age-squared is perfectly predicted by age!
 - By having the untransformed and transformed covariate in the model, they are inherently correlated!
- Best practice to reduce the correlation: center your covariate
 - By centering age, we no longer have a one-to-one connection between age and age-squared
 - If centered at 40yo: a 35 yo and a 45 yo will both have centered age of 5, and age-squared of 25
 $x - \bar{x}$
- Check out the Penn State site for a work through of an example with VIFs

Summary of multicollinearity

- Correlated covariates/predictors will hurt our model's precision and interpretations of coefficients
- We need to check for multicollinearity by using VIFs or GVIFs
- If $VIF > 10$ or $GVIF^{1/(2 \cdot df)} > 3.162$, we need to do something about the covariates
 - Data based: remove one the of correlated variables
 - Structural based: centering usually fixes it

Regression analysis process

