## Penguins\_Plot

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#### **Setting up my environment**

Notes: Setting up my R environment by loading the 'tidyverse', and 'Palmer Panguins' packages.

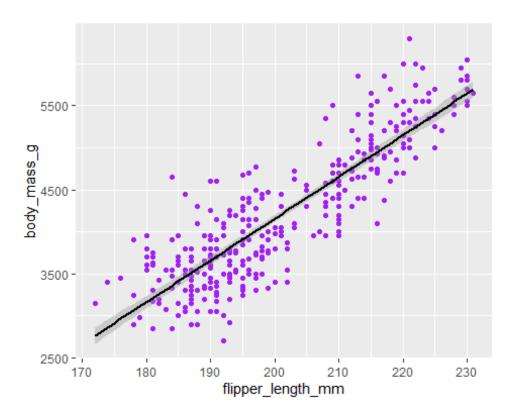
```
library(tidyverse)
library(palmerpenguins)
library(ggplot2)
```

#### View the data

```
glimpse(penguins)
## Rows: 344
## Columns: 8
                      <fct> Adelie, Adelie, Adelie, Adelie, Adelie, Adelie,
## $ species
Adel~
## $ island
                      <fct> Torgersen, Torgersen, Torgersen, Torgersen,
Torgerse~
## $ bill_length_mm
                      <dbl> 39.1, 39.5, 40.3, NA, 36.7, 39.3, 38.9, 39.2,
34.1, ~
## $ bill_depth_mm
                      <dbl> 18.7, 17.4, 18.0, NA, 19.3, 20.6, 17.8, 19.6,
18.1, ~
## $ flipper_length_mm <int> 181, 186, 195, NA, 193, 190, 181, 195, 193, 190,
186~
                      <int> 3750, 3800, 3250, NA, 3450, 3650, 3625, 4675,
## $ body_mass_g
3475, ~
                      <fct> male, female, female, NA, female, male, female,
## $ sex
male~
                      <int> 2007, 2007, 2007, 2007, 2007, 2007, 2007,
## $ year
2007~
```

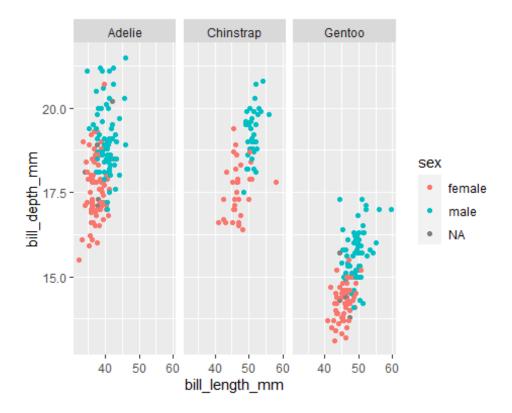
#### Penguins Flipper and body mass comparison in purple

```
penguins %>%
   ggplot(aes(x=flipper_length_mm, y=body_mass_g))+
   geom_point(color="purple") + geom_smooth(method = lm, color = 'Black')
## `geom_smooth()` using formula 'y ~ x'
```



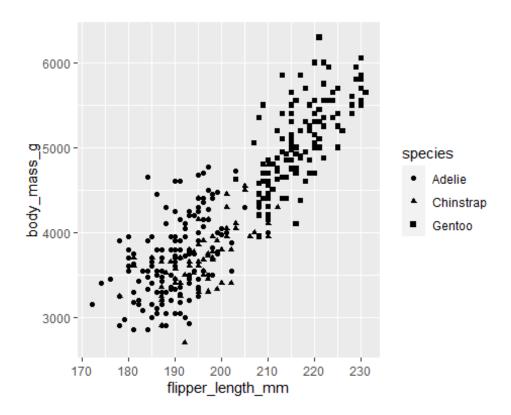
### Bill length and bill depth by species and sex

```
penguins %>%
  ggplot(aes(x=bill_length_mm, y=bill_depth_mm, color=sex))+
  geom_point()+
  facet_wrap(~species)
```



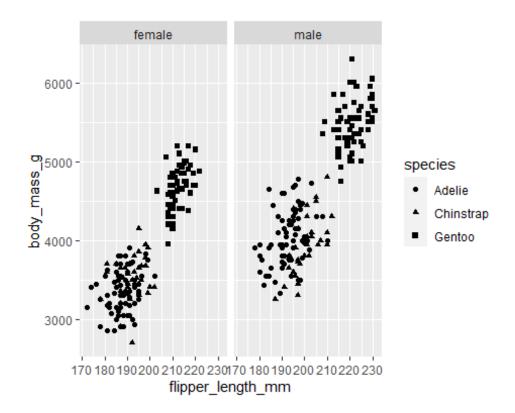
### Fliper and body mass by species

```
penguins %>%
  ggplot(aes(x=flipper_length_mm, y=body_mass_g, shape=species))+
  geom_point()
```



### Flipper and body mass by species and sex

```
penguins %>%
  drop_na(sex) %>%
  ggplot(aes(x=flipper_length_mm, y=body_mass_g, Color=sex, shape=species))+
  geom_point()+
  facet_wrap(~sex)
```



#### **Comparison of Three Species of Penguins**

# Palmer Penguins: Body Mass vs, Flipper Length

Sample of Three Penguins

