# Assignment 2: Summarize global cesarean delivery rates and GDP across 137 countries

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- Solutions will be released on Tuesday, September 8.
- This semester, homework assignments are for practice only and will not be turned in for marks.

#### Helpful hints:

- Every function you need to use was taught during lecture! So you may need to revisit the lecture code to help you along by opening the relevant files on Datahub. Alternatively, you may wish to view the code in the condensed PDFs posted on the course website. Good luck!
- Knit your file early and often to minimize knitting errors! If you copy and paste code for the slides, you are bound to get an error that is hard to diagnose. Typing out the code is the way to smooth knitting! We recommend knitting your file each time after you write a few sentences/add a new code chunk, so you can detect the source of knitting errors more easily. This will save you and the GSIs from frustration! You must knit correctly before submitting.
- If your code runs off the page of the knitted PDF then you will LOSE POINTS! To avoid this, have a look at your knitted PDF and ensure all the code fits in the file (you can easily view it on Gradescope via the provided link after submitting). If it doesn't look right, go back to your .Rmd file and add spaces (new lines) using the return or enter key so that the code runs onto the next line.

## Summarizing global cesarean delivery rates and GDP across 137 countries

#### Introduction

Recall from this week's lab that we constructed bar charts and histograms to explore a data set that looked at global rates of cesarean delivery and GDP. If you need a refresher, you can view your knitted file from lab and remind yourself what you found.

In this week's assignment, you will describe these distributions using numbers. You will investigate the **mean** and **median** of the distribution of GDP. You will also examine the distribution of cesarean delivery separately for countries of varying income levels. Lastly, you will describe the **spread** of the distributions using **quartiles** and make a **box plot**.

Execute this code chunk to load the required libraries:

CS\_data <- CS\_data %>% mutate(CS\_rate\_100 = CS\_rate\*100)

```
library(readr)
library(dplyr)
library(ggplot2)
```

Just like last time, read in the data that is stored as a .csv file and assign it to an object called CS\_data. We will also use dplyr's mutate() to create the new cesarean delivery variable that ranges between 0 and 100:

```
CS_data <- read_csv("cesarean.csv")</pre>
## Parsed with column specification:
## cols(
##
     Country_Name = col_character(),
##
     CountryCode = col character(),
     Births_Per_1000 = col_double(),
##
##
     Income_Group = col_character(),
##
     Region = col character(),
     GDP_2006 = col_double(),
##
     CS_rate = col_double()
##
# This code reorders the Factor variable `Income Group` in the
# order specified in this function. This will affect the order the ggplot
# panels are shown in question 8 when we use `facet_wrap()`.
CS_data$Income_Group <- forcats::fct_relevel(CS_data$Income_Group,</pre>
                                             "Low income", "Lower middle income",
                                             "Upper middle income", "High income: nonOECD",
                                             "High income: OECD")
```

1. [1.5 points] Fill in the blanks indicated by "—-" by saving the answer to each blank in the code chunk below. Make sure you capitalize correctly, as R is case-sensitive.

The function  $\mathtt{mutate}()$  takes the old variable called  $-(\mathbf{a})-$  and multiplies it by — to make a new variable called  $-(\mathbf{b})-$ .

Investigate what would have happened had we not assigned the data using <- to CS\_data? Re-run the code without the assignment and see what happens.

```
# First, let's re-read in the data as we did in the previous chunk
CS_data <- read_csv("cesarean.csv")</pre>
## Parsed with column specification:
## cols(
##
     Country_Name = col_character(),
##
     CountryCode = col_character(),
##
     Births_Per_1000 = col_double(),
##
     Income Group = col character(),
     Region = col_character(),
##
##
     GDP 2006 = col double(),
##
     CS_rate = col_double()
## )
CS_data$Income_Group <- forcats::fct_relevel(CS_data$Income_Group,
                                             "Low income", "Lower middle income",
                                             "Upper middle income", "High income: nonOECD",
                                             "High income: OECD")
# Now, we try again without the re-assignment to CS_data
CS_data %>% mutate(CS_rate_100 = CS_rate*100)
## # A tibble: 137 x 8
      Country_Name CountryCode Births_Per_1000 Income_Group Region GDP_2006
##
##
      <chr>>
                   <chr>
                                          <dbl> <fct>
                                                              <chr>
                                                                        <dbl>
##
    1 Albania
                   ALB
                                             46 Upper middl~ Europ~
                                                                        3052.
##
    2 Andorra
                   AND
                                              1 High income~ Europ~
                                                                       42417.
   3 United Arab~ ARE
                                             63 High income~ Middl~
                                                                       42950.
  4 Argentina
                                            689 High income~ Latin~
                                                                        6649.
##
                   ARG
   5 Armenia
                   ARM
                                             47 Lower middl~ Europ~
                                                                        2127.
##
##
  6 Australia
                   AUS
                                            267 High income~ East ~
                                                                       36101.
  7 Austria
                   AUT
                                             76 High income~ Europ~
                                                                       40431.
##
   8 Azerbaijan
                   AZE
                                            166 Upper middl~ Europ~
##
                                                                        2473.
                                            119 High income~ Europ~
## 9 Belgium
                   BEL
                                                                       38936.
## 10 Benin
                   BEN
                                            342 Low income
                                                              Sub-S~
                                                                         557.
## # ... with 127 more rows, and 2 more variables: CS_rate <dbl>,
       CS_rate_100 <dbl>
# check the variables on CS_data
names(CS data)
## [1] "Country_Name"
                          "CountryCode"
                                            "Births_Per_1000" "Income_Group"
```

Did CS\_rate\_100 get added to the data set? No. You can tell by using head(CS\_data) to view the first few rows and notice that the variable hasn't been added. This is because when we don't assign the output to anything, it just prints it out for us to see. Nothing is saved. So, we want to save the output by assigning the result of the code to a variable, which in this case, we used CS\_data. In general, you want to use new variable names at every significant step in your analysis as you work with your data, so that you have access to the data at all those significant stages. However, if you are performing multiple small operations on the same dataset, you can overwrite the original variable, since you know you won't be needing the in-between steps anyway.

"CS rate"

"GDP 2006"

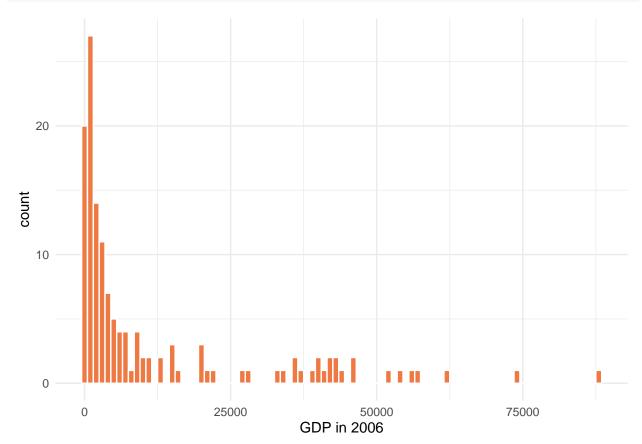
## [5] "Region"

# This overwrites the original CS\_data object
CS\_data <- CS\_data %>% mutate(CS\_rate\_100 = CS\_rate\*100)

#### GDP: Summarizing measures of centrality

Recall your histogram of GDP from this week's lab:

```
ggplot(data = CS_data, aes(x = GDP_2006)) +
  geom_histogram(col = "white", fill = "sienna2", binwidth = 1000) +
  xlab("GDP in 2006") +
  theme_minimal()
```



2. [1 point] Describe the shape of this distribution. Is it "skewed left", "skewed right", "symmetric", or "bimodal"? Uncomment one of the possible choices.

```
# p2 <- "skewed left"
p2 <- "skewed right"
# p2 <- "symmetric"
# p2 <- "bimodal"

check_problem2()</pre>
```

```
## [1] "Checkpoint 1 Passed: You chose the correct shape of distribution!"
##
## Problem 2
## Checkpoints Passed: 1
## Checkpoints Errored: 0
## 100% passed
## ------
## Test: PASSED
```

3. [1 point] Based on your answer, will the mean be approximately the "same", "larger than", or "smaller than" the median?

```
# p3 <- "same"
p3 <- "larger than"
# p3 <- "smaller than"

# This only checks that you've selected an answer, not its correctness.
check_problem3()

## [1] "Checkpoint 1 Passed: Correct choice!"

##
## Problem 3
## Checkpoints Passed: 1
## Checkpoints Errored: 0
## 100% passed
## -------
## Test: PASSED</pre>
```

#### 4. [3 points] Describe, in words, how the mean and median are calculated:

The mean is calculated by adding up all the numbers and dividing that sum by how many numbers there are. The median is calculated by ordering a list of numbers from smallest to largest, and then finding the number at the very middle or the list.

To calculate the mean and median in R, we can use the summarize() function from the dplyr package. The summarize() function is used anytime we want to take a variable and summarize something about it into one number, like it's mean or median. Here is the code to summarize GDP\_2006's mean and print it out to the screen. In the code, we name the mean mean\_GDP and output the result to the screen:

```
GDP_summary <- CS_data %>% summarize(mean_GDP = mean(GDP_2006))
GDP_summary
```

```
## # A tibble: 1 x 1
## mean_GDP
## <dbl>
## 1 11791.
```

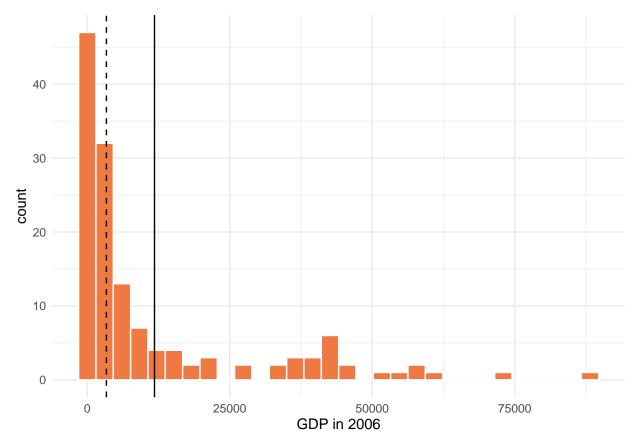
5. [1 point] Extend the above code to also summarize the median. Call the median summary median\_GDP. Assign this summary to GDP\_summary (it will overwrite the previous version):

```
GDP_summary <- CS_data %>% summarize(mean_GDP = mean(GDP_2006), median_GDP = median(GDP_2006))
GDP_summary
## # A tibble: 1 x 2
##
    mean_GDP median_GDP
##
                   <dbl>
        <dbl>
## 1
       11791.
                   3351.
check_problem5()
## [1] "Checkpoint 1 Passed: Correct mean value has been calculated!"
## [1] "Checkpoint 2 Passed: Correct median value has been calculated!"
## Problem 5
## Checkpoints Passed: 2
## Checkpoints Errored: 0
## 100% passed
## -----
## Test: PASSED
```

6. [2 points] geom\_vline() can be used to add the mean and the median to the histogram shown above. This geom\_vline() adds a vertical line to the graph. You need to specify where to add the line by passing it an "x-intercept" argument. Remove the comments (i.e., the three "#") from the code below and update the geom\_vline() code to plot lines at the mean and median by telling it the mean and median estimates. The argument lty=1 (standing for line type) will plot a solid line and lty=2 will plot a dashed line.

For the purposes of this question, please assign xintercept to a plain NUMERIC, not a variable or expression

## `stat\_bin()` using `bins = 30`. Pick better value with `binwidth`.



```
# REMINDER: this is only a sanity check; it doesn't check for accuracy
check_problem6()
```

```
## [1] "Checkpoint 1 Passed: A ggplot has been defined"
## [1] "Checkpoint 2 Passed: A vertical line of the correct mean added!"
## [1] "Checkpoint 3 Passed: A vertical line of the correct median added!"
##
## Problem 6
```

```
## Checkpoints Passed: 3
## Checkpoints Errored: 0
```

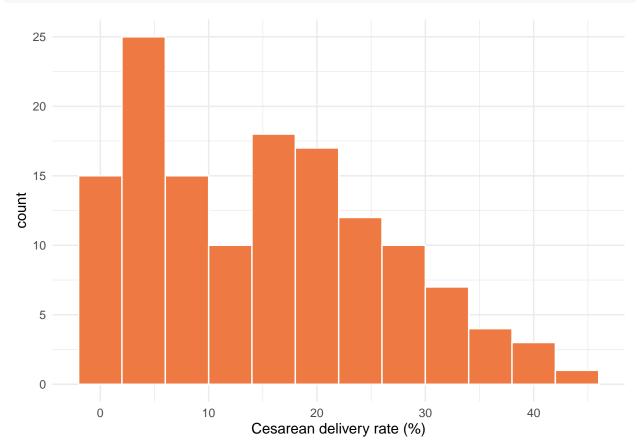
## 100% passed ## -----

## Test: PASSED

#### Summarizing the distribution of cesarean delivery

Recall the distribution of cesarean delivery rates across countries:

```
ggplot(data = CS_data, aes(x = CS_rate_100)) +
geom_histogram(binwidth = 4, col = "white", fill = "sienna2") +
xlab("Cesarean delivery rate (%)") +
theme_minimal()
```



7. [1 point] Describe the shape of this distribution. Is it "skewed left", "skewed right", "symmetric", or "bimodal"?

```
# p7 <- "skewed left"
# p7 <- "symmetric"
p7 <- "bimodal"

# This only checks that you've selected an answer, not its correctness.
check_problem7()

## [1] "Checkpoint 1 Passed: You chose the correct shape of distribution!"

## ## Problem 7

## Checkpoints Passed: 1

## Checkpoints Errored: 0

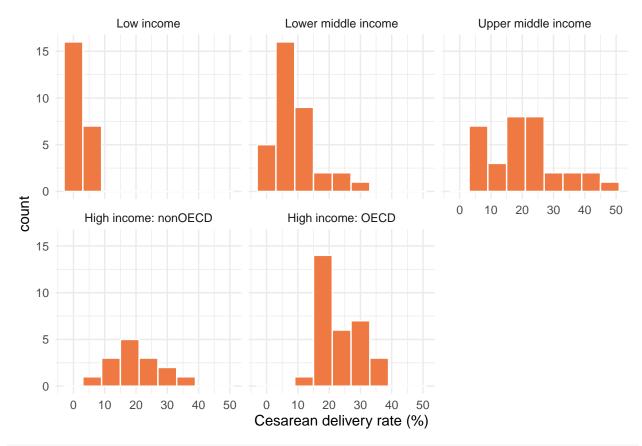
## 100% passed

## -------

## Test: PASSED</pre>
```

There appears to be multiple peaks which sometimes points to there being another variable that might explain the peaks. We can make a separate histogram for each income group using the facet\_wrap() function.

8. [1 point] Extend the ggplot code given below using the facet\_wrap() statement to make a separate histogram for each level of the Income\_Group variable:



#### check\_problem8()

## Test: PASSED

```
## [1] "Checkpoint 1 Passed: A ggplot has been defined"
## [1] "Checkpoint 2 Passed: A separate histogram for each level of the `Income_Group` variable has been
## Problem 8
## Checkpoints Passed: 2
## Checkpoints Errored: 0
## 100% passed
```

9.	2	[points]	Based	on	$\mathbf{this}$	plot	and	the	previous	one,	$\operatorname{describe}$	why	the the	data	had	$\mathbf{two}$	peaks
----	---	----------	-------	----	-----------------	------	-----	-----	----------	------	---------------------------	-----	---------	------	-----	----------------	-------

The ceserean delivery rate peaks of lower income versus higher income countries are different, and therefore create two peaks in the combined data histogram.

#### 10. [1 point] Why might lower income countries have lower rates of cesarean delivery?

Lower income countries might not have the proper medical equipment to carry out ceserean deliveries. These countries could be struggling financially so ceserean deliveries aren't as available as in higher income countries.

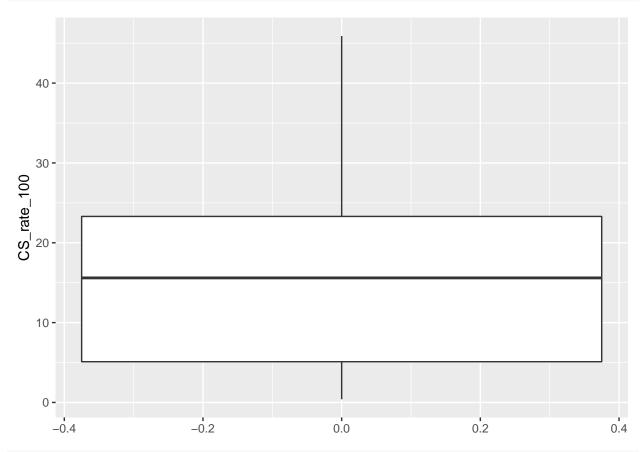
11. [2 points] Calculate the mean\_CS and median\_CS of CS\_rate\_100 using only one summarize() command. Assign this summary to the name CS\_summary and then print the results by typing CS\_summary so you can see the contents.

```
CS_summary <- CS_data %>% summarize(mean_CS = mean(CS_rate_100), median_CS = median(CS_rate_100))
CS_summary
## # A tibble: 1 x 2
##
    mean_CS median_CS
##
       <dbl>
                 <dbl>
## 1
        15.3
                  15.6
check problem11()
## [1] "Checkpoint 1 Passed: Correct mean value has been calculated!"
## [1] "Checkpoint 2 Passed: Correct median value has been calculated!"
##
## Problem 11
## Checkpoints Passed: 2
## Checkpoints Errored: 0
## 100% passed
## -----
## Test: PASSED
```

#### Measures of variation

12. [2 marks] Use ggplot2 to make a boxplot of the distribution of CS\_rate\_100

```
p12 <- ggplot(CS_data, aes(y = CS_rate_100)) + geom_boxplot()
p12</pre>
```



#### check\_problem12()

```
## [1] "Checkpoint 1 Passed: A ggplot has been defined"
## [1] "Checkpoint 2 Passed: The correct y axis has been defined!"
## [1] "Checkpoint 3 Passed: A box plot has been defined!"
##
## Problem 12
## Checkpoints Passed: 3
## Checkpoints Errored: 0
## 100% passed
## -------
## Test: PASSED
```

Recall that the box plot summarizes the distribution in five numbers: the minimum, the first quartile (with 25% of the data below it), the median, the third quartile (with 75% of the data below it), and the maximum. Each of these numbers has at least one corresponding R function:

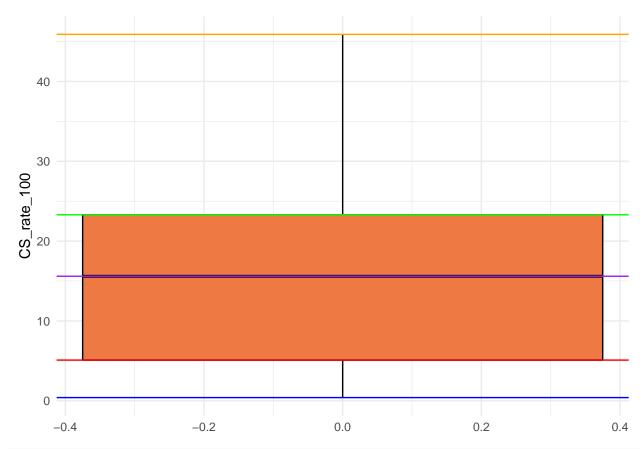
Number	R function
Minimum First quartile Median Third quartile Maximum	<pre>min(variable) quantile(variable, probs = 0.25) median(variable) or quantile(variable, probs = 0.5) quantile(variable, probs = 0.75) max(variable)</pre>

13. [2 points] Use a combination of dplyr's summarize function and the above functions to compute the five number summary of CS\_rate\_100. Assign the summary to the name five\_num\_summary, which should contain values for min, Q1, median, Q3, and max.

```
five_num_summary <- CS_data %>% summarize(min = min(CS_rate_100),
Q1 = quantile(CS_rate_100, 0.25), median = median(CS_rate_100), Q3 = quantile(CS_rate_100, 0.75),
max = max(CS rate 100))
five_num_summary
## # A tibble: 1 x 5
##
      min
              Q1 median
                           Q3
     <dbl> <dbl> <dbl> <dbl> <dbl> <
##
      0.4
           5.1
                   15.6 23.3 45.9
check_problem13()
## [1] "Checkpoint 1 Passed: Correct min value has been computed!"
## [1] "Checkpoint 2 Passed: Correct value of the first quartile has been computed!"
## [1] "Checkpoint 3 Passed: Correct median value has been computed!"
## [1] "Checkpoint 4 Passed: Correct value of the third quartile has been computed!"
## [1] "Checkpoint 5 Passed: Correct max value has been computed"
## Problem 13
## Checkpoints Passed: 5
## Checkpoints Errored: 0
## 100% passed
## -----
## Test: PASSED
```

Double check that <code>geom\_boxplot()</code> is making the box plot correctly. You can do this by adding horizontal lines to the plot at each number in your five number summary using <code>geom\_hline()</code>. Because horizontal lines intercept the y-axis, <code>geom\_hline()</code> requires the <code>yintercept</code> argument that you can set to each number in your summary.

### 14. [2 points] The code below includes one horizontal line at the minimum shown in blue. Add the rest of the lines:



#### check\_problem14()

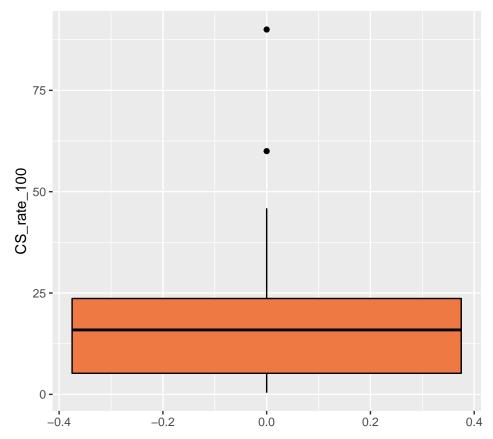
```
## [1] "Checkpoint 1 Passed: A geom_hline has been defined!"
## [1] "Checkpoint 2 Passed: A geom_hline has been defined!"
## [1] "Checkpoint 3 Passed: A geom_hline has been defined!"
## [1] "Checkpoint 4 Passed: A geom_hline has been defined!"
## [1] "Checkpoint 5 Passed: A geom_hline has been defined!"
##
## Problem 14
## Checkpoints Passed: 5
```

## Checkpoints Errored: 0

## 100% passed ## -----

## Test: PASSED

15. [4 marks] Compile the following code which adds two points to the CS\_data, makes a new dataset called CS\_data\_plus\_2, and redraws the box plot. How did the box plot change? Perform a calculation to justify why it changed. What are the newly-added features on the plot called?



## 1

0.4 5.20

15.9 23.6

```
new_IQR <- CS_data_plus_2 %>% summarize(min = min(CS_rate_100),
Q1 = quantile(CS_rate_100, 0.25), median = median(CS_rate_100), Q3 = quantile(CS_rate_100, 0.75),
max = max(CS_rate_100))
new_IQR

## # A tibble: 1 x 5
## min Q1 median Q3 max
## <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> </dbl>
```

The Q1, median, Q3, and max of the dataset all increased. The newly-added features to the box plot are called outliers. These outliers shifted all the quartile values and max higher. The box plot got flatter/more "squished," and the two new points were also plotted at the top of the box plot.

#### Check your score

Click on the middle icon on the top right of this code chunk (with the downwards gray arrows and green bar) to run all your code in order. Then, run this chunk to check your score.

```
# Just run this chunk.
total_score()
```

Туре	Points_Possible	Test					##
autograded	2	PASSED			1	${\tt Problem}$	##
autograded	1	PASSED			2	${\tt Problem}$	##
autograded	1	PASSED			3	${\tt Problem}$	##
free-response	3	${\tt GRADED}$	YET	NOT	4	${\tt Problem}$	##
autograded	1	PASSED			5	${\tt Problem}$	##
autograded	2	PASSED			6	${\tt Problem}$	##
autograded	1	PASSED			7	${\tt Problem}$	##
autograded	1	PASSED			8	${\tt Problem}$	##
free-response	2	GRADED	YET	NOT	9	${\tt Problem}$	##
free-response	1	GRADED	YET	NOT	10	${\tt Problem}$	##
autograded	1	PASSED			11	${\tt Problem}$	##
autograded	2	PASSED			12	${\tt Problem}$	##
autograded	2	PASSED			13	${\tt Problem}$	##
autograded	2	PASSED			14	${\tt Problem}$	##
free-response	4	GRADED	YET	NOT	15	Problem	##