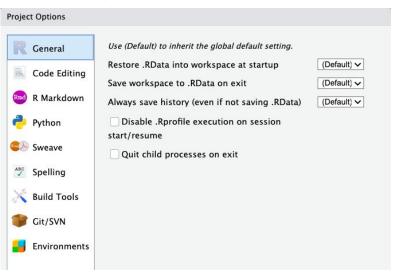
Data Visualization

Jen Cruz, Zichao Li, Isobel McEwen, Claire McLaughlin, Sanjana Srinivasan, Julien Chen

Set Up

- set up a new directory --> new project --> work in script --> save script regularly and especially before closing session
- use console for commands such as install
- use sections, section labels, and comments with command + shift + r and the #



Data Cleaning

- 1. Familiarize yourself with the data set
- 2. Check for structural errors
- 3. Check for data irregularities
- 4. Decide how to deal with missing values
- 5. Document data versions and changes made

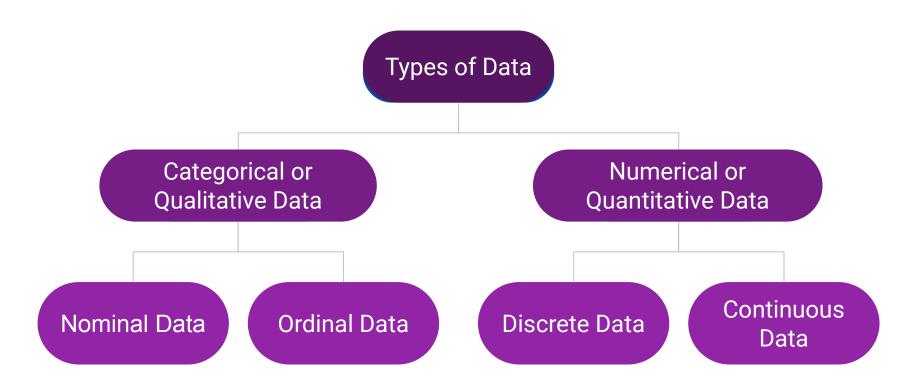
Dplyr

"Dplyr changed my life!" -Jarvis

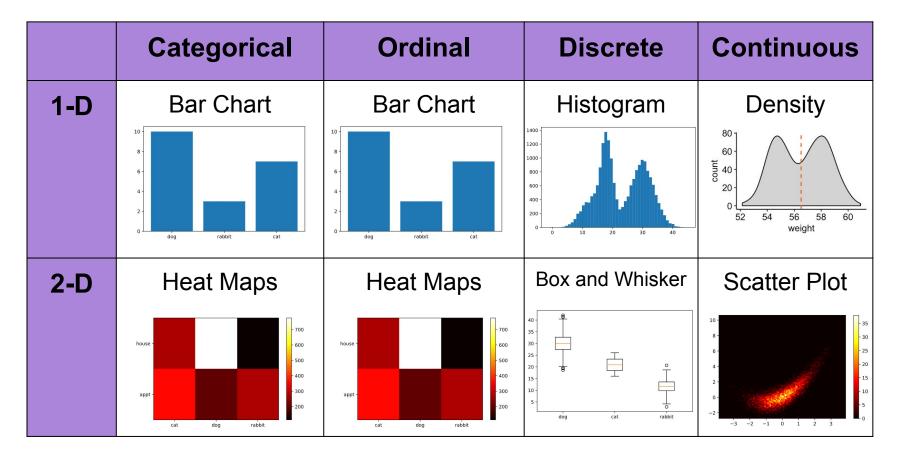
```
popsize_and_poverty <- popsize_and_poverty |>
   mutate(
    proportion_in_poverty = in_poverty / total_for_povert
)

popsize_and_poverty <- popsize_and_poverty |>
   select(GEOID, popsize, proportion_in_poverty)
```

What type of data do you have?



Picking a way to visualize your data



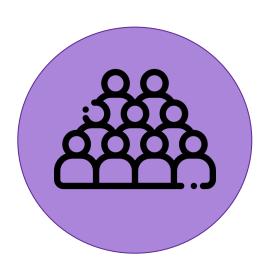
Goals of data visualization



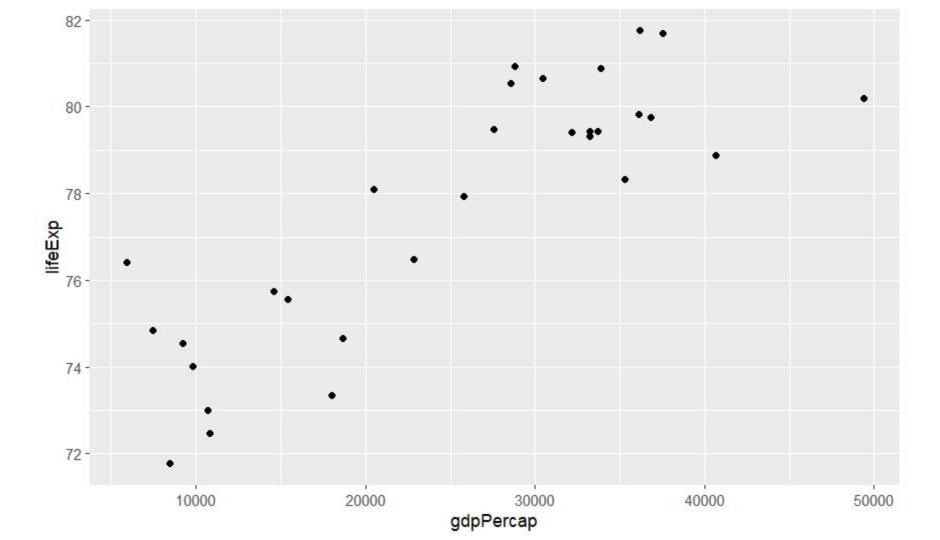




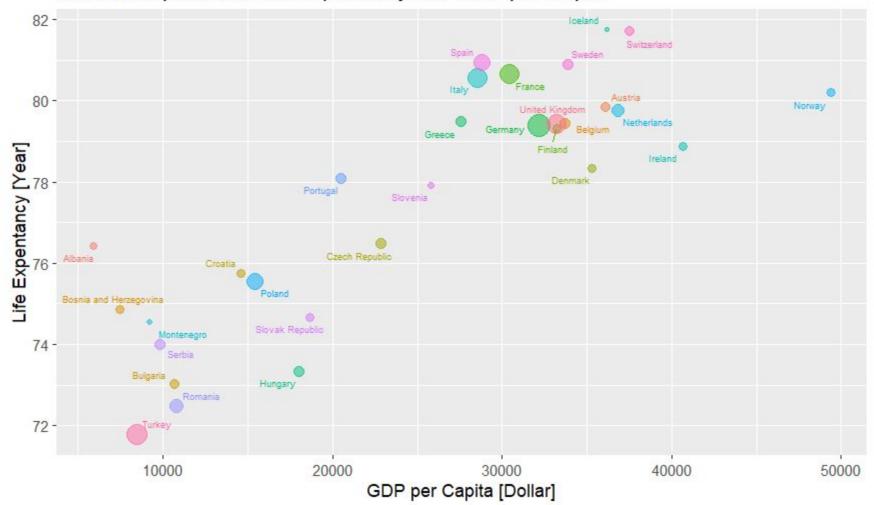
Storytelling



Accessibility



Relationship between life expentancy and GDP per Capita

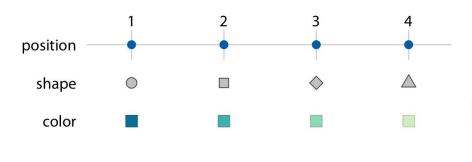


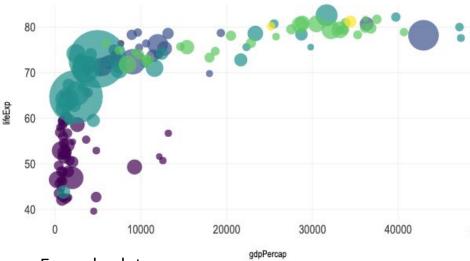
Scale

"A scale defines a unique mapping between data and aesthetics."

Rule of thumb: Each data value must have a unique scale.

3 different scales:



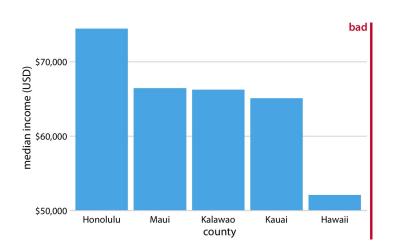


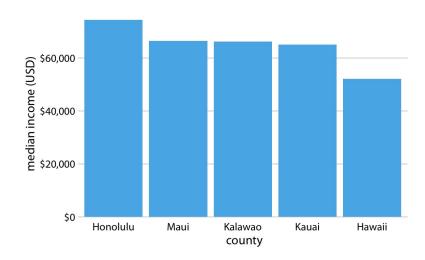
Source: https://clauswilke.com/dataviz/aesthetic-mapping.html
Interactive bubble chart – the R Graph Gallery (r-graph-gallery.com)

Example plot:

Axis

Quantitative axes:





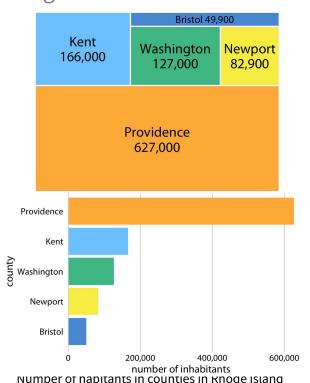
Example plot: 2 bar graphs of median income of 5 counties of Hawaii the state

Why do the income gaps look so different in the two graphs?

Source: https://clauswilke.com/dataviz/proportional-ink.html

Axis

Categorical axes:

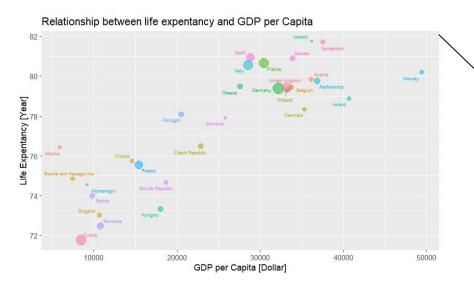


Source: https://clauswilke.com/dataviz/proportional-ink.html https://clauswilke.com/dataviz/proportional-ink.html</a

Dual axes: two different y axes

```
scale = 15
pkpd \leftarrow ggplot(res, aes(x = time, y = CP)) +
   geom_line(aes(color = "Drug Concentration")) +
  geom line(aes(y = RESP/scale, color = "Biomarker (IU/mL")) +
   scale_x_continuous(breaks = seq(0, 336, 24)) +
   scale_y_continuous(sec.axis = sec_axis(~.*scale, name="Biomarker (IU/mL)")) +
   labs(x = "Time (hr)", y = "Concentration (mg/L)", color = "") +
   scale_color_manual(values = c("orange2", "gray30"))
print(pkpd)
Concentration (mg/L)
                                                                         Biomarker (IU/mL
                                                              (IU/mL)
                                                                         Drug Concentration
        24 48 72 96 120 144 168 192 216 240 264 288 312 336
                            Time (hr)
```

Style for Accessibility



devtools::install_github("AndreaCirilloAC/paletter")

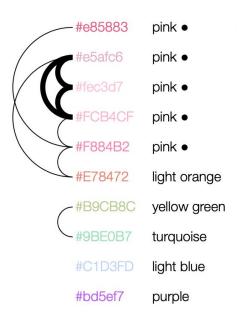
#E8588E	#E5AFC6	#FEC3D7	#BD5EF7
#FCB4CF	#B9CB8C	#E78472	#F884B2
#9BE0B7	#C1D3FD		

Viz Palette by Elijah Meeks & Susie Lu

COLOR REPORT

Arcs link colors difficult to tell apart as:

- Lines or small points
- Medium areas
- Large areas



 Minimize name conflicts for categorical palettes

```
install.packages("RColorBrewer")
library(RColorBrewer)
display.brewer.all(colorblindFriendly = TRUE)
```



install.packages("wesanderson") library(wesanderson)

```
> names(wes_palettes)
                                       "Rushmore1"
 [1] "BottleRocket1"
                      "BottleRocket2"
                                                         "Rushmore"
                      "Royal2"
                                       "Zissou1"
                                                         "Darjeeling1"
 [5] "Royal1"
 [9] "Darjeeling2" "Chevalier1"
                                       "FantasticFox1"
                                                         "Moonrise1"
[13] "Moonrise2"
                      "Moonrise3"
                                       "Cavalcanti1"
                                                         "GrandBudapest1"
[17] "GrandBudapest2" "IsleofDogs1"
                                        "IsleofDogs2"
```

wes_palette("FantasticFox1")



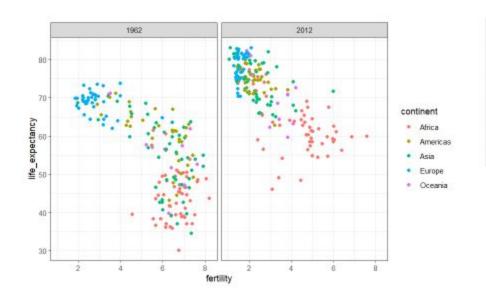
Style for Accessibility (cont.)



```
## This chart has title 'Relationship between life expentancy and GDP per Capita'.
## It has x-axis 'GDP per Capita [Dollar]' with labels 10000, 20000, 30000, 40000 and 50000.
## It has y-axis 'Life Expentancy [Year]' with labels 72, 74, 76, 78, 80 and 82.
## In this chart colour is used to show country. The legend that would normally indicate this has been hidden.
## In this chart size is used to show pop. The legend that would normally indicate this has been hidden.
## Layer 1 is a set of 30 points.
## Layer 1 has alpha set to 0.5.
## Layer 2 is a textrepel graph that VI can not process.
## Layer 2 has size set to 2.
```

facet_wrap

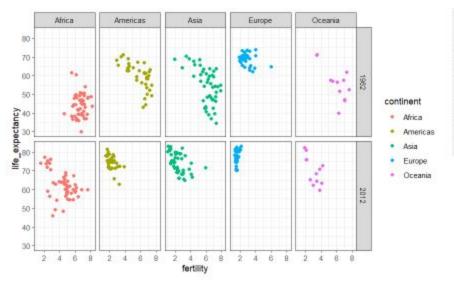
Faceting can group and visualize each subset. Facet_wrap can only visualize non-empty univariate.



```
filter(gapminder, year%in%c(1962, 2012)) |> ggplot(aes(fertility, life_expectancy, col = continent)) + geom_point() + facet_wrap(.~year)
```

facet_grid

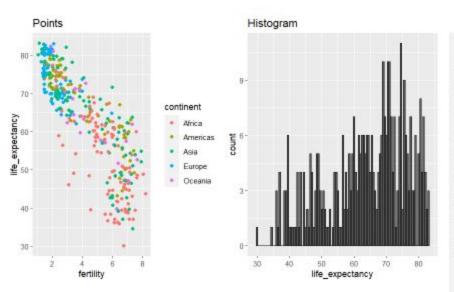
Facet_grid can visualize bivariate even some are empty.



```
filter(gapminder, year%in%c(1962, 2012)) |> ggplot(aes(fertility, life_expectancy, col = continent)) + geom_point() + facet_grid(year~continent)
```

grid.arrange

grid.arrange can layout multiple graphs on the same page

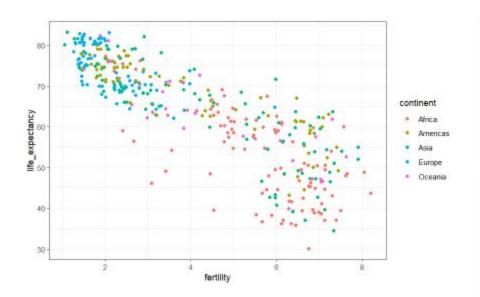


```
p1=filter(gapminder, year%in%c(1962, 2012)) |> ggplot(aes(fertility, life_expectancy, col = continent)) + geom_point() + ggtitle("Points")

p2=filter(gapminder, year%in%c(1962, 2012)) |> ggplot(aes(life_expectancy)) + geom_histogram(binwidth = 0.5, color = "black") + ggtitle("Histogram")

grid.arrange(p1, p2, ncol = 2)```
```

ggplot2 can build a complex plot a layer at a time. Each layer can come from a different dataset and have a different aesthetic mapping. Layers are created using geom_* or stat_*



```
filter(gapminder, year%in%c(1962,
2012)) |> ggplot(aes(fertility,
life_expectancy, col = continent)) +
geom_point()
```

```
p + layer(
mapping = NULL,
data = NULL,
geom = "point",
stat = "identity",
position = "identity"
)
```

Geom*

One variable:

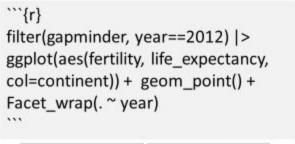
- · geom_bar(): display distribution of discrete variable.
- geom_histogram(): bin and count continuous variable, display with bars.
- geom_density(): smoothed density estimate.
- geom_dotplot(): stack individual points into a dot plot.
- · geom_freqpoly(): bin and count continuous variable, display with lines.

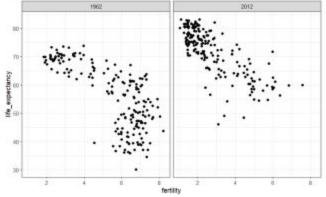
Two variables:

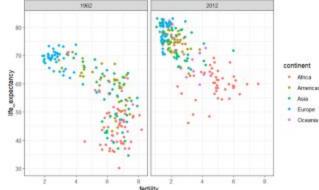
- geom_point(): scatterplot.
- geom_quantile(): smoothed quantile regression.
- geom_rug(): marginal rug plots.
- · geom_smooth(): smoothed line of best fit.
- geom_text(): text labels.

Aesthetic Mapping*

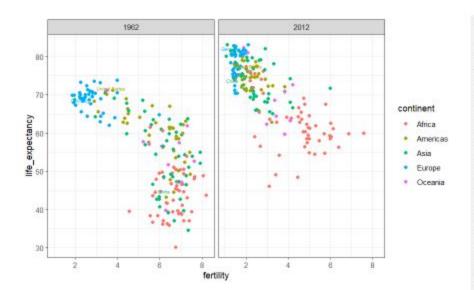
```
filter(gapminder, year==2012) |>
ggplot(aes(fertility, life_expectancy)) +
geom_point() +
Facet_wrap(. ~ year)
```







Data



```
```{r}
highlight <- c("Germany", "China",
"United States")
filter(gapminder, year %in% c(1962,
2012)) %>% ggplot(aes(fertility,
life expectancy, color = continent)) +
geom point() +
geom_text_repel(size = 2,
show.legend = FALSE, aes(fertility,
life expectancy, label=country), data
= filter(gapminder, year %in% c(1962,
2012) & country %in% highlight)) +
facet_grid(. ~ year)
```

#### Stats\*

```
stat_bin(): geom_bar(), geom_freqpoly(), geom_histogram()
stat_bin2d(): geom_bin2d()
stat_bindot(): geom_dotplot()
stat_binhex(): geom_hex()
stat_boxplot(): geom_boxplot()
stat_contour(): geom_contour()
stat_quantile(): geom_quantile()
stat_smooth(): geom_smooth()
stat_sum(): geom_count()
```

#### Position\*

#### Three position adjustments for points:

- position\_nudge(): move points by a fixed offset.
- position\_jitter(): add a little random noise to every position.
- position\_jitterdodge(): dodge points within groups, then add a little random noise.

#### Three positions adjustments for bars:

- · position\_stack(): stack overlapping bars (or areas) on top of each other.
- position\_fill(): stack overlapping bars, scaling so the top is always at 1.
- position\_dodge(): place overlapping bars (or boxplots) side-by-side.

### Saving and Exporting Visualizations

If using ggplot:

```
ggsave (
 ← The name of the file locally
 ex. "mygraphic.png"
 filename,
 plot = last_plot(), \leftarrow \underline{Plot to export}
 ex. plot1
 device = \overline{\text{NULL}}, \leftarrow Type of export
 ex. "Png", "ipeg"
 ← File location to be saved
 ex. "mydocuments/folder1"
 path = NULL,
 ← Multiplicative factor
 scale = 1.
 width = NA,
 ← Dimensions of graphic
 ex. 3,5,10 (inches, pixels...)
 height = NA,
 units = c("in", "cm", "mm", "px"), \leftarrow unit of size
 ex. "retina",300, "screen"
 dpi = 300, \leftarrow Set resolution
 limitsize = TRUE, ← max image size of 50 x 50 inches
 bg = NULL, ← Background color
 ex. "gray"
```

# **Interactability**

If looking for individuals to be able to change elements in your visualization, try creating an **RShiny App** 

#### **Shiny App Example**

- Easy way to share ggplot figures, and others, in a way that allows users to change elements of the visualization easily
- See more information here: R Shiny Tutorial



## **Key Takeaways**

- Don't take shortcuts in your code
- Be intentional with your visualizations
- Making good visualizations of data takes time
- Making accessible visualizations increases impact by broadening audience