

Descriptive Statistic, ggplot2, and a Little Bit of LM

Summarizing Commands

- `max(x, na.rm = FALSE)` – It shows the maximum value. By default, NA values are not removed. NA is considered the largest unless `na.rm=true` is used.
- `min(x, na.rm = FALSE)` – Shows minimum value in a vector. If there are na values, NA is returned unless `na.rm=true` is used.
- `length(x)` – Gives length of the vector and includes na values. `Na.rm=instruction` does not work with this command.
- `sum(x, na.rm = FALSE)` – Shows the sum of the vector elements.
- `mean(x, na.rm = FALSE)` – We obtain an arithmetic mean with this.
- `median(x, na.rm = FALSE)` – Shows the median value of the vector.
- `sd(x, na.rm = FALSE)` – Shows the standard deviation.
- `var(x, na.rm = FALSE)` – Shows the variance.

Summarizing Commands

- `mad(x, na.rm = FALSE)` – Shows the median absolute deviation. This first finds the median of the data set, then computes the absolute differences between each data point and the median, and finally takes the median of these absolute differences.
- `log(dataset)` – Shows log value for each element.
- `summary(dataset)` – We have seen how it shows a summary of dataset like maximum value, minimum value, mean, etc.
- `quantile()` – Shows the quantiles by default—the 0%, 25%, 50%, 75%, and 100% quantiles. You can select other quantiles also.

Descriptive Summary Table

① Option 1: stargazer

- `stargazer::stargazer(DATAFRAME)`: will generate a nice table
- ONLY DATAFRAME: `as.data.frame()`
- output options: `type = "text" / "html" / "latex"` "latex" is the default
- save as file: `out = {path} -> extension!`

② Option 2: modelsummary

- `modelsummary::datasummary()`

Histograms and Scatter Plots

```
hist(worldTFR$GDPpc)
plot(x = worldTFR$GDPpc, y = worldTFR$Year)
```

In-class exercises:

We will continue using the worldTFR dataset.

- 1 Load the dataset and omit all NA's in the dataframe.
- 2 What are the column names and row names?
- 3 What is the max and min of Years?

Introduction to ggplot2

- ggplot2 is a powerful visualization package
- Shares design philosophy with dplyr -> piping and chaining with +
- `geom_graphtype()`
 - ▶ `geom_point()`
 - ▶ `geom_smooth()`
 - ▶ `geom_histogram()`
 - ▶ `geom_line()`
- `aes()`: aesthetics, x and y value, colors
- ggplot cheat sheet: <https://posit.co/wp-content/uploads/2022/10/data-visualization-1.pdf>
- We need to first install and load the package.

```
# install.packages("ggplot2")  
library(ggplot2)
```

How to Make a Simple Scatterplot

- `geom_point()`

```
data = USArrests # built-in dataset
```

```
ggplot(data, aes(x = UrbanPop, y = Assault)) +  
  geom_point()
```

```
# alternatively
```

```
ggplot() +  
  geom_point(data, aes(x = UrbanPop, y = Assault))
```

```
# sort of equivalent to doing this with the base R plot  
plot(data$UrbanPop, data$Assault)
```


With dplyr

```
library(dplyr)

USArrests %>%
  group_by(row.names(USArrests)) %>%
  filter(Murder < 9) %>%
  ggplot() + # piping -> +
  geom_point(aes(x = UrbanPop, y = Rape)) +
  theme_minimal() # minimal theme
```

Histograms with ggplot

We can use `geom_histogram`.

```
ggplot(data = worldTFR) +  
  geom_histogram(aes(GDPpc))
```

Adjusting the X and Y Axis Limits

- `xlim()` and `ylim()`
- Accepts a vector of length 2 (minimum and maximum)

```
ggplot(data, aes(x=UrbanPop, y=Assault)) +  
  geom_point() +  
  xlim(c(40, 80)) +  
  ylim(c(0, 350))    # deletes some points
```

Change the Title and Axis Labels

- `labs(title, subtitle, x, y, legend)`

```
ggplot(data, aes(x=UrbanPop, y=Assault)) +  
  geom_point() +  
  geom_smooth(method="lm") +  
  xlim(c(30, 90)) + ylim(c(0, 350)) +  
  labs(title="Urban Population VS Assault",  
        subtitle="Using the built-in dataset USArrests",  
        y="Assault", x="Urban Population") # adding labels here
```

Change the Title and Axis Labels

- `xlab()`
- `ylab()`

```
ggplot(data, aes(x=UrbanPop, y=Assault)) +  
  geom_point() +  
  geom_smooth(method="lm") +  
  xlim(c(30, 90)) + ylim(c(0, 350)) +  
  ggtitle("Urban Population Vs Assault",  
          subtitle="Using the built-in dataset USArrests") +  
  xlab("Urban Population") +  
  ylab("Assault")
```

Change the Color and Size of Points

- `aes()`

```
ggplot(data, aes(x=UrbanPop, y=Assault)) +  
  geom_point(col="pink", size=2) + # Set static color and size for points  
  xlim(c(30, 90)) + ylim(c(0, 350)) +  
  labs(title="Urban Population VS Assault",  
        subtitle="Using the built-in dataset USArrests",  
        y="Assault", x="Urban Population") +
```

Saving your plot

- `ggsave()` is a convenient function for saving a plot.
- By default, save the last plot using the current size

The following code chunk is not executable.

```
ggsave(  
  filename,  
  plot = last_plot(), # you can provide plot as an object  
  device = NULL, # automatically determined by path  
  path = NULL, # include EXTENSION!  
  scale = 1,  
  width = NA,  
  height = NA,  
  units = c("in", "cm", "mm", "px"),  
  dpi = 300,  
  limitsize = TRUE,  
  bg = NULL,  
  ...  
)
```

Saving your plot

```
ggplot(data, aes(x=UrbanPop, y=Assault)) +  
  geom_point() +  
  geom_smooth(method="lm", col="skyblue") +  
  xlim(c(30, 90)) + ylim(c(0, 350)) +  
  labs(title="Urban Population VS Assault",  
        subtitle="Using the built-in dataset USArrests",  
        y="Assault", x="Urban Population") +  
  geom_point(col="pink", size=3)  
  
ggsave("myplot.png")
```


Saving your plot

```
g1 <- ggplot(data, aes(x=UrbanPop, y=Assault)) +  
  geom_point() +  
  geom_smooth(method="lm", col="skyblue") +  
  xlim(c(30, 90)) + ylim(c(0, 350)) +  
  labs(title="Urban Population VS Assault",  
        subtitle="Using the built-in dataset USArrests",  
        y="Assault", x="Urban Population") +  
  geom_point(col="pink", size=3)  
  
ggsave("myplot.pdf", plot = g1)
```

In-class exercises:

- 1 Omit all NA's from worldTFR.
- 2 Make a plot where the horizontal axis is LifeExpB and vertical axis is TFR using `ggplot`
- 3 Set the color to red, and size for points as 0.5
- 4 Add appropriate label for the axis and the plot.
- 5 Save your plot

Simple Linear Regression

- Foundations for linear regressions will be covered more in QPM 1.
- `lm(formula, data)`
- formula has a form $y \sim x_1 + x_2$
- interaction term: $x_1 * x_2 \rightarrow x_1 + x_2 + x_1:x_2$

Simple Linear Regression

- As always, we can store the output as an object.
- `summary()`

```
# Bivariate
```

```
model1 <- lm(formula = LifeExpB ~ Pop1564Female, data = worldTFR)  
summary(model1)
```

```
# Multivariate
```

```
model2 <- lm(formula = LifeExpB ~ Pop1564Female + InfMRateUN, data = worldTFR)  
summary(model2)
```

```
# Interaction
```

```
model3 <- lm(formula = LifeExpB ~ Pop1564Female*InfMRateUN, data = worldTFR)  
summary(model3)
```

Plot the Fitted Line

- `geom_smooth()` with `method = "lm"`

```
g1 <- ggplot(data, aes(x = UrbanPop, y = Assault)) +  
  geom_point() +  
  geom_smooth(  
    method = "lm", # default is LOESS  
    formula = "y ~ x",  
    data = df,  
    se = TRUE,  
  )
```

Robust Standard Errors

- Robust standard errors are not the default.
- sandwich and lmtest

```
library(sandwich)
library(lmtest)

# Robust standard errors with lm()
modell1_robust <- coeftest(modell1, vcov = vcovHC)

# summary may not work here
print(modell1_robust)

# Clustered robust standard errors with lm()
modell1_robust_clustered <-
  coeftest(modell1,
            vcov = vcovCL,
            type = "HC1",
            cluster = ~Country)

# summary may not work here
print(modell1_robust_clustered)
```

Visual Diagnosis of Linear Model

- `plot()` function provides a quick way to visually check Gauss-Markov assumptions and outliers / high leverage points.
- This will be covered in more detail in QPM 1.

```
plot(model1, which = 1) # Residuals and Fitted values
plot(model1, which = 2) # Q-Q plot
plot(model1, which = 3) # Standardized Residuals and Fitted values
plot(model1, which = 4) # Cook's distance
plot(model1, which = 5) # Residuals and Leverages
plot(model1, which = 6) # Cook's distance and Leverages
```

Summarize Regression Outputs with `modelsummary`

- `modelsummary::modelsummary()` provides some easy ways to convert the regression model outputs into other formats such as latex and docx.

```
library(modelsummary)
```

```
# By default as markdown table
```

```
modelsummary(model3)
```

```
# Latex
```

```
modelsummary(model3, output="latex")
```

```
# or you can provide output file name such as model3.tex
```

```
# HTML
```

```
modelsummary(model3, output="html")
```

```
# or you can provide output file name such as model3.html
```

```
# Multiple models as list
```

```
modelsummary(list(model1, model2, model3))
```


Visualizing Regression Outputs with modelsummary

- modelsummary package also provides easy and beautiful ways to plot regression outputs.
- modelplot()
- a nice coefficient plot in a ggplot object

```
modelplot(model1) +  
  labs(title = "Coefficient Plot for Model1") +  
  aes(color = ifelse(p.value < 0.001,  
                     "Significant", "Not significant")) +  
  scale_color_manual(values = c("grey", "black"))
```

More on modelsummary

- modelsummary package has more useful features
 - ▶ datasummary() to generate a beautiful descriptive statistics table
- Check more details here!
<https://modelsummary.com/vignettes/modelsummary.html>

In-class exercises:

We will continue using the worldTFR dataset.

- 1 Run a simple linear regression where: GDPpc as an outcome and MtoFbirth, Pop1564, and LifeExpB as predictors.
- 2 Get robust SE for the model.
- 3 Summarize the model in tables and figures using `modelsummary`.