

Lesson 15: MLR Model Diagnostics

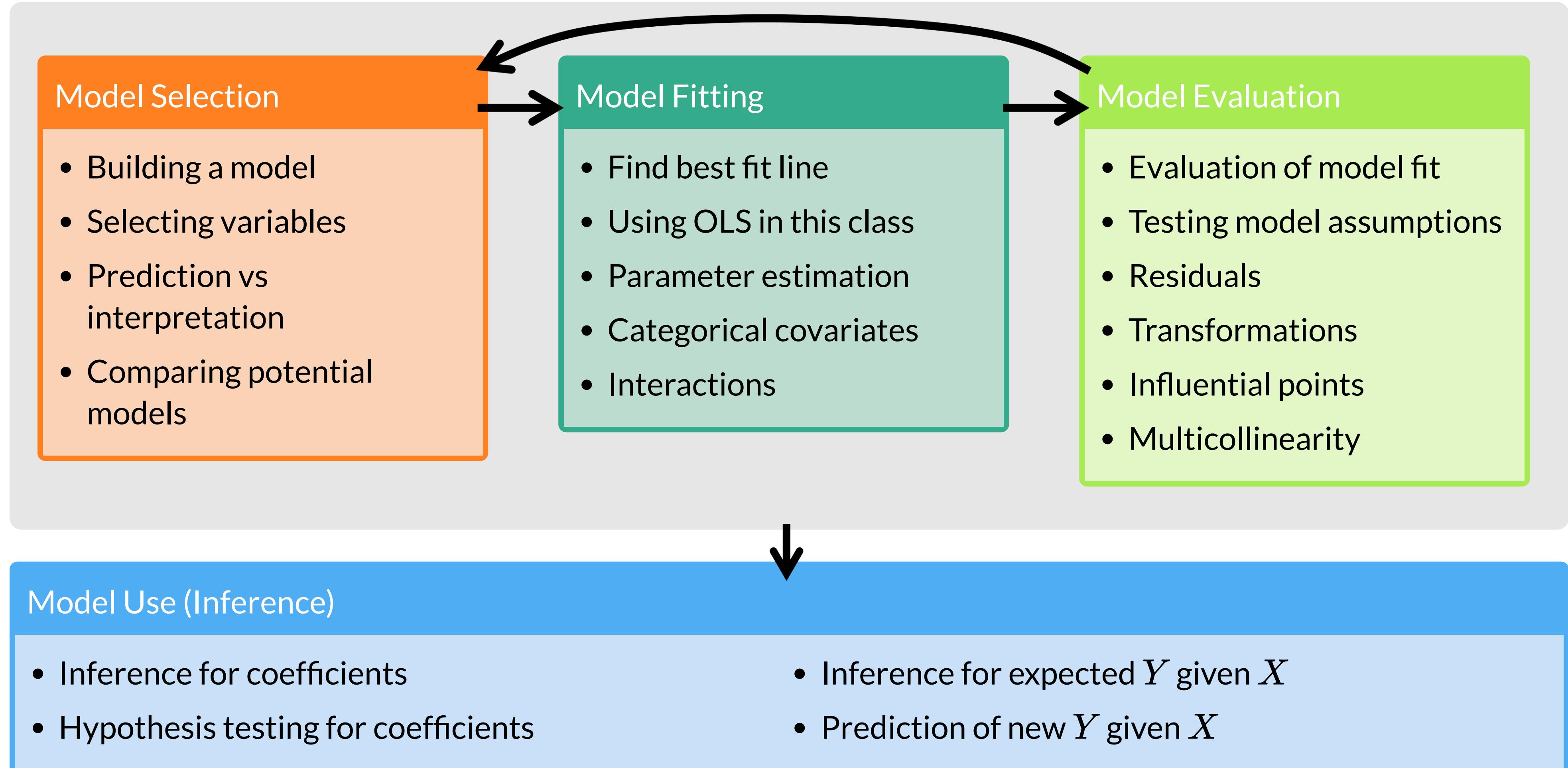
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Learning Objectives

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



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**
- No interactions

► Display regression table for final model

Characteristic	Beta	95% CI ¹	p-value
Female literacy rate (%)	-0.07	-0.17, 0.02	0.13
CO2 emissions quartiles			
[0.0439,0.806]	—	—	
(0.806,2.54]	1.1	-2.7, 4.9	0.6
(2.54,4.66]	-0.29	-5.1, 4.6	>0.9
(4.66,35.2]	-0.60	-5.6, 4.5	0.8
Income levels			
Low income	—	—	
Lower middle income	5.4	0.75, 10	0.024
Upper middle income	6.1	0.20, 12	0.043
High income	8.0	1.4, 15	0.018
World region			
Africa	—	—	
Americas	9.0	4.9, 13	<0.001
Asia	5.3	2.0, 8.5	0.002
Europe	6.9	1.1, 13	0.020
Access to improved water (%)	0.17	0.03, 0.30	0.015
Food supply (kcal PPD)	0.00	0.00, 0.01	0.073
Intergovernmental group			
g77	—	—	
oecd	1.1	-4.2, 6.5	0.7
others	1.0	-4.0, 6.1	0.7

¹ CI = Confidence Interval

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 the SLR lessons ([SLR: Checking model assumptions](#) and [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 (`.cooksdi`), \hat{Y}_i (`.fitted`), leverage (`.hat`), residuals (`.resid`), std residuals (`.std.resid`)

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

# A tibble: 6 × 14
#>   LifeExpectancyYrs .fitted .resid .std.resid .hat .cooksdi FemaleLiteracyRate
#>   <dbl>     <dbl>   <dbl>      <dbl> <dbl>       <dbl>                 <dbl>
#> 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      76.0      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: C02_q <fct>, income_levels1 <fct>, four_regions <fct>,
#> # WaterSourcePrct <dbl>, FoodSupplykcPPD <dbl>, members_oecd_g77 <chr>,
#> # .sigma <dbl>
```

RDocumentation on the `augment()` function.

Learning Objectives

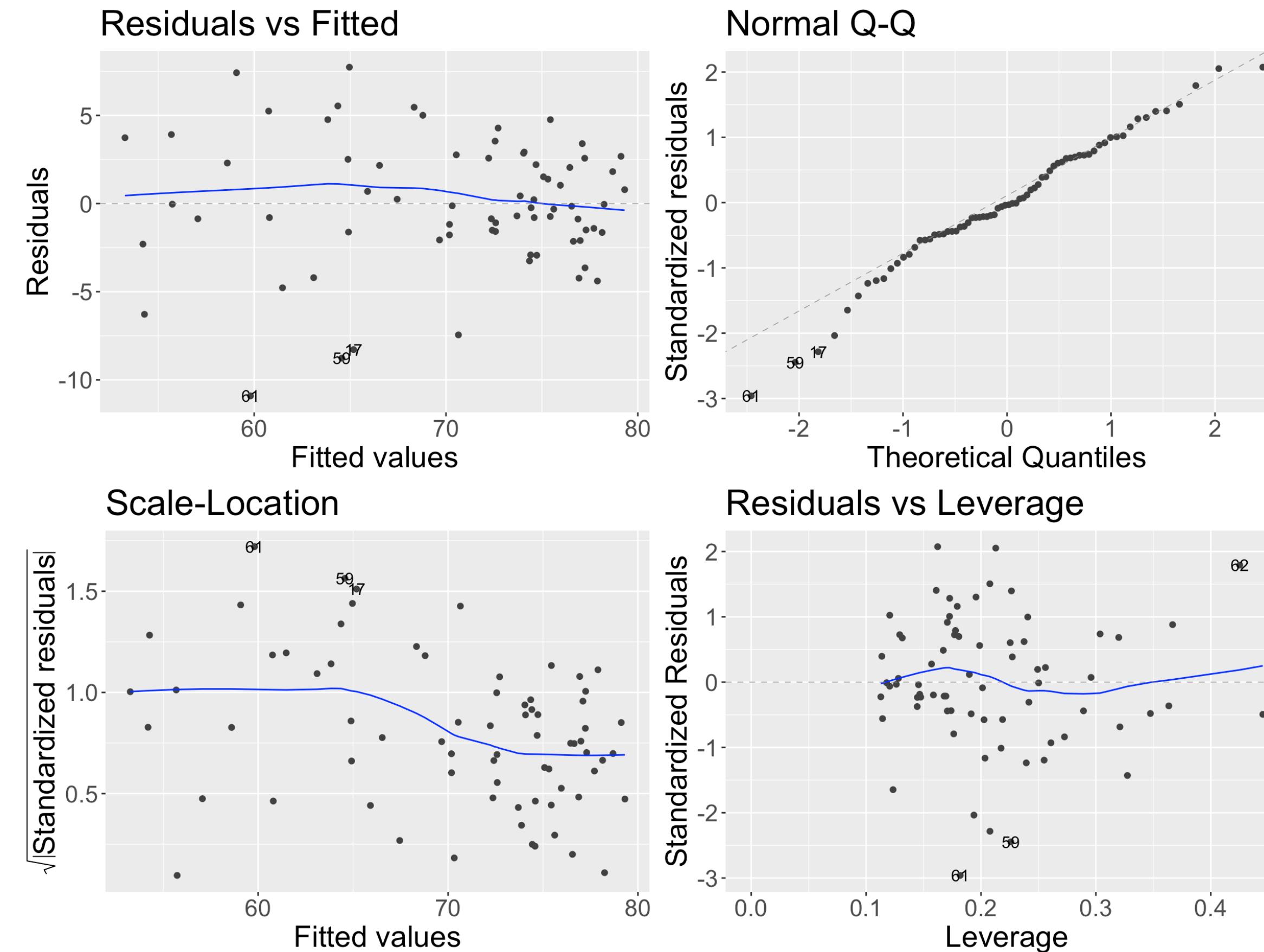
1. Apply tools from SLR (Lesson 6: SLR Diagnostics) in MLR to **evaluate LINE assumptions**, including residual plots and QQ-plots
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Summary of the assumptions and their diagnostic tool

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">Residuals (and thus $Y X_1, X_2, \dots, X_p$) are normally distributed	<ul style="list-style-type: none">QQ plot of residualsDistribution of residuals
Equality of variance	<ul style="list-style-type: none">Variance of residuals (and thus $Y X_1, X_2, \dots, X_p$) is same across fitted values (homoscedasticity)	<ul style="list-style-type: none">Residual plot

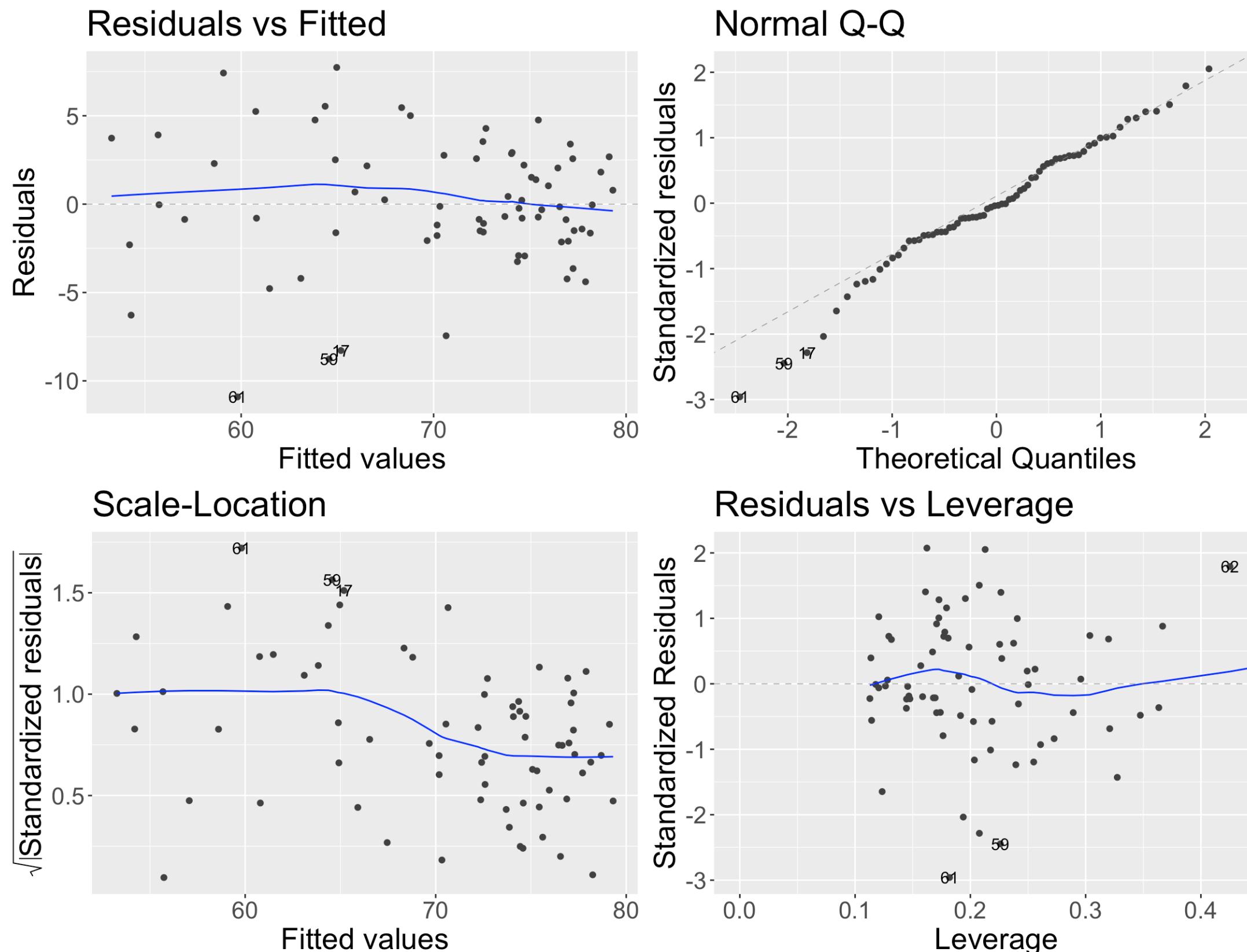
autoplot() to examine equality of variance and Normality

```
1 library(ggfortify)  
2 autoplot(final_model) + theme(text=element_text(size=20))
```



autoplot() to examine equality of variance and Normality

```
1 library(ggfortify)  
2 autoplot(final_model) + theme(text=element_text(size=12))
```



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

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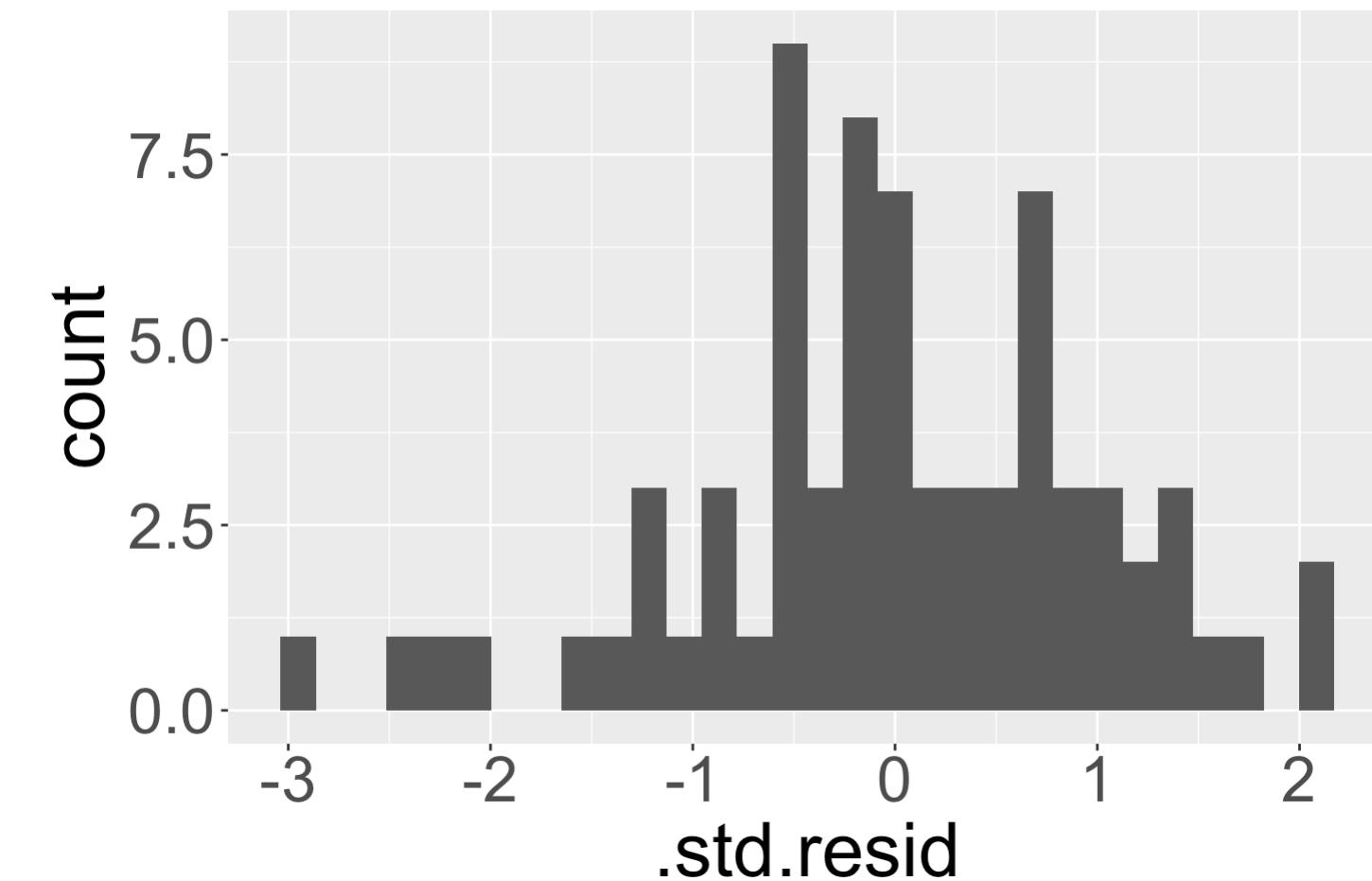
Identifying outliers

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 ($|r_i| > 3$)

- We can identify the countries that are outliers

```
1 aug %>% relocate(.std.resid, .after = country) %>%  
2   filter(abs(.std.resid) > 3) %>% arrange(desc(abs(.std.resid)))  
  
# A tibble: 0 × 15  
# i 15 variables: country <chr>, .std.resid <dbl>, LifeExpectancyYrs <dbl>,  
#   FemaleLiteracyRate <dbl>, C02_q <fct>, income_levels1 <fct>,  
#   four_regions <fct>, WaterSourcePrct <dbl>, FoodSupplykcPPD <dbl>,  
#   members_oecd_g77 <chr>, .fitted <dbl>, .resid <dbl>, .hat <dbl>,  
#   .sigma <dbl>, .cooksdi <dbl>
```

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	C02_q	income_levels1	four_regions	WaterSourcePrct	FoodSupplykcPPD	members_oecd_g77	.fitted	.resid	.hat	.sigma	.cooksdi
	<chr>	<dbl>	<dbl>	<dbl>	<dbl>	<fct>	<fct>	<dbl>	<dbl>	<chr>	<dbl>	<dbl>	<dbl>	<dbl>	<dbl>
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>, FoodSupplykcPPD <dbl>, members_oecd_g77 <chr>, .fitted <dbl>, .resid <dbl>, .hat <dbl>, .sigma <dbl>, .cooksdi <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
 - **Only good for SLR:** Some people suggest $h_i > 6/n$
 - **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)) %>% head(5)
```

```
# A tibble: 5 × 15
  country      LifeExpectancyYrs FemaleLiteracyRate .hat C02_q income_levels1
  <chr>           <dbl>             <dbl>        <dbl> <dbl> <fct> <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 He...   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 ...
# i 9 more variables: four_regions <fct>, WaterSourcePrct <dbl>,
# FoodSupplykcPPD <dbl>, members_oecd_g77 <chr>, .fitted <dbl>, .resid <dbl>,
# .sigma <dbl>, .cooksdi <dbl>, .std.resid <dbl>
```

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>, C02_q <fct>, income_levels1 <fct>,
#   four_regions <fct>, WaterSourcePrct <dbl>, FoodSupplykcPPD <dbl>,
#   members_oecd_g77 <chr>, .fitted <dbl>, .resid <dbl>, .sigma <dbl>,
#   .cooksdi <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

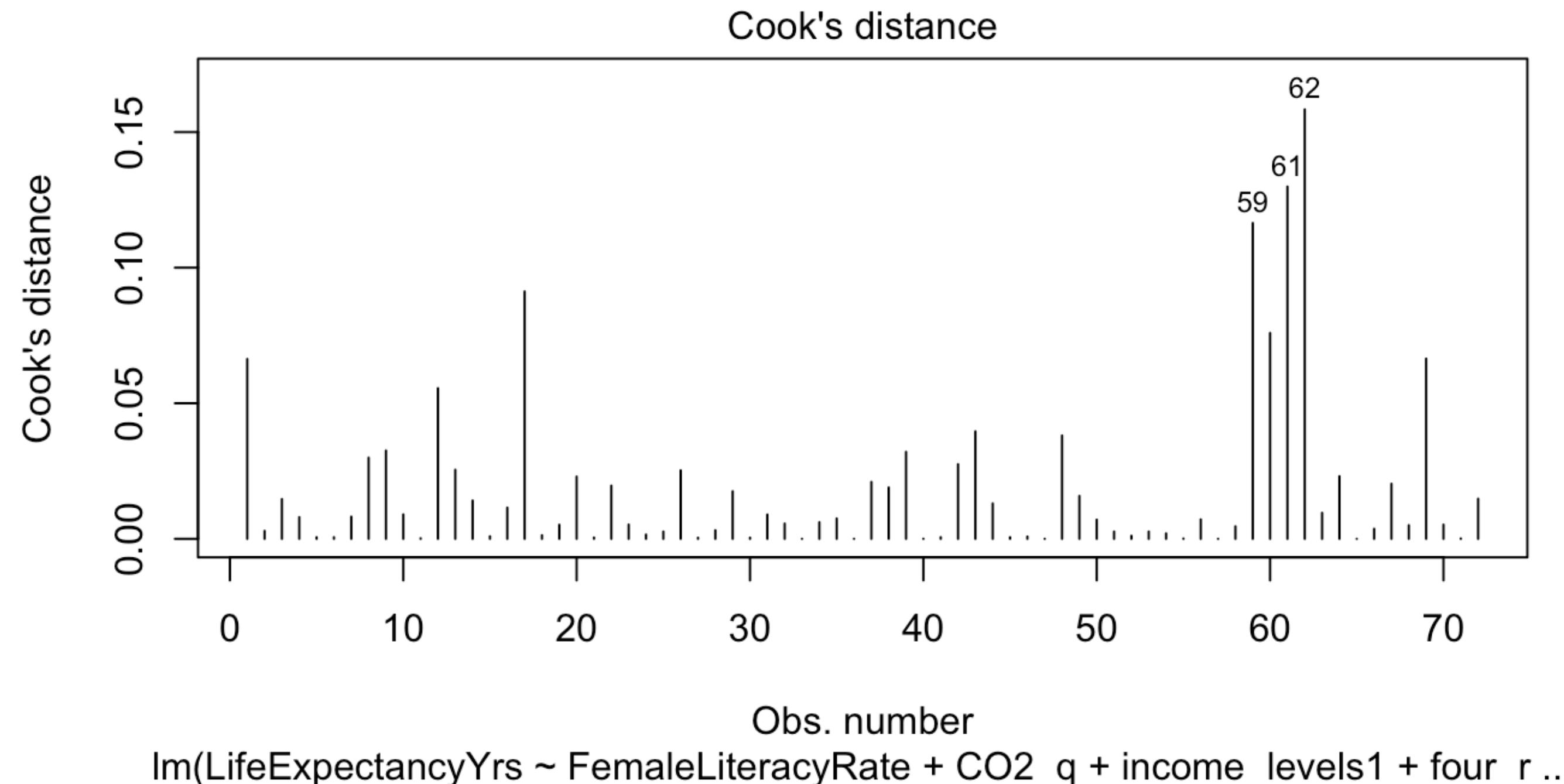
```
1 aug = aug %>% relocate(.cooksdi, .after = country)
2 aug %>% arrange(desc(.cooksdi)) %>% filter(.cooksdi > 1)
```

```
# A tibble: 0 × 15
# i 15 variables: country <chr>, .cooksdi <dbl>, LifeExpectancyYrs <dbl>,
#   FemaleLiteracyRate <dbl>, .hat <dbl>, C02_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() but adds Cook's distance  
2 plot(final_model, which = 4)
```



- Identify 3 highest Cook's distance: 59, 61, 62 (South Africa, Kingdom of Eswatini, Tajikistan)

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 categorization or transformation (or numeric variables) 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!
 - **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
 - Or we are using two variables that are practically measuring the same thing

Poll Everywhere Question 3

Why is multicollinearity a problem?

In linear regression, depending on the other predictors in the model, the following will change:

- Estimated regression coefficient of any one variable
 - Not necessarily bad, but a big change might be an issue
- Hypothesis tests for any coefficient may yield different conclusions
- Contribution of any one predictor variable in reducing sum of squared errors

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 on **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
 - $VIF < 4$: Good!
 - $4 < VIF < 10$: 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

Let's apply it to our final model

- Naive way to calculate this in R:

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

FemaleLiteracyRate	C02_q(0.806,2.54]
4.863139	2.979224
C02_q(2.54,4.66]	C02_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
WaterSourcePrct	FoodSupplykcPPD
4.824266	3.499250
members_oecd_g77oecd	members_oecd_g77others
2.720955	5.125196

- 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)
```

	GVIF	Df	GVIF ^{1/(2*Df)}
FemaleLiteracyRate	4.863139	1	2.205253
C02_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, $GVIF^{1/(2\cdot df)} = \sqrt{GVIF}$
- So we can square $GVIF^{1/(2\cdot df)}$ and set VIF rules
- OR: we can correct any $GVIF^{1/(2\cdot df)} > \sqrt{10} = 3.162$

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

	GVIF	Df	GVIF^(1/(2*Df))
FemaleLiteracyRate	4.863139	1	2.205253
C02_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

- 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 GVIFs 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
- 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 of the correlated variables
 - Structural based: centering usually fixes it

Regression analysis process

