# Summarising and Plotting Data in R

Visualising Data

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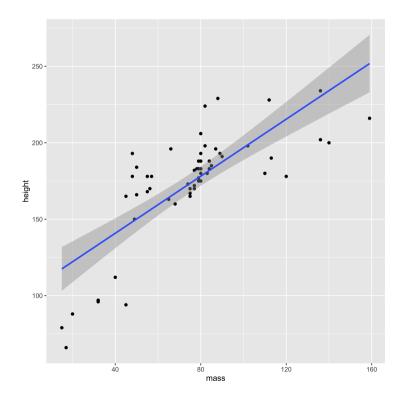
#### **Data Visualisation**

What's a better way to summarise the relationship between height and mass in our star wars data set?

We could report a correlation and **some averages**, though that's going to be hard to picture.

```
## # A tibble: 31 x 3
     species mean_height mean_weight
     <chr>
                    <dbl>
                                 <dbl>
##
## 1 Aleena
                                    15
## 2 Besalisk
                       198
                                   102
## 3 Cerean
                       198
                                    82
## 4 Clawdite
                       168
                                     55
## # ... with 27 more rows
```

So **plots** do a lot of the work.



#### A Grammar of Graphics

R has an inbuilt (base) plotting system, but we'll use **ggplot2** from the tidyverse. ggplot2 has a consistent method of building up plots layer by layer.

#### We have to:

- Define how we want to **map** the data onto the plot.
- Define the **geom**etric shapes we want to use to draw the data.
- Define any **aes**thetics for the look for our plot.

# Grammar of Graphics with ggplot2

As with all R functions, we need to pass arguments to ggplot2. Let's start with our data:

```
ggplot(data = starwars)
```

That's not too exciting, but we can see we've made the beginning of a plot (the background).

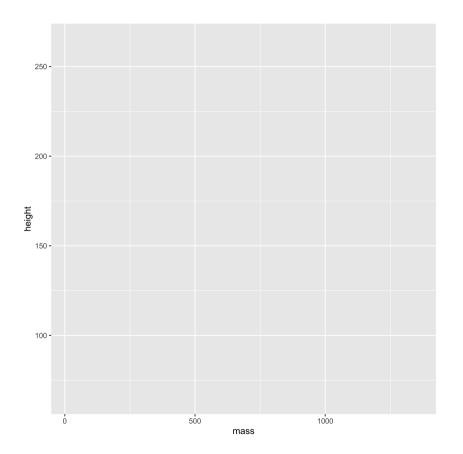
## Mapping our Data

Then we can define how we want to map data onto aesthetics.

Here, we tell R to map the mass and height data from starwars onto the X and Y axes respectively.

```
ggplot(
  data = starwars,
  mapping = aes(x = mass, y = height)
)
```

That looks a little better. Now we know the ranges of our data!



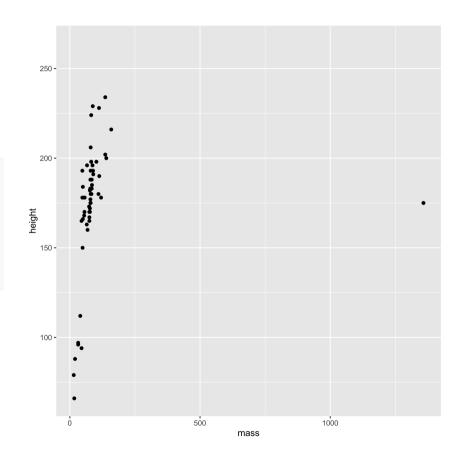
# Adding Geoms

Next, we can tell R how we want to present the data on screen in a new **layer**.

We do this by defining the **geometric shapes** of the data, or geoms.

```
ggplot(
  data = starwars,
  mapping = aes(x = mass, y = height)
) +
  geom_point()
```

Now we can see how mass and height relate to one another.

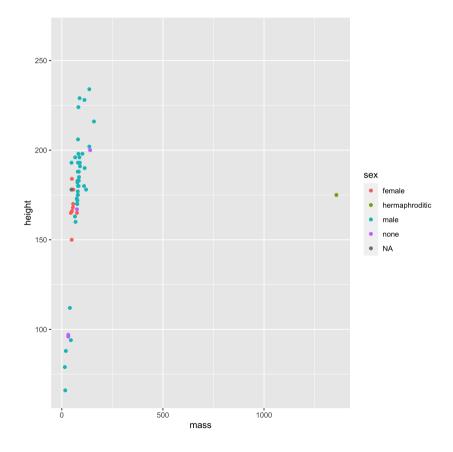


## Differentiating Data

We can add additional arguments to our mappings, for example **mapping colour onto sex**.

```
ggplot(
  data = starwars,
  mapping = aes(x = mass, y = height, colour = sex)
) +
  geom_point()
```

As you can see, we get different groups of data depending upon the sex of the individual.

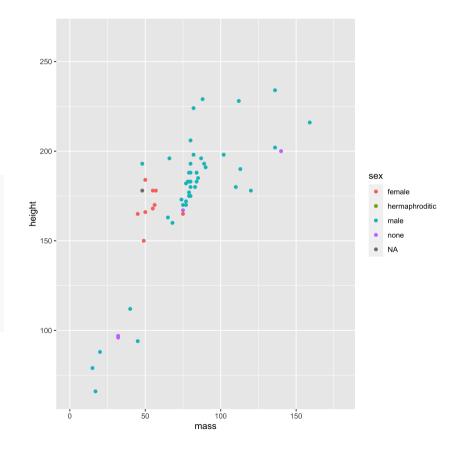


# Filtering and Zooming

It's difficult to see the pattern between height and mass due to an extreme outlier.

We can exclude this individual either by (a) Filtering observations first, (b) restricting limits on the axis of the plot using, e.g. xlim() in coord\_cartesian().

```
ggplot(
  data = starwars,
  mapping = aes(x = mass, y = height, colour = sex)
) +
  geom_point() +
  coord_cartesian(xlim = c(0, 180))
```



# Filtering and Zooming

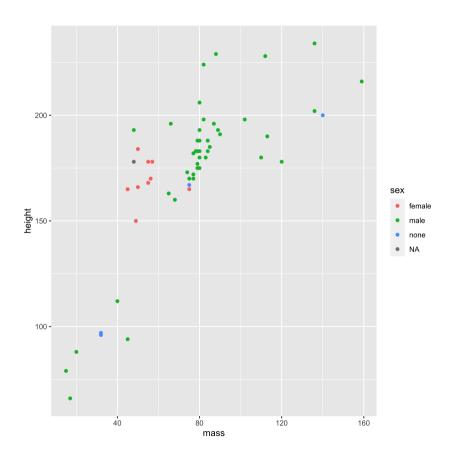
We can exclude this individual either by:

• Chaining tidyverse functions together. Here, we'll filter Jabba from the data before plotting.

```
starwars %>%
  filter(mass < 180) %>%
  ggplot(
    mapping = aes(x = mass, y = height, colour = sex)
) +
  geom_point()
```

We pipe the data into the filter function which passes the filtered data to ggplot.

Warning: we pipe data using %>% but build plot layers with +. Don't confuse the two.



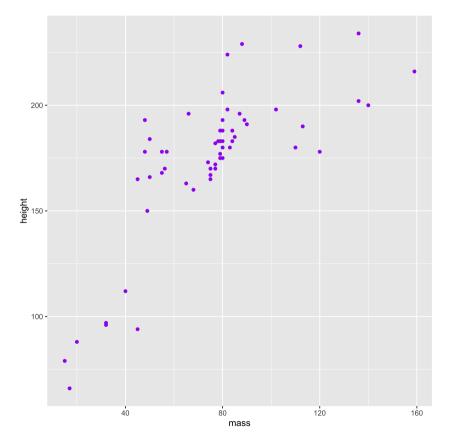
## **Setting Fixed Colours**

What if you want all points to be the same colour? Don't map it to a group. Set it outside the mapping.

```
starwars %>%
  filter(mass < 180) %>%
  ggplot(mapping = aes(x = mass, y = height)) +
  geom_point(colour = "purple")
```

Here, colour is defined in the geom\_point() call. Colour only applies to this geometric shape.

This overrides any specific group mappings.

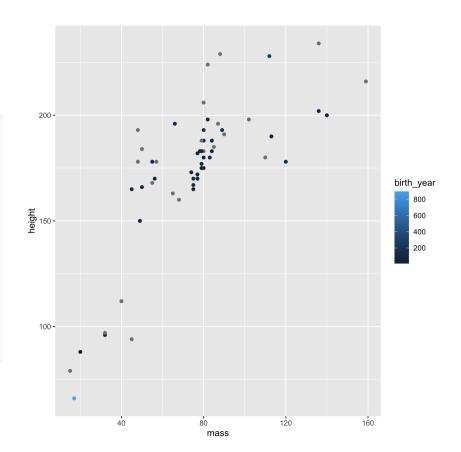


#### **Continuous Colour**

ggplot is pretty smart in how it handles colour. Categorical variables get indiviudal colours, while scales get continuous palettes.

```
starwars %>%
  filter(mass < 180) %>%
  ggplot(
    mapping = aes(
        x = mass,
        y = height,
        colour = birth_year
    )
) +
geom_point()
```

Now the legend doesn't have categories, but a scale.

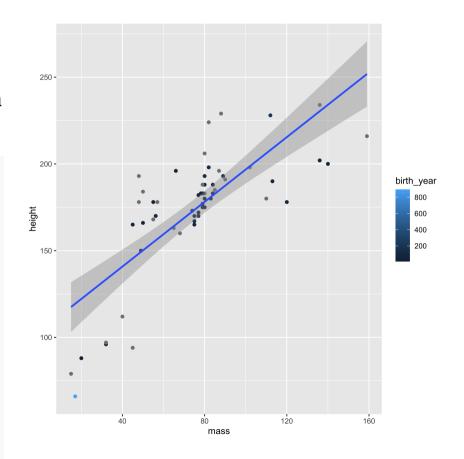


## Combining Geoms

We can add geoms like a line of best fit with 95% confidence intervals!

Without getting too into it, this is defined as a **linear model** in a **smoothing function**, hence we use geom\_smooth().

```
starwars %>%
 filter(mass < 180) %>%
 ggplot(
   mapping = aes(
     x = mass,
      y = height,
      colour = birth_year
 geom_point() +
 geom_smooth(
   method = "lm",
   formula = "y ~ x"
```



#### Some Other Geoms

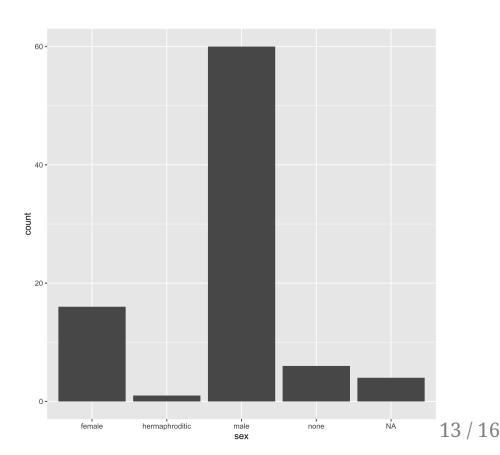
#### Bar Plots for Count Data

For a **bar plot of count data**, we just need to pass a column of data to the aesthetics, and say it should appear on the x-axis.

We then use geom\_bar() to make the bar plot.

This is what this looks like for counting up characters of different sexes.

```
ggplot(data = starwars, mapping = aes(x = sex)) +
  geom_bar()
```



#### Some Other Geoms

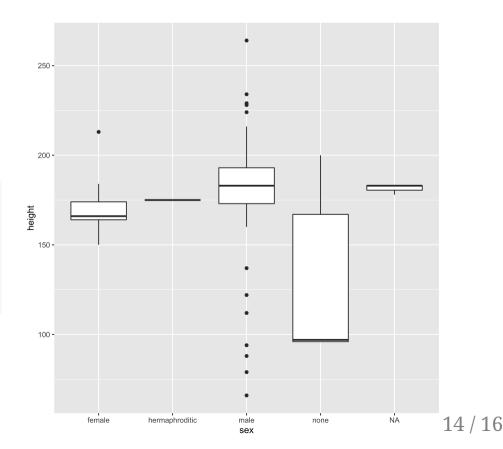
#### **Box Plots**

What if we want to display a **continuous variable** across groups? A **boxplot** is handy here.

Let's get heights of characters from each sex. Now, we just add height to the y-axis, and change geom\_bar() to geom\_boxplot()

```
ggplot(
  data = starwars,
  mapping = aes(x = sex,y = height)
) +
  geom_boxplot()
```

Remember, the dark line is the median, the box the middle 50% of scores.



#### Saving Plots

We can save graphics from R using a number of methods, but for plots produced in ggplot2, we can use ggsave().

We can either make our plot without assigning it to a variable, and then save it as follows:

```
ggplot(data = starwars, aes(x = height)) +
  geom_density()

ggsave(here("myplot.png"), last_plot())
```

Or we can make a plot and assign it to a variable, and save it as follows:

```
my_plot <- ggplot(data = starwars, aes(x = height)) +
  geom_density()

ggsave(here("myplot.png"), my_plot)</pre>
```

I prefer the latter. Why? Once a plot is stored as a variable, you can change it by adding ggplot arguments! For example...

```
my_plot + coord_cartesian(xlim = c(0, 180))
```

#### Recap

#### We've learned...

- About the importance of graphs for communicating findings.
- About a **grammar of graphics** for defining plots in code.
- How to **map** data onto a plot, how to display that data with **geom**etric shapes, and how to control the **aes**thetics of the plot.
- How to build up a plot layer by layer in ggplot2.
- How to use ggplot functions to **differentiate data**.
- How to use ggplot functions to filter data, or how to **chain functions together** from other packages.
- How to **use different geoms** to summarise your data in different ways.
- How to save your plots.