# Data exploration using the palmerpenguins dataset

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# Data exploration

# Motivation

In this section, we **explore** the data from package **palmerpenguins**. A recent publication from the researcher, Dr Kristen Gorman, who shared the data is Connors et al. (2020).

#### Data

The data are displayed below (first 10 rows):

```
penguins %>%
  slice(1:10) %>%
  knitr::kable()
```

species	island	bill_length_mm bill_	_depth_mmflipper_	_lengthmm body_	_mass_	_g sex	year
Adelie	Torgersen	39.1	18.7	181	3750	male	2007
Adelie	Torgersen	39.5	17.4	186	3800	female	2007
Adelie	Torgersen	40.3	18.0	195	3250	female	2007
Adelie	Torgersen	NA	NA	NA	NA	NA	2007
Adelie	Torgersen	36.7	19.3	193	3450	female	2007
Adelie	Torgersen	39.3	20.6	190	3650	$_{\mathrm{male}}$	2007
Adelie	Torgersen	38.9	17.8	181	3625	female	2007
Adelie	Torgersen	39.2	19.6	195	4675	$_{\mathrm{male}}$	2007
Adelie	Torgersen	34.1	18.1	193	3475	NA	2007
Adelie	Torgersen	42.0	20.2	190	4250	NA	2007

#### Numerical exploration

There are 344 penguins in the dataset, and 3 different species. The data were collected in 3 islands of the Palmer archipelago in Antarctica.

The mean of all traits that were measured on the penguins are:

```
## # A tibble: 3 x 6
##
     species
               bill_length_mm bill_depth_mm flipper_length_mm body_mass_g year
     <fct>
                         <dbl>
                                       <dbl>
                                                          <dbl>
                                                                      <dbl> <dbl>
## 1 Adelie
                         38.8
                                        18.3
                                                           190.
                                                                      3701. 2008.
## 2 Chinstrap
                         48.8
                                        18.4
                                                           196.
                                                                      3733. 2008.
## 3 Gentoo
                         47.5
                                        15.0
                                                                      5076. 2008.
                                                           217.
```

# Graphical exploration

A histogram of body mass per species:

## Warning: Removed 2 rows containing non-finite values (stat\_bin).

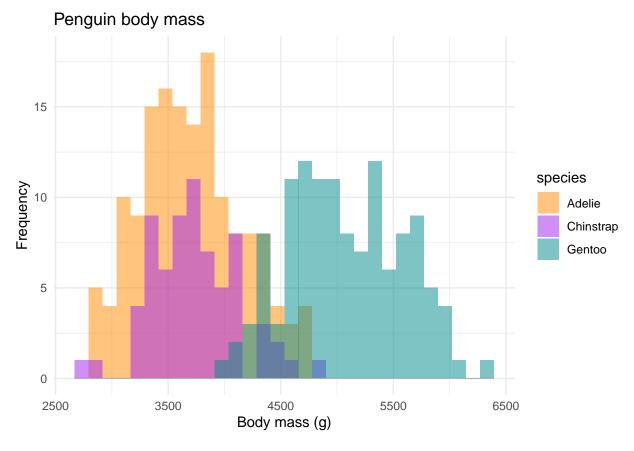


Figure 1: Distribution of body mass by species of penguins

# Linear regression

And here is a nice model with graphical output

```
m1 <- lm(bill_length_mm ~ flipper_length_mm + body_mass_g + sex, data = penguins)
summary(m1)</pre>
```

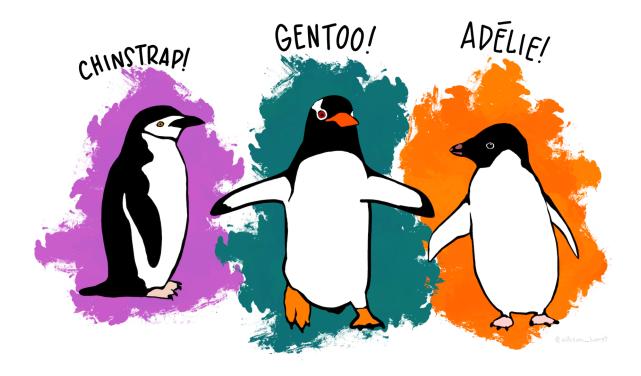
##

```
## Call:
## lm(formula = bill_length_mm ~ flipper_length_mm + body_mass_g +
##
        sex, data = penguins)
##
##
   Residuals:
##
       Min
                  1Q Median
                                    3Q
                                            Max
   -9.6131 -2.8005 -0.6307 2.1699 20.1682
##
## Coefficients:
##
                          Estimate Std. Error t value Pr(>|t|)
## (Intercept)
                       -8.0754117
                                     4.6732129
                                                 -1.728
                                                            0.0849 .
## flipper_length_mm 0.2672650
                                     0.0335203
                                                   7.973 2.57e-14 ***
                                                            0.2853
                       -0.0006670
                                     0.0006232
                                                  -1.070
## body_mass_g
## sexmale
                         2.3047154
                                     0.5055670
                                                   4.559 7.27e-06 ***
##
                     0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
## Signif. codes:
##
## Residual standard error: 4.029 on 329 degrees of freedom
      (11 observations deleted due to missingness)
## Multiple R-squared: 0.4621, Adjusted R-squared: 0.4572
## F-statistic: 94.2 on 3 and 329 DF, p-value: < 2.2e-16
par(mfrow= c(2,2))
plot(m1)
                                                     Standardized residuals
                                                                         Normal Q-Q
                   Residuals vs Fitted
        20
                 0294
                                                                                              2940
   Residuals
                                                           ^{\circ}
        2
        -10
                                                           7
                     40
                               45
                                         50
                                                               -3
                                                                     -2
                                                                                0
                                                                                           2
                                                                                                3
                        Fitted values
                                                                      Theoretical Quantiles
  (Standardized residuals)
                                                     Standardized residuals
                     Scale-Location
                                                                    Residuals vs Leverage
                                                                             2940
        1.5
                                                           \alpha
        0.0
                                         50
                                                              0.00
                     40
                               45
                                                                       0.01
                                                                               0.02
                                                                                       0.03
                                                                                               0.04
                        Fitted values
                                                                             Leverage
```

Figure 2: Checking assumptions of the model

# The end

The 3 species of penguins:



# References

Connors, Brendan, Michael J. Malick, Gregory T. Ruggerone, Pete Rand, Milo Adkison, James R. Irvine, Robert Campbell, and Kristen Gorman. 2020. "Climate and Competition Influence Sockeye Salmon Population Dynamics Across the Northeast Pacific Ocean." Canadian Journal of Fisheries and Aquatic Sciences 77 (6): 943–49. https://doi.org/10.1139/cjfas-2019-0422.