# Assignment 4: Mind the Gap

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## Exercise 1

## #View(gapminder)

- i. Which variables in the dataset are categorical?
- : country,continent, region, year
  - ii. Which variables in the dataset are continuous (i.e. numerical)?
- : infant\_mortality, life\_expectancy, fertility, population, gdp
  - iii. What does each row in the dataset represent?
- : each country's information

```
gapminder %>%
  group_by(continent) %>%
  summarize(
    count = n()
)
```

continent	count
Africa	2907
Americas	2052
Asia	2679
Europe	2223
Oceania	684

```
gapminder %>%
  summarize(
    mean = mean(infant_mortality, na.rm = TRUE),
    median = median(infant_mortality, na.rm = TRUE),
    standard_deviation = sd(infant_mortality, na.rm = TRUE),
    interquartile_range = IQR(infant_mortality, na.rm = TRUE)
)
```

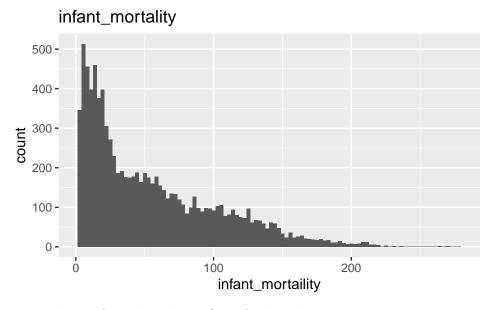
mean	median	$standard\_deviation$	interquartile_range
55.30862	41.5	47.72805	69.1

## Exercise 3

i.

```
gapminder %>%
  ggplot() +
  geom_histogram(
    mapping = aes(x = infant_mortality),
    bins = 100
) +
  labs(
    title = "infant_mortality",
    x = "infant_mortality"
)
```

## Warning: Removed 1453 rows containing non-finite values (stat\_bin).



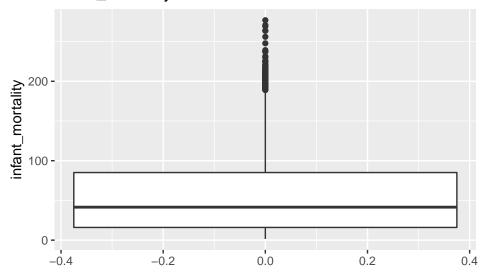
- 1) What is the shape of the distribution? : left skewed
- 2) Why do you think that most of the data points occur where they do (i.e. what is the real-world interpretation of this graph)? : infant\_mortaility is barely appearing in the real-world
- ii. box plot

```
gapminder %>%
  ggplot() +
  geom_boxplot(
    mapping = aes(y = infant_mortality)
) +
  labs(
```

```
title = "infant_mortality",
   y = "infant_mortality"
)
```

## Warning: Removed 1453 rows containing non-finite values (stat\_boxplot).

## infant\_mortality

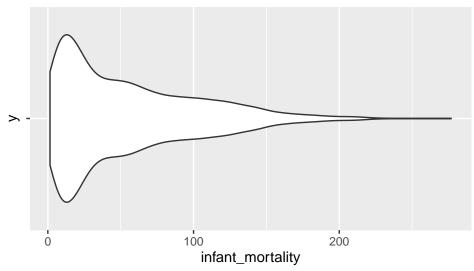


- 1) What is the shape and where is center of this distribution?
- $\rightarrow$  symmetric  $\rightarrow$  0.0 is the center
  - iii. violin plot

```
gapminder %>%
  ggplot() +
  geom_violin(
    mapping = aes(x = infant_mortality, y="")
) +
  labs(
    title = "infant_mortality",
    x = "infant_mortality"
)
```

## Warning: Removed 1453 rows containing non-finite values (stat\_ydensity).

# infant\_mortality



1) What is the shape and where is the center of this distribution? left skewed, center is 150

- i. histogram
- 1) **Describe each graph separately (shape and center).** Right skewed histogram/center is at around 60

```
gapminder %>%
  ggplot() +
  geom_histogram(
    mapping = aes(x = life_expectancy),
    bins = 100
) +
  labs(
    title = "life_expectancyy",
    x = "life_expectancy")
```

# life\_expectancyy 400300100200 40 60 80 life\_expectancy

- ii. boxplot
- 1) Describe each graph separately (shape and center). Symmetrical, center = 0.0

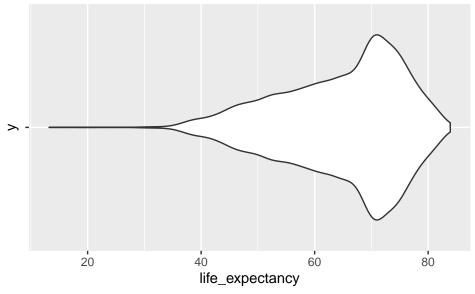
```
gapminder %>%
  ggplot() +
  geom_boxplot(
    mapping = aes(y = life_expectancy)
) +
  labs(
    title = "life_expectancy",
    y = "life_expectancy"
)
```

# 

- iii. violin plot
- 1) Describe each graph separately (shape and center) right skewed, center is at 60.

2) Do all three graphs show the same pattern(s), or do any of the graphs display patterns in the The histogram and violin plot show the right skewed plot while box plot showed the symmetrial shape. Since the box plot is graphed under then condition that y is the parameter unlike other two plots are drawn upon  $x = life_e$ expectancy.

```
gapminder %>%
  ggplot() +
  geom_violin(
    mapping = aes(x = life_expectancy, y="")
)
```



```
labs(
   title = "life_expectancy",
   x = "life_expectancy"
)

## $x

## [1] "life_expectancy"

##

## $title

## [1] "life_expectancy"

##

## attr(,"class")

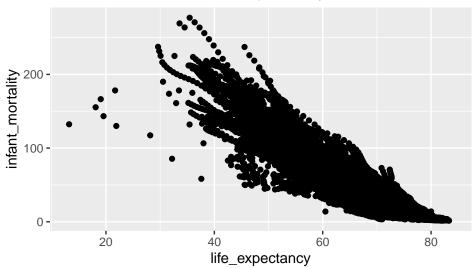
## [1] "labels"
```

```
gapminder %>%
  ggplot() +
  geom_point(
    mapping = aes(
        x = life_expectancy,
```

```
y = infant_mortality
)
) +
labs(
  title = "covariation between life_expectancy and infant_mortality",
  x = "life_expectancy",
  y = "infant_mortality"
)
```

## Warning: Removed 1453 rows containing missing values (geom\_point).

## covariation between life\_expectancy and infant\_mortality



Describe any patterns you see in this graph.

infant\_mortality and life\_expectancy show the negative covariation.

```
gapminder %>%
  ggplot() +
  geom_bar(mapping = aes(x = continent))
```

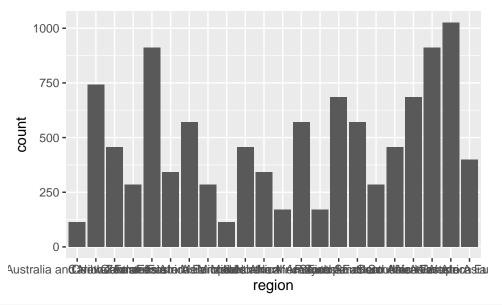
```
2000 - Africa Americas Asia Europe Oceania continent
```

```
theme(axis.text.x =element_text(angle = 45, hjust = 1)) +

labs(
   title = "continent",
   x = "continent"
)
```

```
## List of 3
   $ axis.text.x:List of 11
     ..$ family
##
                      : NULL
     ..$ face
##
                      : NULL
     ..$ colour
##
                      : NULL
##
     ..$ size
                      : NULL
     ..$ hjust
##
                      : num 1
     ..$ vjust
                      : NULL
##
     ..$ angle
##
                      : num 45
     ..$ lineheight
                      : NULL
##
     ..$ margin
                      : NULL
##
                      : NULL
##
     ..$ debug
     ..$ inherit.blank: logi FALSE
##
##
     ..- attr(*, "class")= chr [1:2] "element_text" "element"
                 : chr "continent"
##
## $ title
                 : chr "continent"
## - attr(*, "class")= chr [1:2] "theme" "gg"
   - attr(*, "complete")= logi FALSE
   - attr(*, "validate")= logi TRUE
gapminder %>%
  ggplot() +
  geom_bar(mapping = aes(x = region), bins = 150)
```

## Warning: Ignoring unknown parameters: bins



```
theme(axis.text.x =element_text(angle = 45, hjust = 1)) +

labs(
   title = "region",
   x = "region"
)
```

```
## List of 3
    $ axis.text.x:List of 11
     ..$ family
##
                       : NULL
##
     ..$ face
                       : NULL
##
     ..$ colour
                       : NULL
     ..$ size
##
                       : NULL
##
     ..$ hjust
                       : num 1
##
     ..$ vjust
                       : NULL
     ..$ angle
##
                       : num 45
     ..$ lineheight
                       : NULL
##
                       : NULL
##
     ..$ margin
     ..$ debug
                       : NULL
##
     ..$ inherit.blank: logi FALSE
##
     ..- attr(*, "class")= chr [1:2] "element_text" "element"
##
                 : chr "region"
##
    $ title
                 : chr "region"
##
   - attr(*, "class")= chr [1:2] "theme" "gg"
   - attr(*, "complete")= logi FALSE
   - attr(*, "validate")= logi TRUE
```

- explore the covariation between two or more variables
- compare multiple countries at a single time or multiple times within a single country
- add time into analyses as a second variable (year, categorical variable -> create a new column

```
for this) as.factor(...)
```

- use the mutate() to create a new variable called year\_cat, which contains the year variable as a categorical var(using as.factor(year))
- store the output dataframe in a new var, gapminder\_cat

```
gapminder_cat <- gapminder%>%
  mutate(as.factor(year))
```

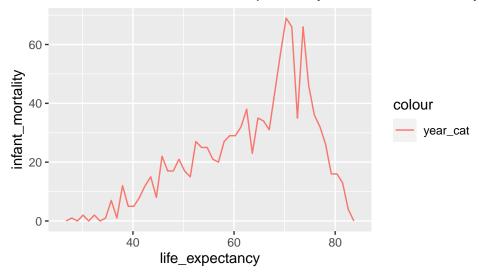
## Exercise 8

```
gapminder_cat %>%
  filter(year%% 10==0) %>%
ggplot() +
    geom_freqpoly(
        mapping = aes(
            x = life_expectancy,
            color = "year_cat",
            ),
        bins = 50,

) +

labs(title = "covariation between life_expectancy and infant_mortality",
        x = "life_expectancy",
        y = "infant_mortality")
```

## covariation between life\_expectancy and infant\_mortality



```
gapminder_cat %>%
    ggplot() +
    geom_histogram(
```

```
mapping = aes(
    x = life_expectancy,
    color = "year_cat",
    ),
    bins=30,

) +
    facet_wrap(~ continent)+

labs(
    title = "covariation between life_expectancy and infant_mortality",
    x = "life_expectancy",
    y = "infant_mortality")
```

## covariation between life\_expectancy and infant\_mortality

