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setting up my environment

Notes: setting up my R environment by loading the 'tidyverse' and 'palmer penguins' packages

```
library("tidyverse")

## Warning: package 'tidyverse' was built under R version 4.2.3
## Warning: package 'ggplot2' was built under R version 4.2.3
## Warning: package 'tibble' was built under R version 4.2.3
## Warning: package 'tidyr' was built under R version 4.2.3
## Warning: package 'readr' was built under R version 4.2.3
## Warning: package 'purrr' was built under R version 4.2.3
## Warning: package 'dplyr' was built under R version 4.2.3
## Warning: package 'stringr' was built under R version 4.2.3
## Warning: package 'forcats' was built under R version 4.2.3
## Warning: package 'lubridate' was built under R version 4.2.3

## — Attaching core tidyverse packages — tidyverse
2.0.0 —
## ✓ dplyr      1.1.1      ✓ readr      2.1.4
## ✓ forcats   1.0.0      ✓ stringr   1.5.0
## ✓ ggplot2   3.4.2      ✓ tibble    3.2.1
## ✓ lubridate 1.9.2      ✓ tidyr     1.3.0
## ✓ purrr     1.0.1
## — Conflicts —
tidyverse_conflicts() —
## ✗ dplyr::filter() masks stats::filter()
## ✗ dplyr::lag()     masks stats::lag()
## ⓘ Use the conflicted package (<http://conflicted.r-lib.org/>) to force all
conflicts to become errors

library("palmerpenguins")

## Warning: package 'palmerpenguins' was built under R version 4.2.3
```

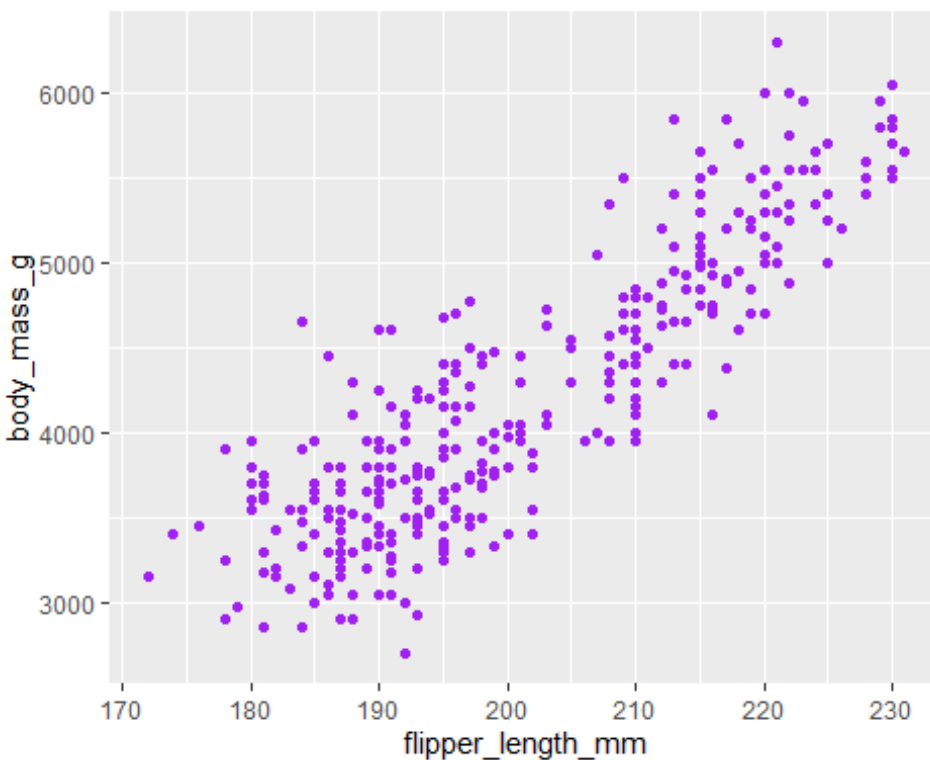
Visualizations

Here we will go through a series of visualizations

Flipper and body mass in purple

Here, we plot flipper length against body mass

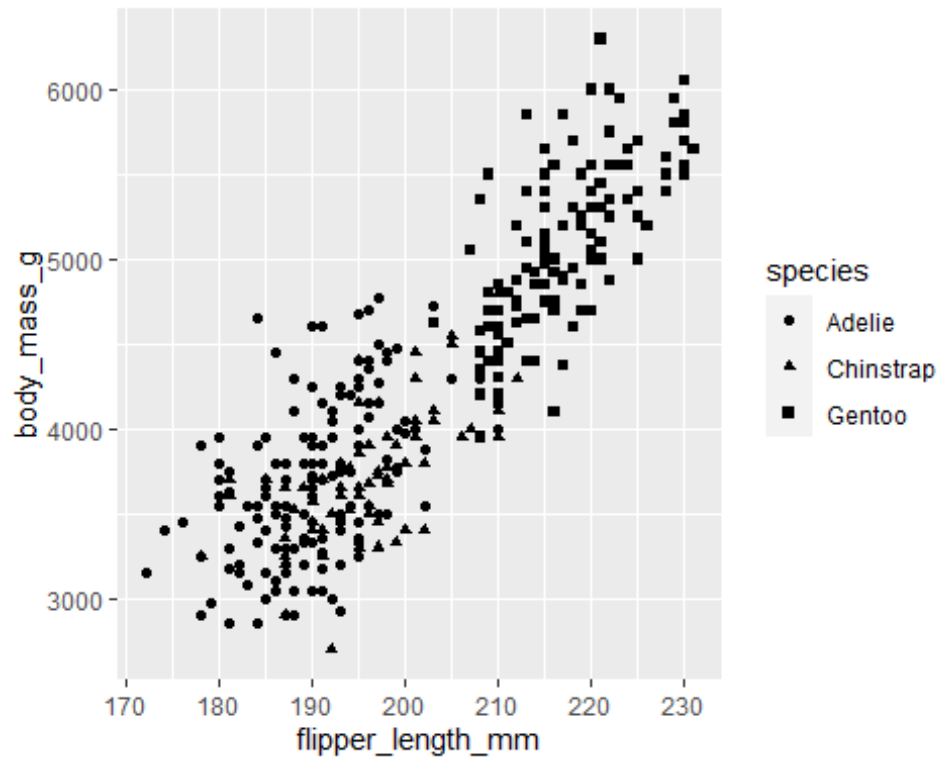
```
ggplot(data=penguins, aes(x=flipper_length_mm, y=body_mass_g)) +  
  geom_point(color="purple")  
## Warning: Removed 2 rows containing missing values (`geom_point()`).
```



Flipper and body mass by species

Here, we plot flipper length against body mass and look at the by species and sex

```
ggplot(data=penguins, aes(x=flipper_length_mm, y=body_mass_g)) +  
  geom_point(aes(shape=species))  
## Warning: Removed 2 rows containing missing values (`geom_point()`).
```



Flipper and body mass by species and sex

Here, we plot flipper length against body mass and look at the breakdown by species and sex

```
ggplot(data = penguins, aes(x = flipper_length_mm, y = body_mass_g)) +  
  geom_point(aes(color = species,  
                 shape = species)) +  
  facet_wrap(~sex)  
## Warning: Removed 2 rows containing missing values (`geom_point()`).
```

