

R you ready?

**IntRo to RStudio and R Markdown
for open data and reproducibility**

Unit 7:

Let's get plotting!

Mason A. Wirtz

 [@WirtzMason](https://twitter.com/WirtzMason)

GGPLOT ALL THE THINGS



ggplot2: What is it?

ggplot2 is a package from the *tidyverse* and is one of the most-used plotting packages in R

ggplot2 plotting system →

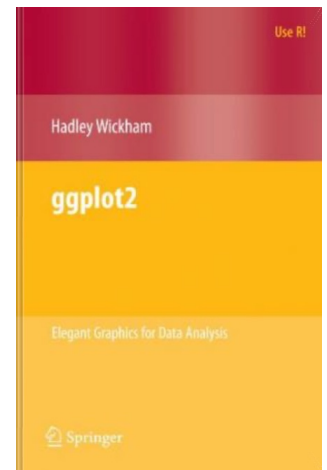
<https://bookdown.org/rdpeng/exdata/the-ggplot2-plotting-system-part-1.html#ggplot2-hello-world>

ggplot2: Elegant Graphics for Data Analysis →

<https://1lib.at/book/704124/daea5e>

ggplot2 cheatsheet →

<https://www.rstudio.com/resources/cheatsheets/>



ggplot2: Why?

- **Set of independent components that can be composed in different ways**
- **Not limited to pre-specified graphics**
- **Plots can be built iteratively**
- **i.e. plots are built using a layering principle**
- **beautiful faceting or multipanel plots**
- **can easily change or update plots**

ggplot2: Terminology

- **Aesthetic mappings (aes())** → describe how variables in the data are mapped to aesthetic attributes
- **Geometric objects (geom)** → represents what we actually see on the plot
- **Statistical transformations (stats)** → summarise data in many useful ways
- **Scales** → map values in the data space to values in an aesthetic space
- **Faceting** → break up the data into subsets

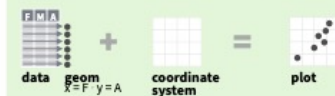
ggplot2: cheat sheet

Data visualization with ggplot2 : : CHEAT SHEET



Basics

ggplot2 is based on the **grammar of graphics**, the idea that you can build every graph from the same components: a **data** set, a **coordinate system**, and **geoms**—visual marks that represent data points.



To display values, map variables in the data to visual properties of the geom (**aesthetics**) like **size**, **color**, and **x** and **y** locations.



Complete the template below to build a graph.

```
ggplot(data = <DATA>) +  
  <GEOM_FUNCTION>(mapping = aes(<MAPPINGS>),  
  stat = <STAT>, position = <POSITION>) +  
  <COORDINATE_FUNCTION> +  
  <FACET_FUNCTION> +  
  <SCALE_FUNCTION> +  
  <THEME_FUNCTION>
```

required
Not required, sensible defaults supplied

ggplot(data = mpg, aes(x = cty, y = hwy)) Begins a plot that you finish by adding layers to. Add one geom function per layer.

last_plot() Returns the last plot.

ggsave("plot.png", width = 5, height = 5) Saves last plot as 5' x 5' file named "plot.png" in working directory. Matches file type to file extension.

Aes Common aesthetic values.

color and **fill** - string ("red", "#RRGGBB")

linetype - integer or string (0 = "blank", 1 = "solid", 2 = "dashed", 3 = "dotted", 4 = "dotdash", 5 = "longdash", 6 = "twodash")

lineend - string ("round", "butt", or "square")

linejoin - string ("round", "mitre", or "bevel")

size - integer (line width in mm)

shape - integer/shape name or a single character ("a")



Geoms

Use a geom function to represent data points, use the geom's aesthetic properties to represent variables. Each function returns a layer.

GRAPHICAL PRIMITIVES

a <- ggplot(economics, aes(date, unemployment))
b <- ggplot(seals, aes(x = long, y = lat))

a + geom_blank() and **a + expand_limits()**
Ensure limits include values across all plots.

b + geom_curve(aes(yend = lat + 1, xend = long + 1), curvature = 1) - x, yend, y, yend, alpha, angle, color, curvature, linetype, size

a + geom_path(lineend = "butt", linejoin = "round", linemitre = 1) - x, y, alpha, color, group, linetype, size

a + geom_polygon(aes(alpha = 50)) - x, y, alpha, color, fill, group, subgroup, linetype, size

b + geom_rect(aes(xmin = long, ymin = lat, xmax = long + 1, ymax = lat + 1)) - xmin, xmax, ymin, ymax, alpha, color, fill, linetype, size

a + geom_ribbon(aes(ymin = unemployment - 900, ymax = unemployment + 900)) - x, ymax, ymin, alpha, color, fill, group, linetype, size

LINE SEGMENTS

common aesthetics: x, y, alpha, color, linetype, size

b + geom_abline(aes(intercept = 0, slope = 1))
b + geom_hline(aes(yintercept = lat))
b + geom_vline(aes(xintercept = long))

b + geom_segment(aes(yend = lat + 1, xend = long + 1))
b + geom_spoke(aes(angle = 1:1155, radius = 1))

ONE VARIABLE continuous

c <- ggplot(mpg, aes(hwy)); c2 <- ggplot(mpg)

c + geom_area(stat = "bin")
x, y, alpha, color, fill, linetype, size

c + geom_density(kernel = "gaussian")
x, y, alpha, color, fill, group, linetype, size, weight

c + geom_dotplot
x, y, alpha, color, fill

c + geom_freqpoly
x, y, alpha, color, group, linetype, size

c + geom_histogram(binwidth = 5)
x, y, alpha, color, fill, linetype, size, weight

c2 + geom_qq(aes(sample = hwy))
x, y, alpha, color, fill, linetype, size, weight

discrete

d <- ggplot(mpg, aes(fit))

d + geom_bar
x, alpha, color, fill, linetype, size, weight

TWO VARIABLES both continuous

e <- ggplot(mpg, aes(cty, hwy))

e + geom_label(aes(label = cty), nudge_x = 1, nudge_y = 1) - x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust

e + geom_point
x, y, alpha, color, fill, shape, size, stroke

e + geom_quantile
x, y, alpha, color, group, linetype, size, weight

e + geom_rug(sides = "bl")
x, y, alpha, color, linetype, size

e + geom_smooth(method = lm)
x, y, alpha, color, fill, group, linetype, size, weight

e + geom_text(aes(label = cty), nudge_x = 1, nudge_y = 1) - x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust

one discrete, one continuous

f <- ggplot(mpg, aes(class, hwy))

f + geom_col
x, y, alpha, color, fill, group, linetype, size

f + geom_boxplot
x, y, lower, middle, upper, ymax, ymin, alpha, color, fill, group, linetype, shape, size, weight

f + geom_dotplot(binaxis = "y", stackdir = "center")
x, y, alpha, color, fill, group

f + geom_violin(scale = "area")
x, y, alpha, color, fill, group, linetype, size, weight

both discrete

g <- ggplot(diamonds, aes(cut, color))

g + geom_count
x, y, alpha, color, fill, shape, size, stroke

e + geom_jitter(height = 2, width = 2)
x, y, alpha, color, fill, shape, size

THREE VARIABLES

sealsSz <- with(seals, sqrt(delta_long^2 + delta_lat^2)); l <- ggplot(seals, aes(long, lat))

l + geom_contour(aes(z = z))
x, y, z, alpha, color, group, linetype, size, weight

l + geom_contour_filled(aes(fill = z))
x, y, alpha, color, fill, group, linetype, size, subgroup

continuous bivariate distribution

h <- ggplot(diamonds, aes(carat, price))

h + geom_bin2d(binwidth = c(0.25, 500))
x, y, alpha, color, fill, linetype, size, weight

h + geom_density_2d
x, y, alpha, color, group, linetype, size

h + geom_hex
x, y, alpha, color, fill, size

continuous function

i <- ggplot(economics, aes(date, unemployment))

i + geom_area
x, y, alpha, color, fill, linetype, size

i + geom_line
x, y, alpha, color, group, linetype, size

i + geom_step(direction = "hv")
x, y, alpha, color, group, linetype, size

visualizing error

df <- data.frame(grp = c("A", "B"), fit = 4.5, se = 1.2)
j <- ggplot(df, aes(grp, fit, ymin = fit - se, ymax = fit + se))

j + geom_crossbar(fatten = 2) - x, y, ymax, ymin, alpha, color, fill, group, linetype, size

j + geom_errorbar
x, y, ymax, ymin, alpha, color, group, linetype, size, width
Also **geom_errorbarh**()

j + geom_linerange
x, y, ymax, ymin, alpha, color, group, linetype, size

j + geom_pointrange
x, y, ymax, ymin, alpha, color, fill, group, linetype, shape, size

maps

data <- data.frame(murder = USArrests\$Murder, state = tolower(rownames(USArrests)))

map <- map_data("state")
k <- ggplot(data, aes(fill = murder))

k + geom_map(aes(map_id = state), map = map) + **expand_limits**(x = map\$long, y = map\$lat)
map_id, alpha, color, fill, linetype, size

ggplot2: Introduction

The workhorse function:

- `ggplot()`



Takes the arguments:

- `data = data frame`
- `mapping = aes()`

ONE variable:

`data frame %>%`

`ggplot(aes(x = variable_x_axis))`

ggplot2: Introduction

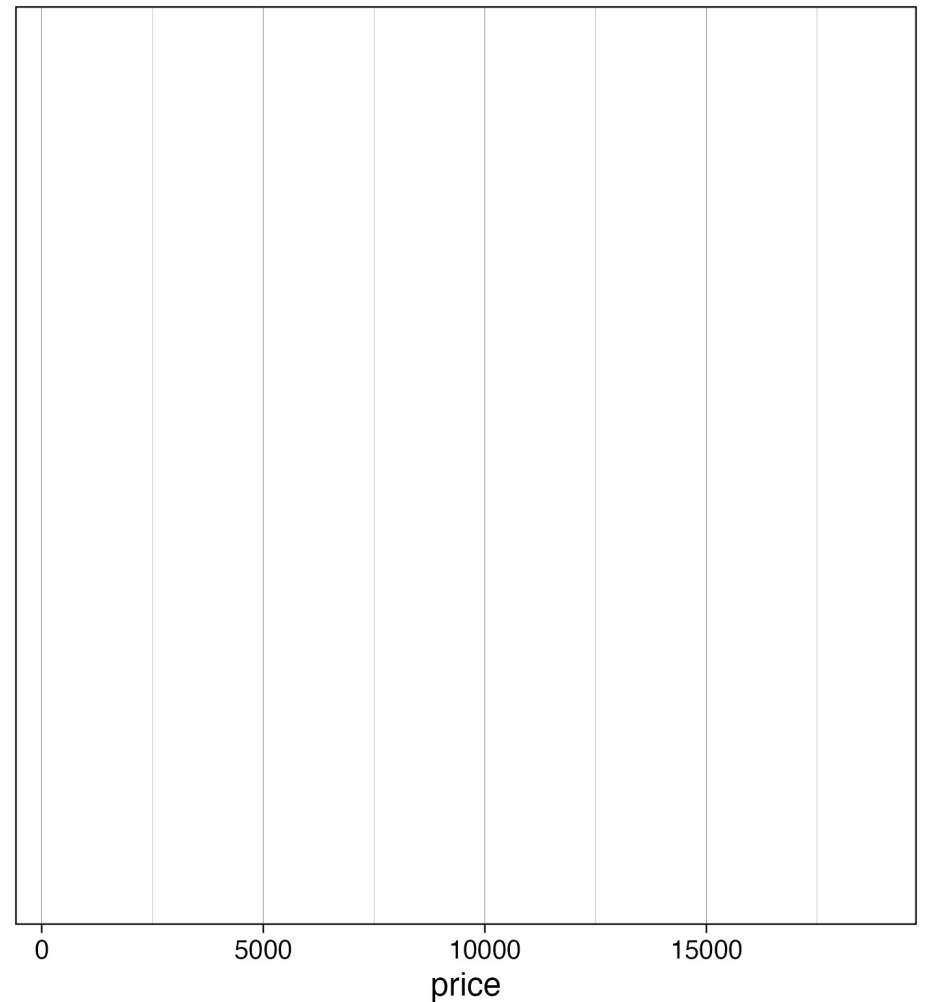
Data frame

```
diamonds %>%
```

```
ggplot(aes(x = price))
```

Add aesthetics

We defined our x-axis,
but we have not **PLOTTED**
anything!



So how do we plot?



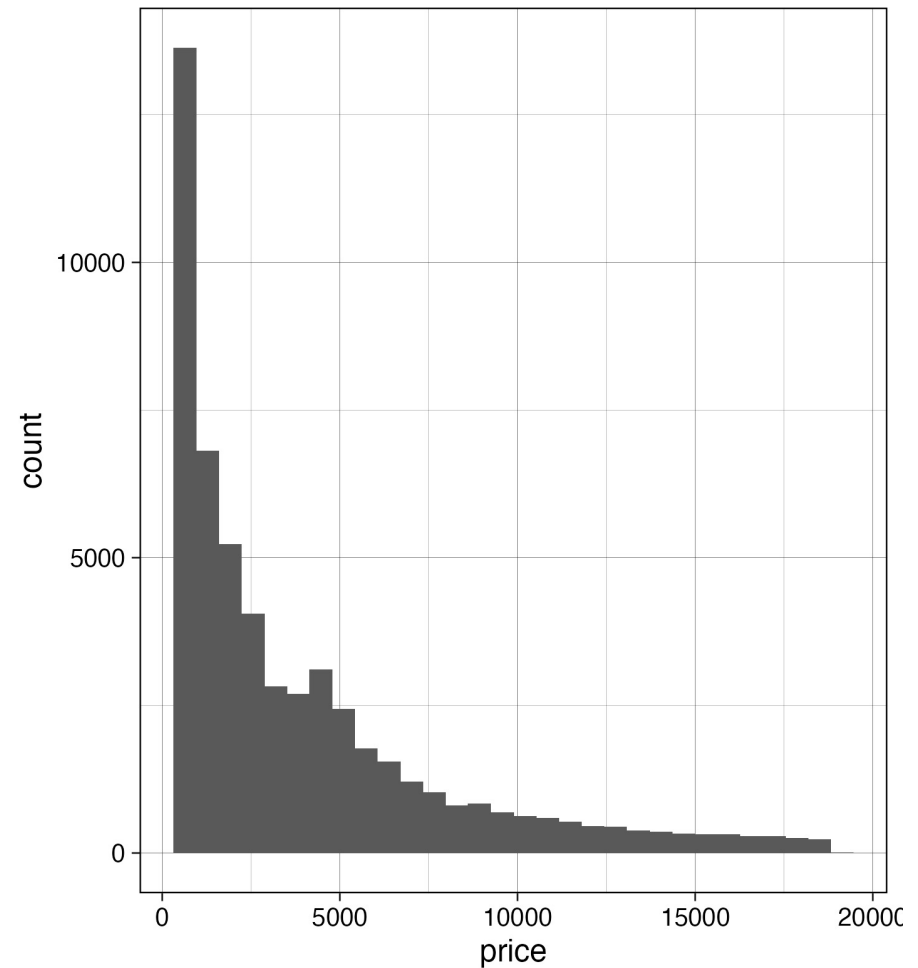
ggplot2: Plots of one variable

One CONTINUOUS variable:

- **Histogram:** *geom_histogram()*
- **Density plot:** *geom_density()*

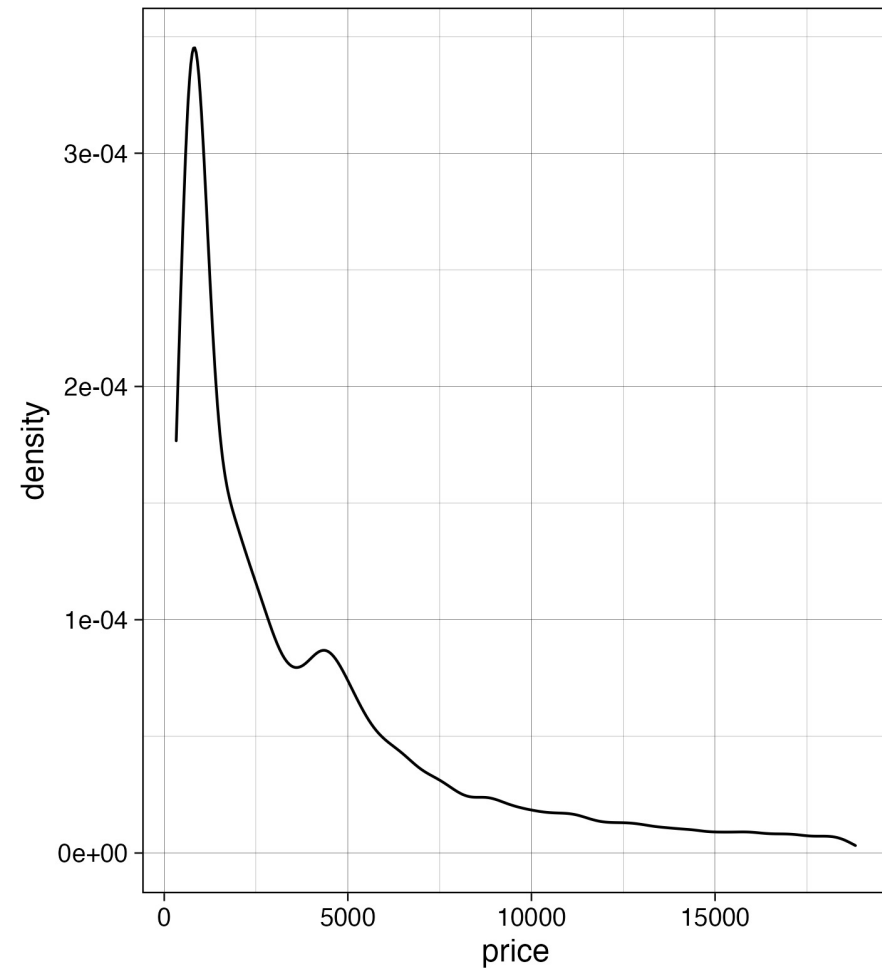
ggplot2: geom_histogram()

```
diamonds %>%  
  ggplot(aes(x = price)) +  
  geom_histogram()
```



ggplot2: geom_density()

```
diamonds %>%  
  ggplot(aes(x = price)) +  
  geom_density()
```



ggplot2: Plots of one variable

One CATEGORICAL variable:

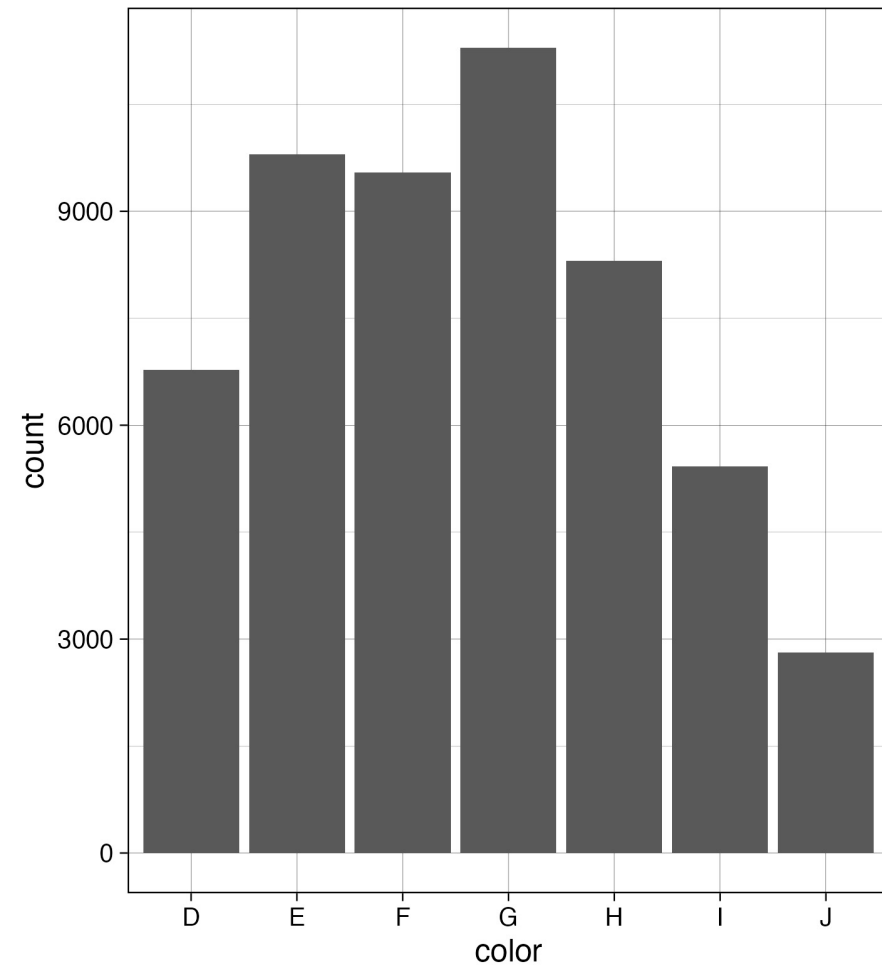
- Barplot: *geom_bar()*

ggplot2: geom_bar()

`diamonds %>%`

`ggplot(aes(x = color)) +`

`geom_bar()`



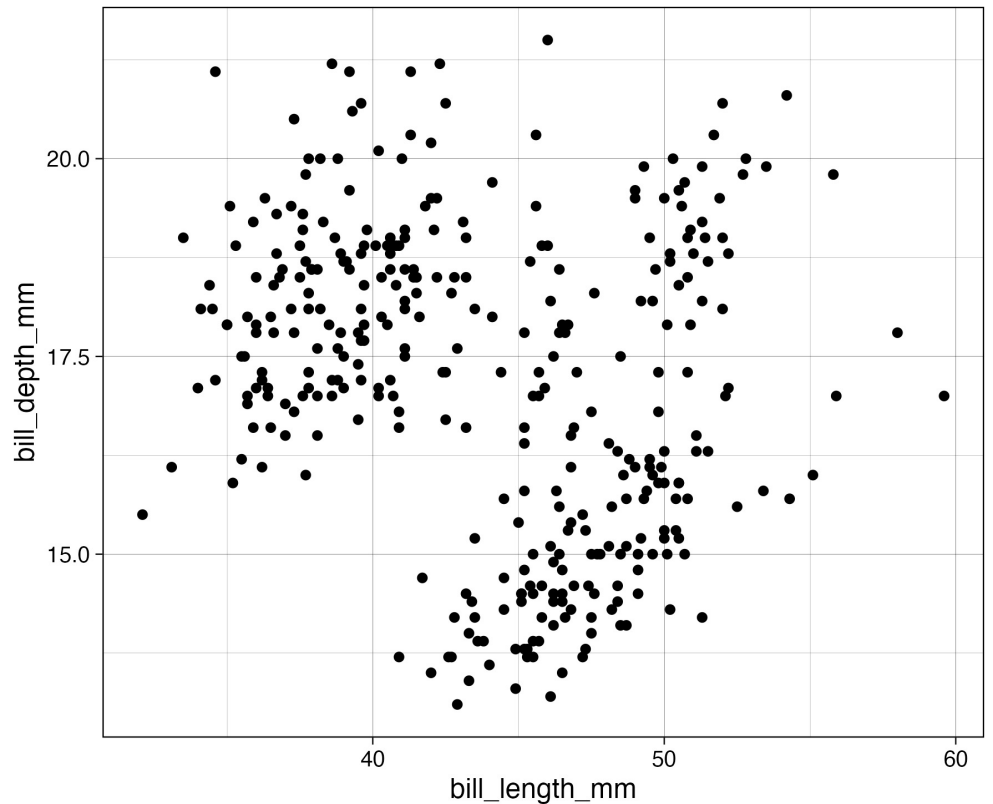
ggplot2: Plots of two variables

Two CONTINUOUS variables:

- **Scatter plot:** *geom_point()*
- **Scatter plot:** *geom_jitter()* → jittered (scattered) data points
- **Smoother:** *geom_smooth()* → linear and nonlinear lines

ggplot2: geom_point()

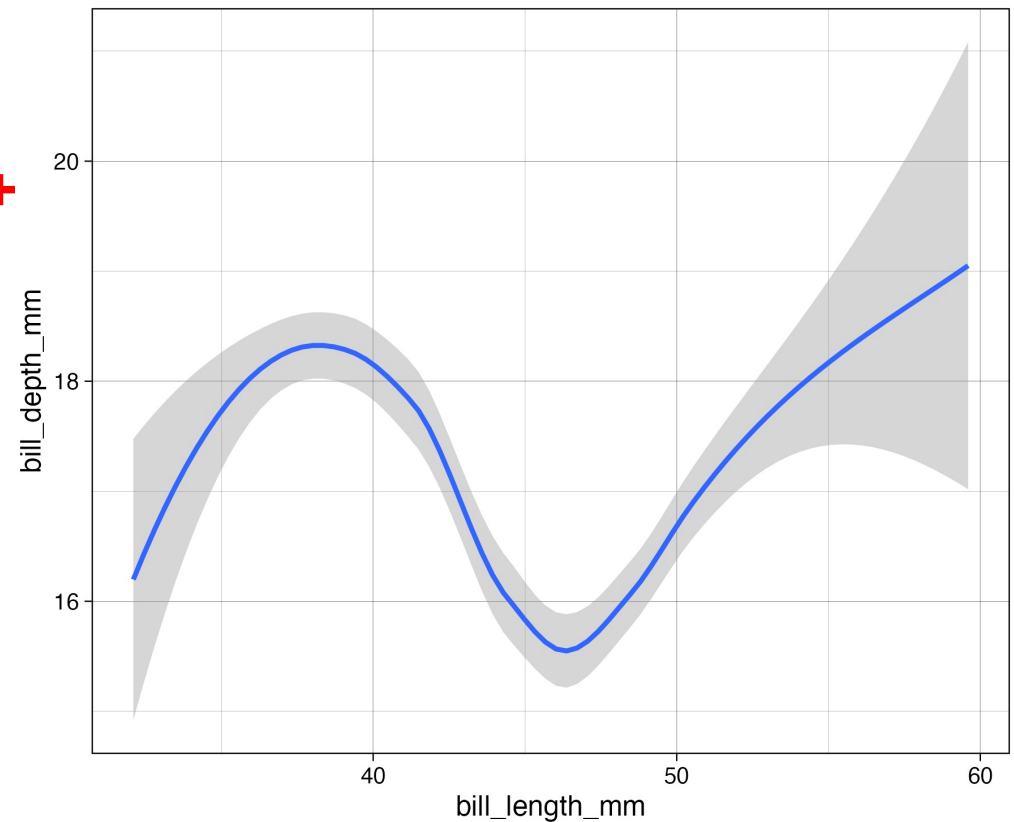
```
penguins %>%  
  ggplot(aes(x = bill_length_mm,  
             y = bill_depth_mm)) +  
  geom_point()
```



ggplot2: geom_smooth()

```
penguins %>%  
  ggplot(aes(x = bill_length_mm,  
             y = bill_depth_mm)) +  
  geom_smooth()
```

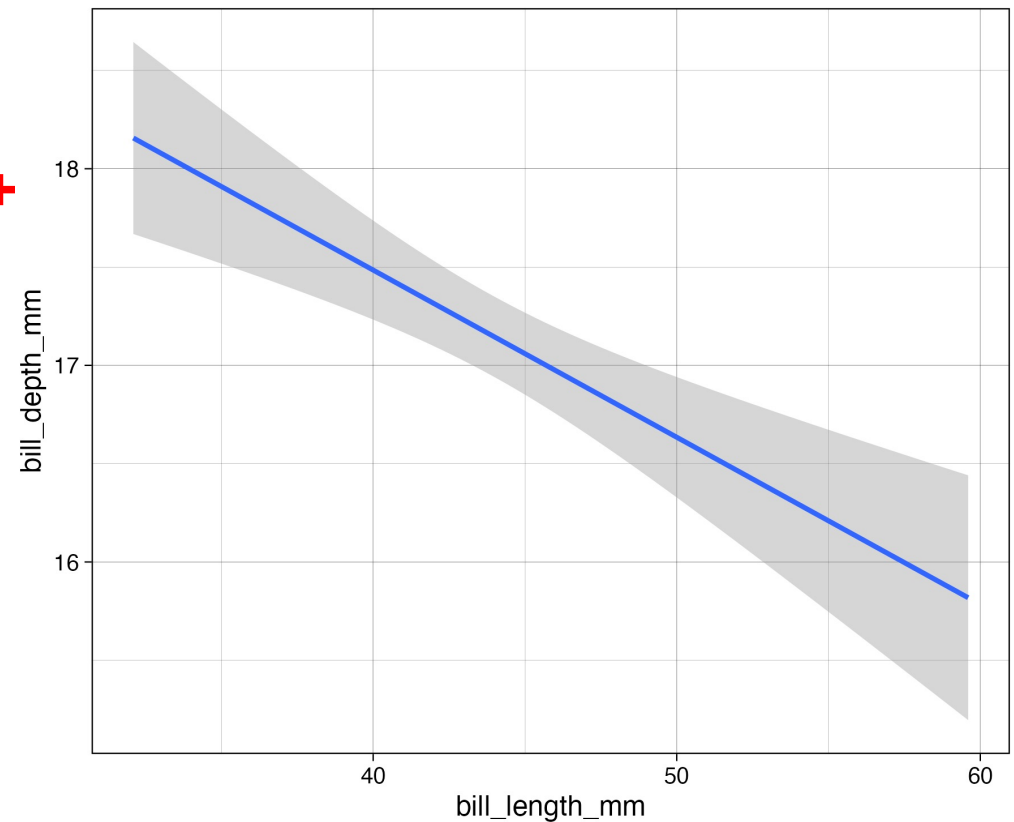
Default method:
Generalized additive model
(GAM) with cubic spline
 $y \sim s(x, bs = "cs")$



ggplot2: geom_smooth()

```
penguins %>%  
  ggplot(aes(x = bill_length_mm,  
             y = bill_depth_mm)) +  
  geom_smooth(method = "lm")
```

Change method:
method = "lm"
→ Linear model



ggplot2: Plots of two variables

CATEGORICAL x and CONTINUOUS y:

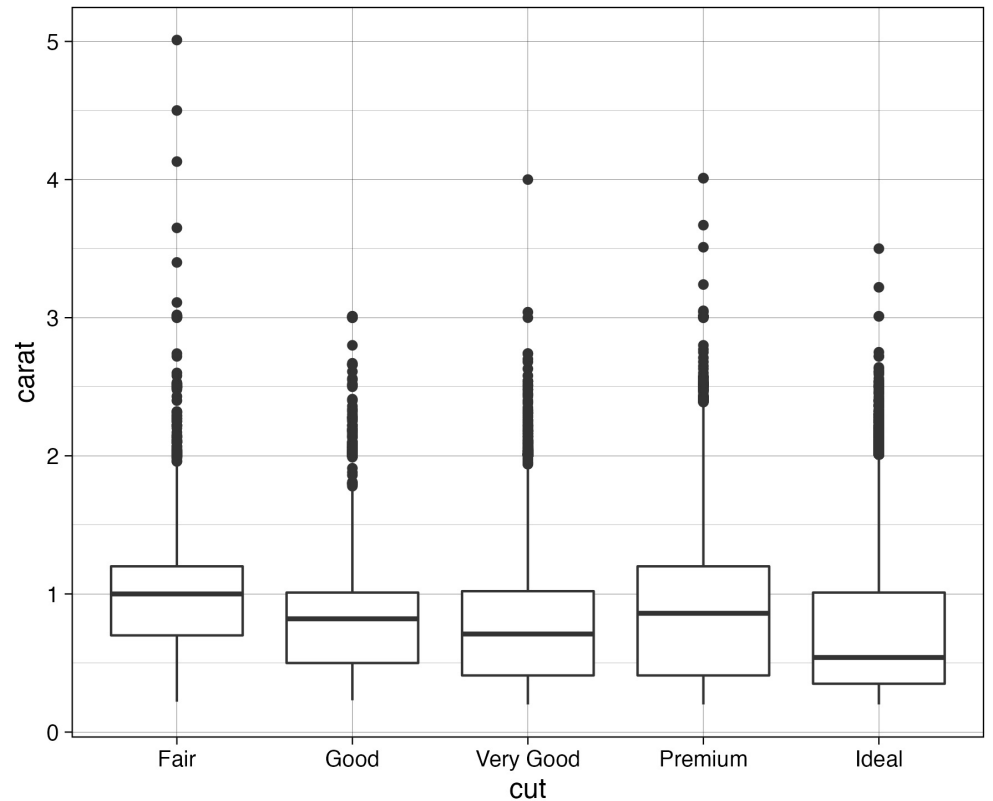
- **Boxplot:** *geom_boxplot()*
- **Barplot:** *geom_bar(stat = "identity")*
- **Violin plot:** *geom_violin()*

ggplot2: geom_boxplot()

```
diamonds %>%
```

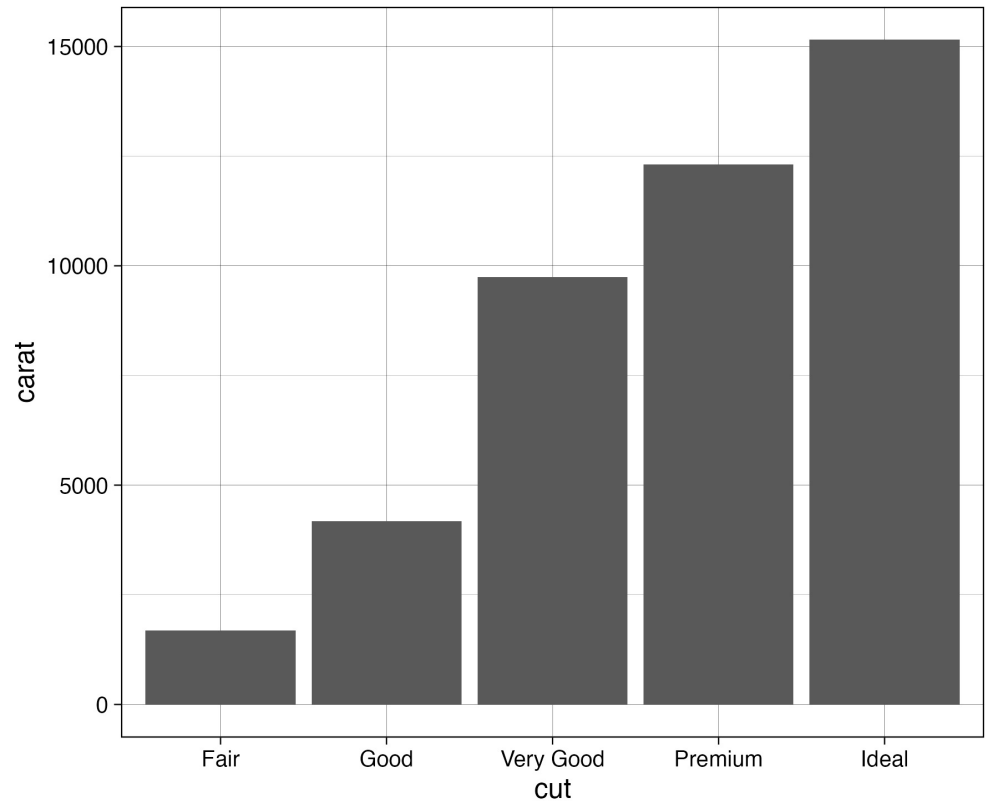
```
  ggplot(aes(x = cut,  
             y = carat)) +
```

```
  geom_boxplot()
```



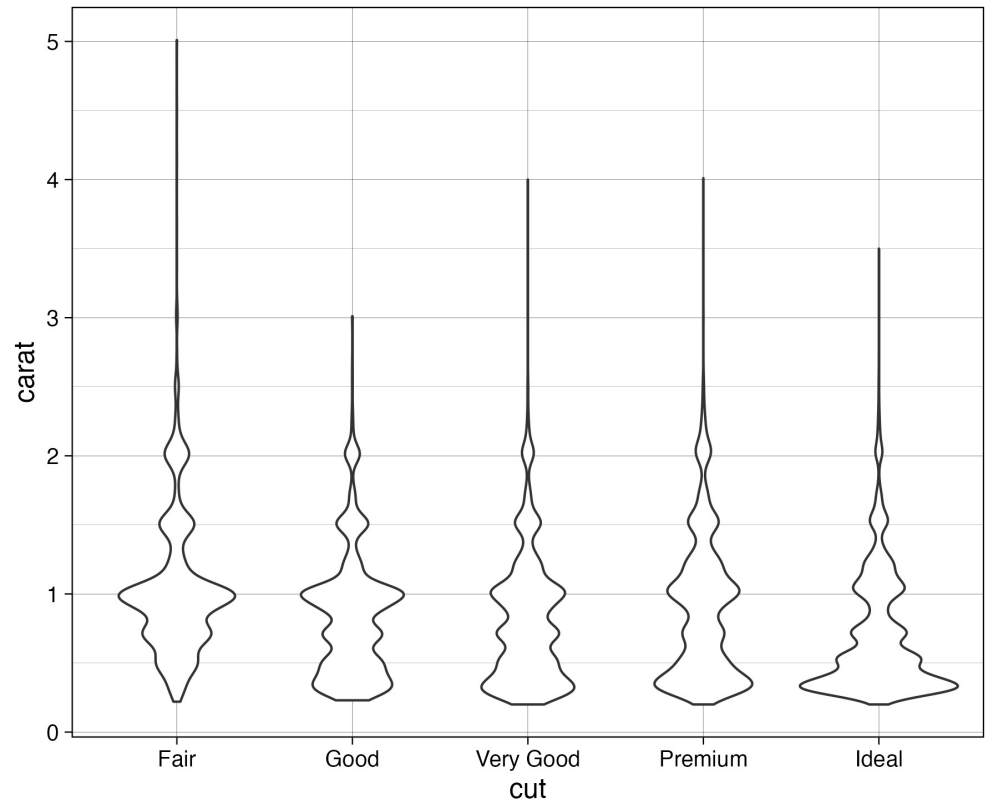
ggplot2: geom_bar()

```
diamonds %>%  
  ggplot(aes(x = cut,  
             y = carat)) +  
  geom_bar(stat = "identity")
```



ggplot2: geom_violin()

```
diamonds %>%  
  ggplot(aes(x = cut,  
             y = carat)) +  
  geom_violin()
```



AESTHETIC S



ggplot2: Aesthetics

Aesthetics:

- **Size**

Change the size of e.g. data points

- **Shape**

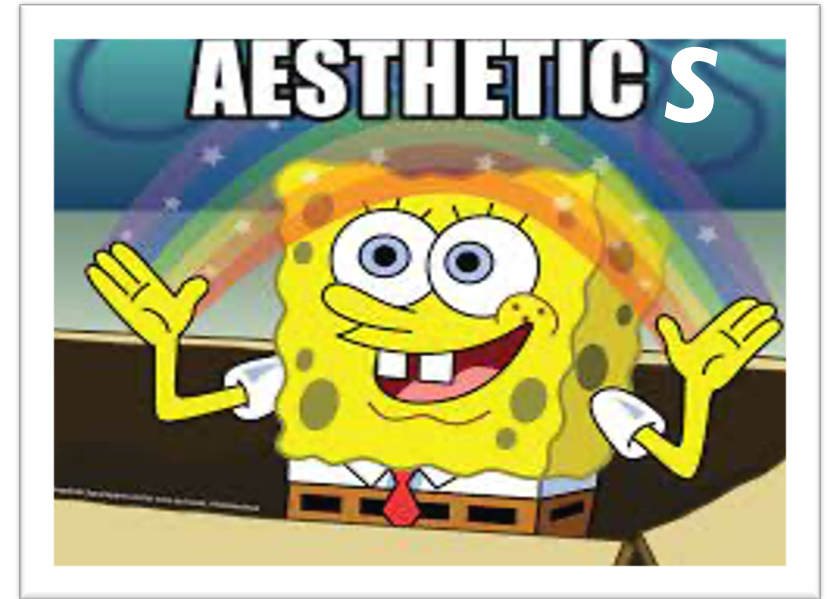
Change the shape of e.g. data points

- **Color**

Change the color of e.g. data points

- **Fill**

Change the fill of geom, e.g. a boxplot

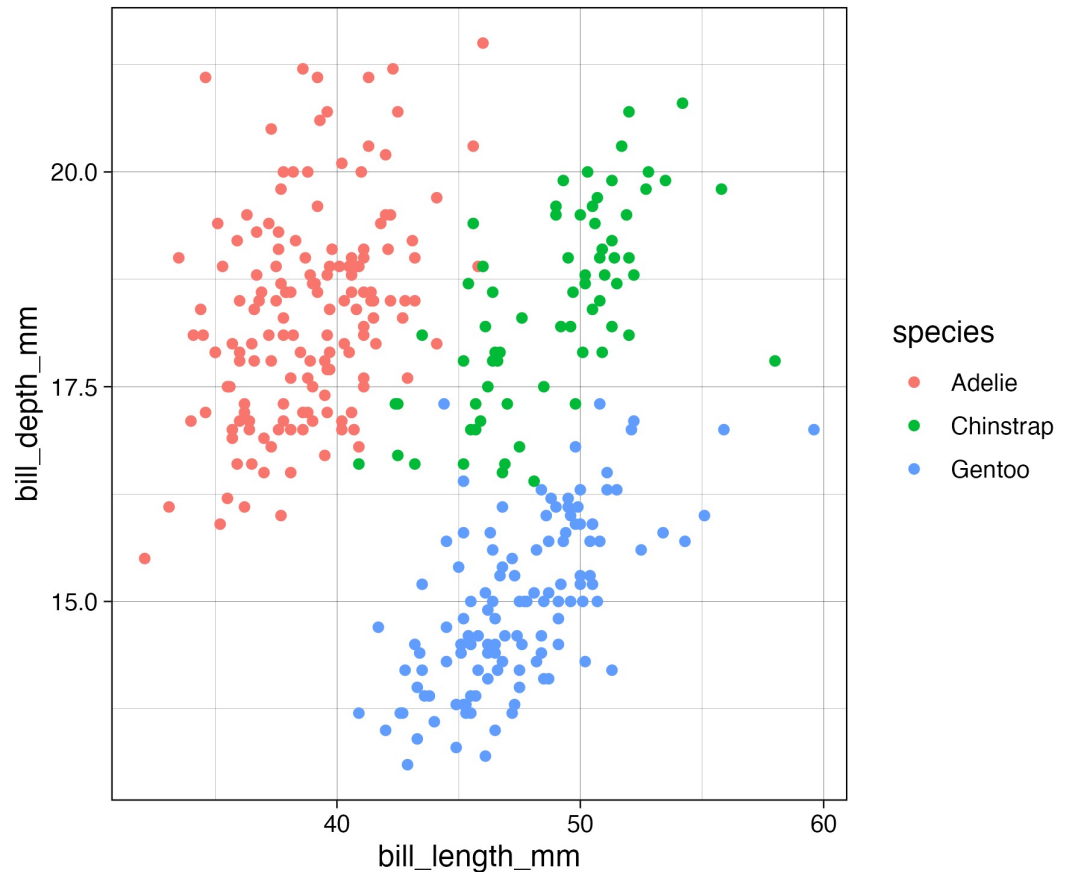


ggplot2: geom_point() + color

```
penguins %>%
```

```
  ggplot(aes(x = bill_length_mm,  
             y = bill_depth_mm,  
             color = species)) +
```

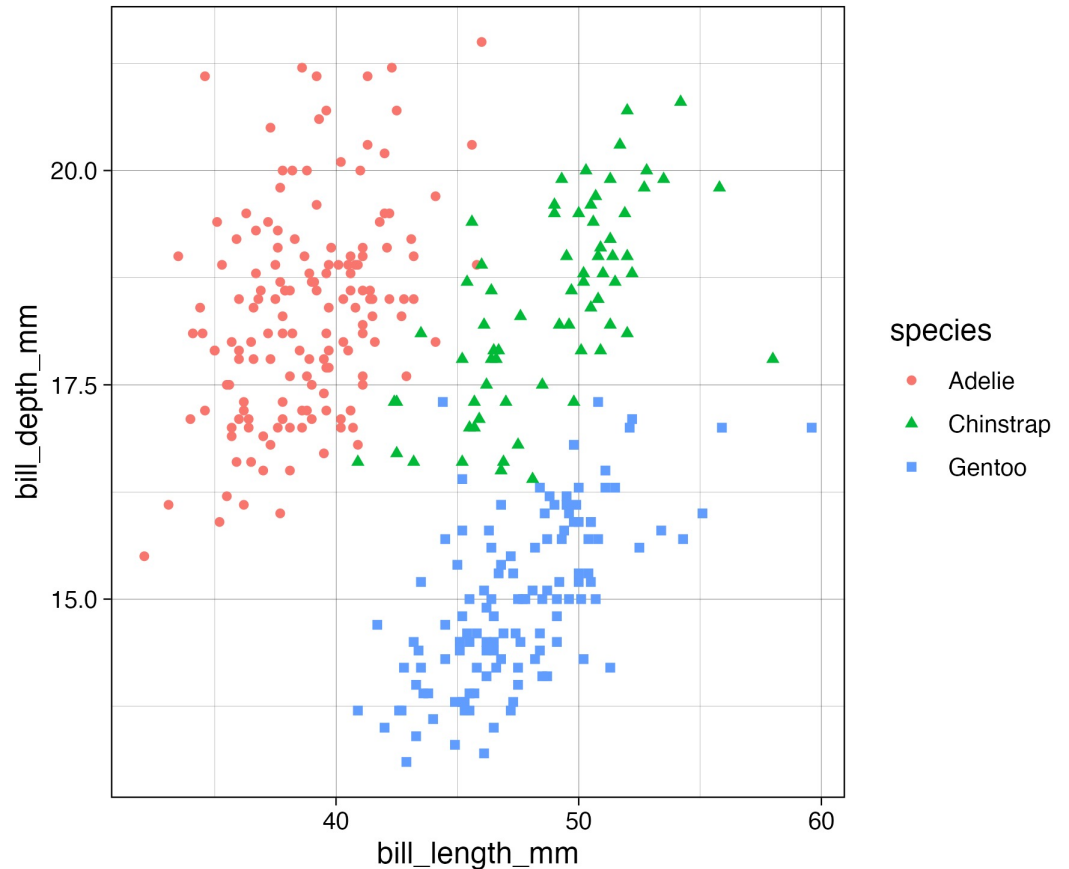
```
  geom_point()
```



ggplot2: geom_point() + color + shape

penguins %>%

```
ggplot(aes(x = bill_length_mm,  
            y = bill_depth_mm,  
            color = species,  
            shape = species)) +  
geom_point()
```

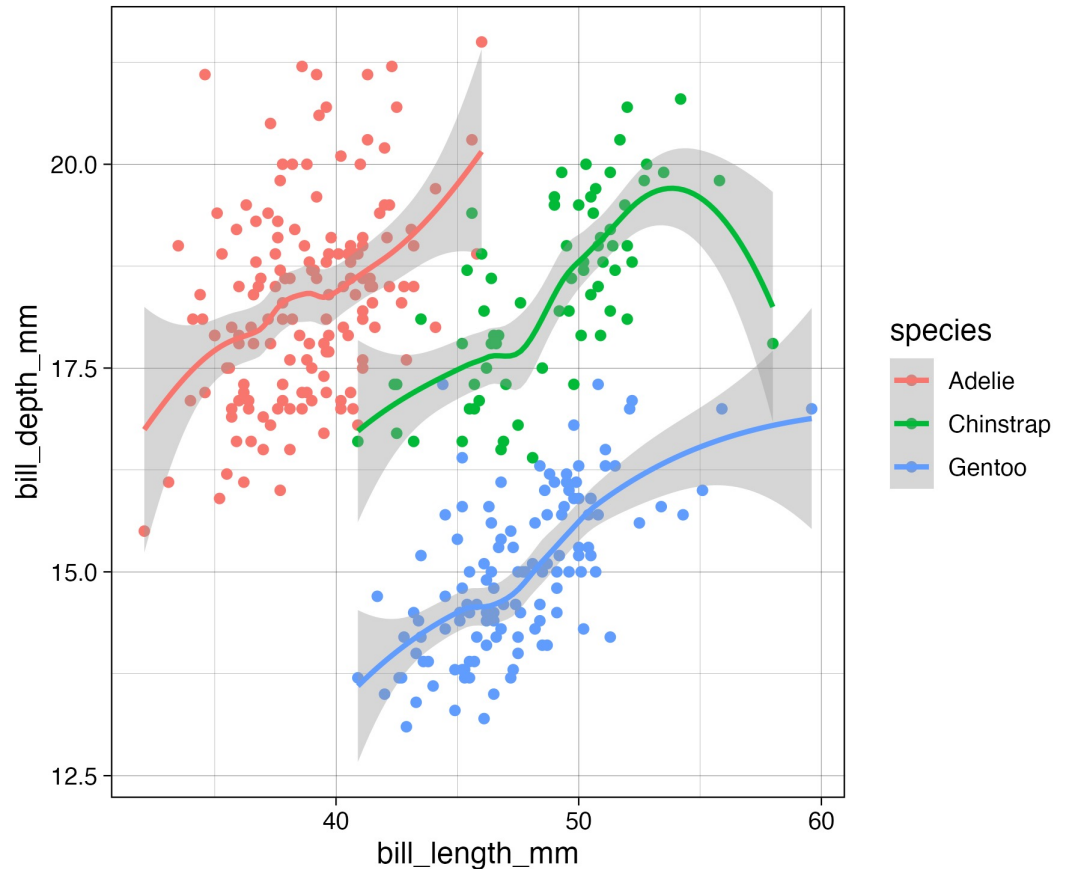


ggplot2: geom_point() + color + geom_smooth

penguins %>%

```
ggplot(aes(x = bill_length_mm,  
           y = bill_depth_mm,  
           color = species)) +
```

```
geom_point() +  
geom_smooth()
```



ggplot2: geom_point() + color + geom_smooth + geom_smooth()

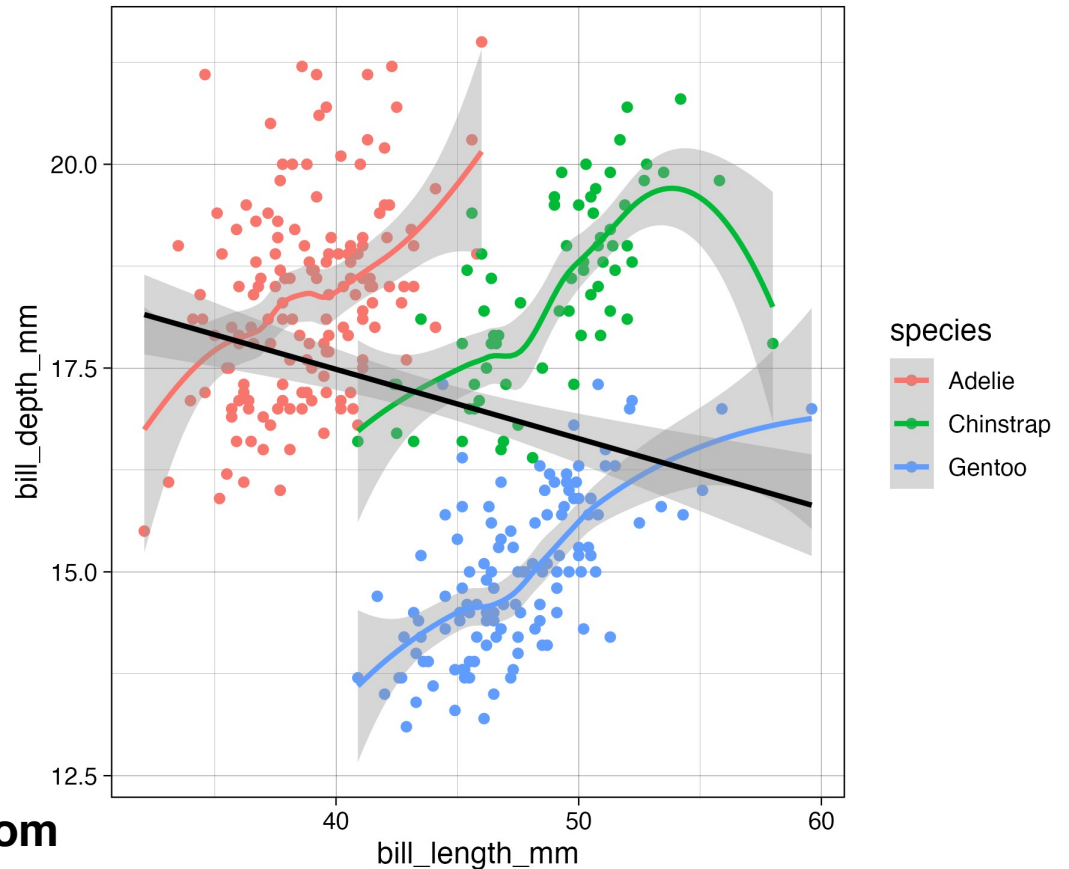
penguins %>%

```
ggplot(aes(x = bill_length_mm,  
           y = bill_depth_mm,  
           color = species)) +
```

```
geom_point() +  
geom_smooth() +  
geom_smooth(method = "lm",  
            color = "black")
```



Overall between-species trend different from
individual species group trends;
i.e. *Simpson's paradox*



ggplot2: Where to put the arguments?

OVERALL aesthetics:

- Place in the `ggplot(aes(...))`

SPECIFIC aesthetics (i.e. only apply to ONE geom)

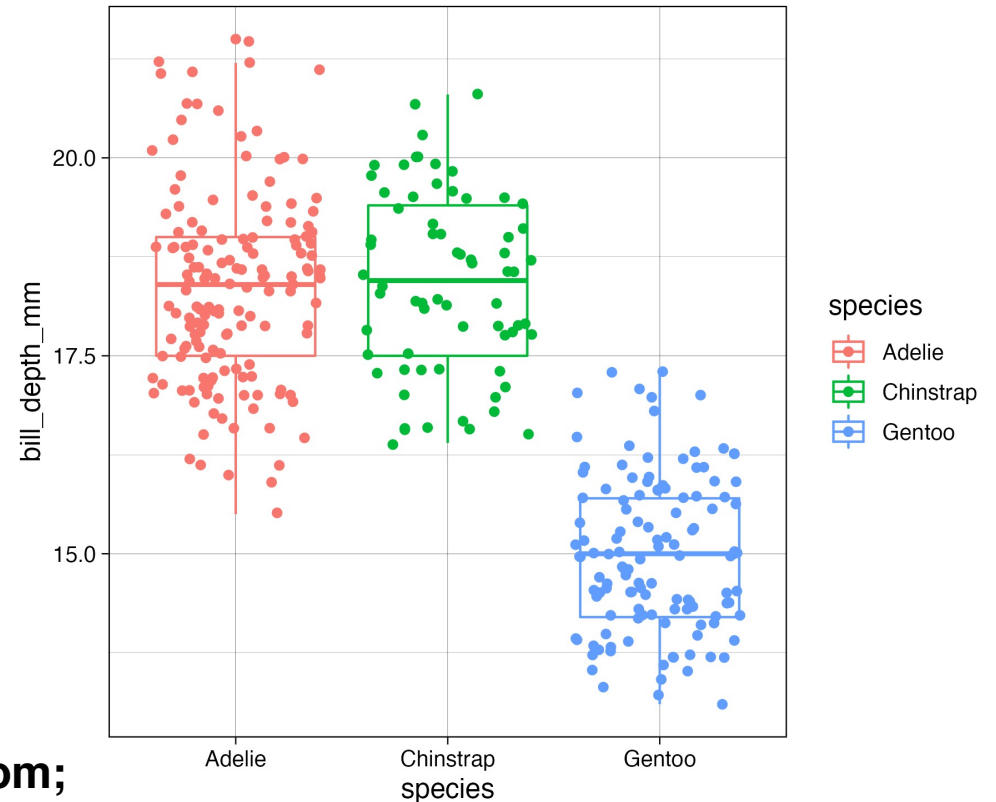
- Place in respective geom

ggplot2: geom_boxplot() + geom_jitter()

penguins %>%

```
ggplot(aes(x = species,  
           y = bill_depth_mm,  
           color = species)) +
```

```
  geom_boxplot() +  
  geom_jitter()
```

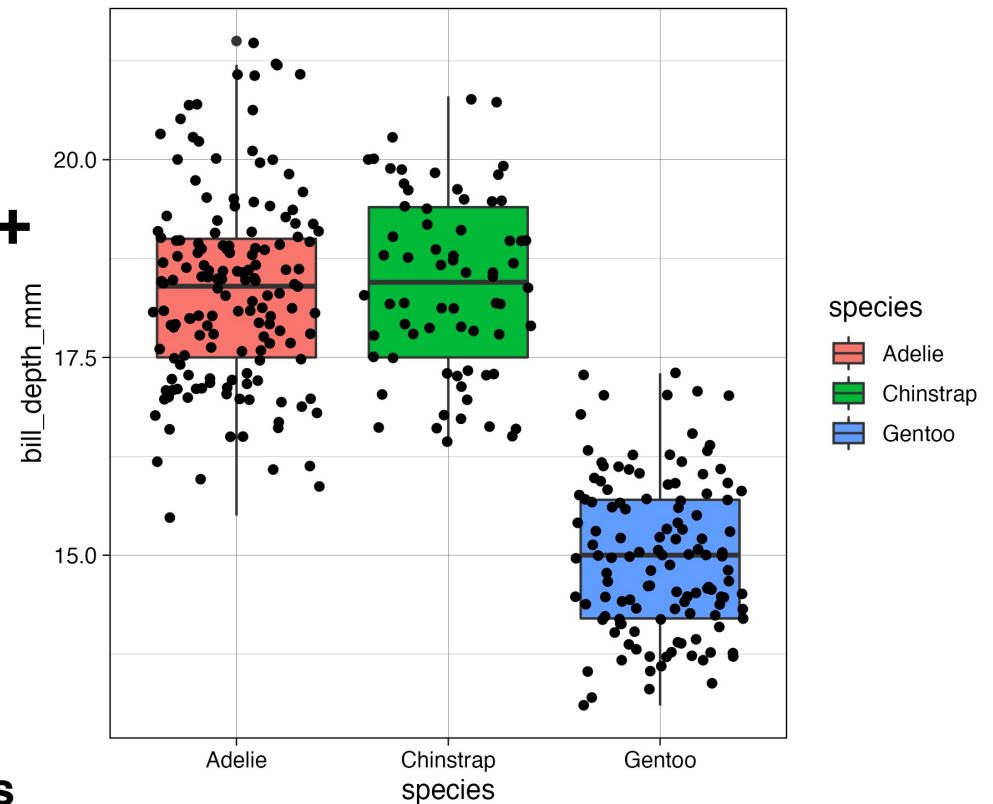


COLOR changes the **OUTER** color of a geom;
FILL changes how e.g. boxplot and violinplots
are filled

ggplot2: geom_boxplot() + color + geom_jitter()

```
penguins %>%
```

```
  ggplot(aes(x = species,  
             y = bill_depth_mm)) +  
    geom_boxplot(aes(fill = species)) +  
    geom_jitter()
```

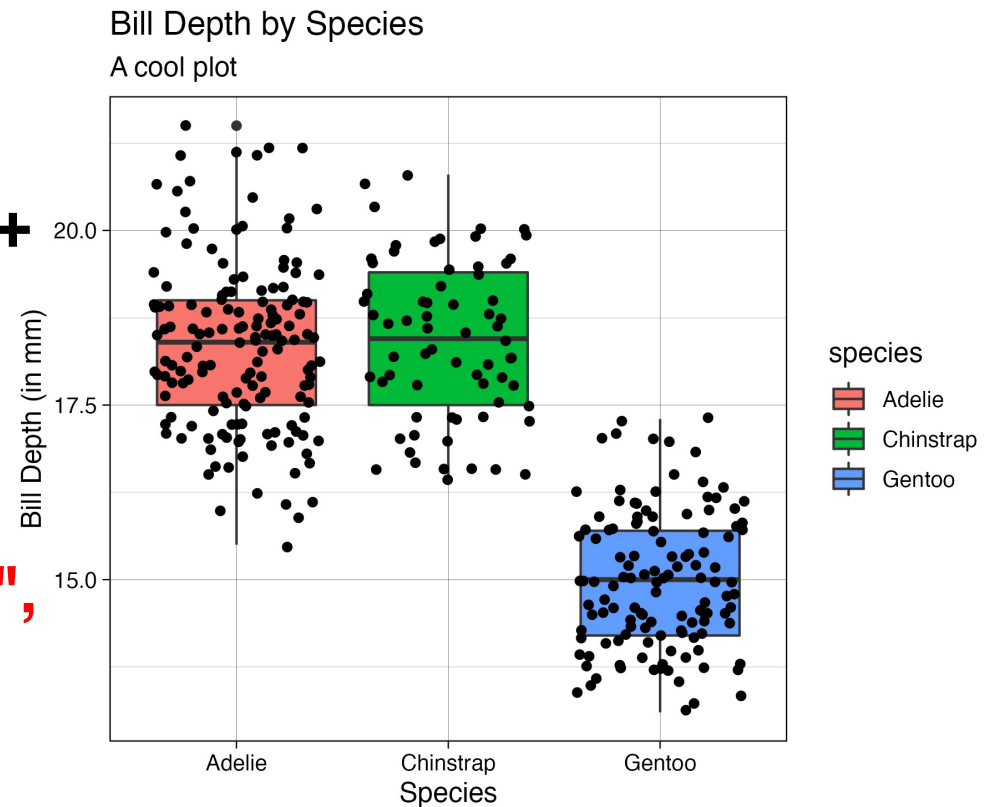


COLOR changes the OUTER color of a geom;
FILL changes how e.g. boxplot and violinplots
are filled

ggplot2: Add plot labels via labs()

penguins %>%

```
ggplot(aes(x = species,  
           y = bill_depth_mm)) +  
  geom_boxplot(aes(fill = species)) +  
  geom_jitter() +  
  labs(x = "Species",  
       y = "Bill Depth (in mm)",  
       title = "Bill Depth by Species",  
       subtitle = "A cool plot")
```



WITH GREAT POWER COMES

A LOT OF CUSTOMIZATION

ggplot2: Saving ggplots

SAVE your plot as an object (just like usual, using the = or -> operator)

e.g.

```
billDepth = penguins %>%  
  ggplot(aes(x = species,  
             y = bill_depth_mm)) +  
  geom_boxplot(aes(fill = species)) +  
  geom_jitter() +  
  labs(x = "Species",  
       y = "Bill Depth (in mm)",  
       title = "Bill Depth by Species",  
       subtitle = "A cool plot")
```

```
ggsave(plot = ...,  
        filename = "RELATIVE PATH")
```

e.g.

```
ggsave(plot = billDepth,  
        filename = "../Figures/billDepth.png")
```



**DON'T FORGET TO SPECIFY
WHAT TYPE OF FILE YOU ARE
SAVING THE PLOT AS (.png, .pdf,
.eps, etc.)**

LET'S GET OUR HANDS DIRTY



makeameme.org