

Using grattantheme

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This vignette explains how to use `grattantheme` to quickly and consistently apply Grattan chart formatting to charts made in R using `ggplot`.

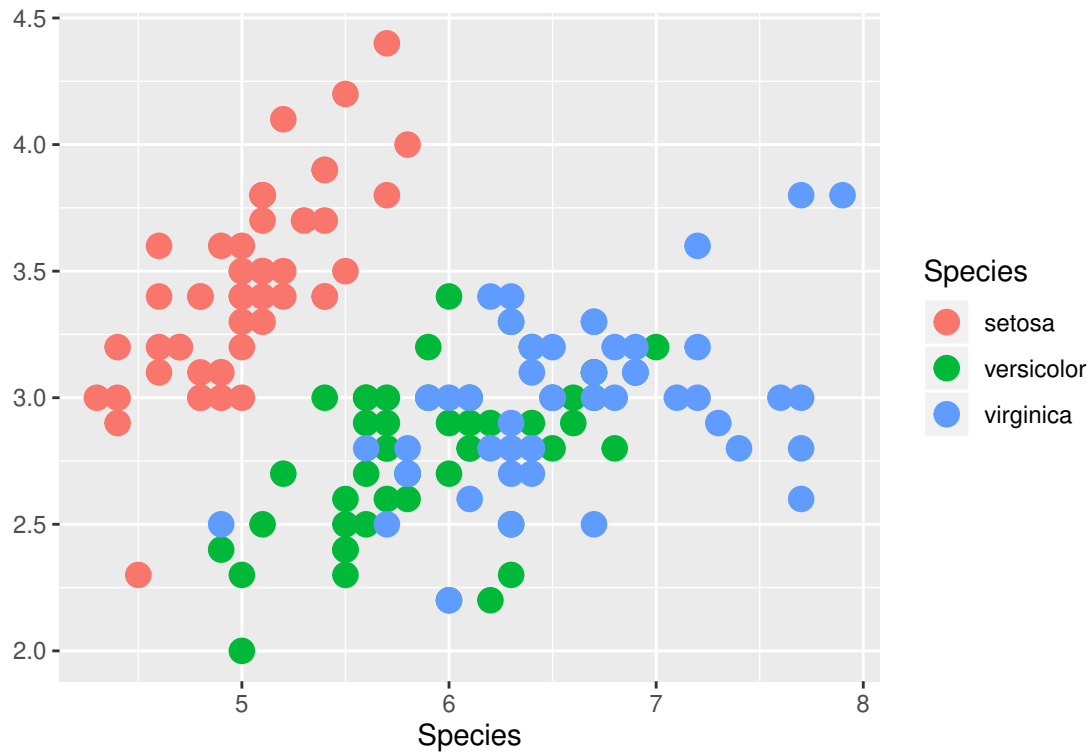
When creating a chart using `ggplot` we have to:

- Choose a dataset;
- Map variables to chart aesthetics `aes()`;
- Choose a `geom_`.

For example, using the in-built `iris` dataset:

```
plot <- ggplot(iris,
               aes(x = Sepal.Length,
                   y = Sepal.Width,
                   colour = Species)) +
  geom_point(size = 4) +
  labs(x = "Species",
       y = "",
       colour = "Species")
```

This successfully plots the data we want to plot:

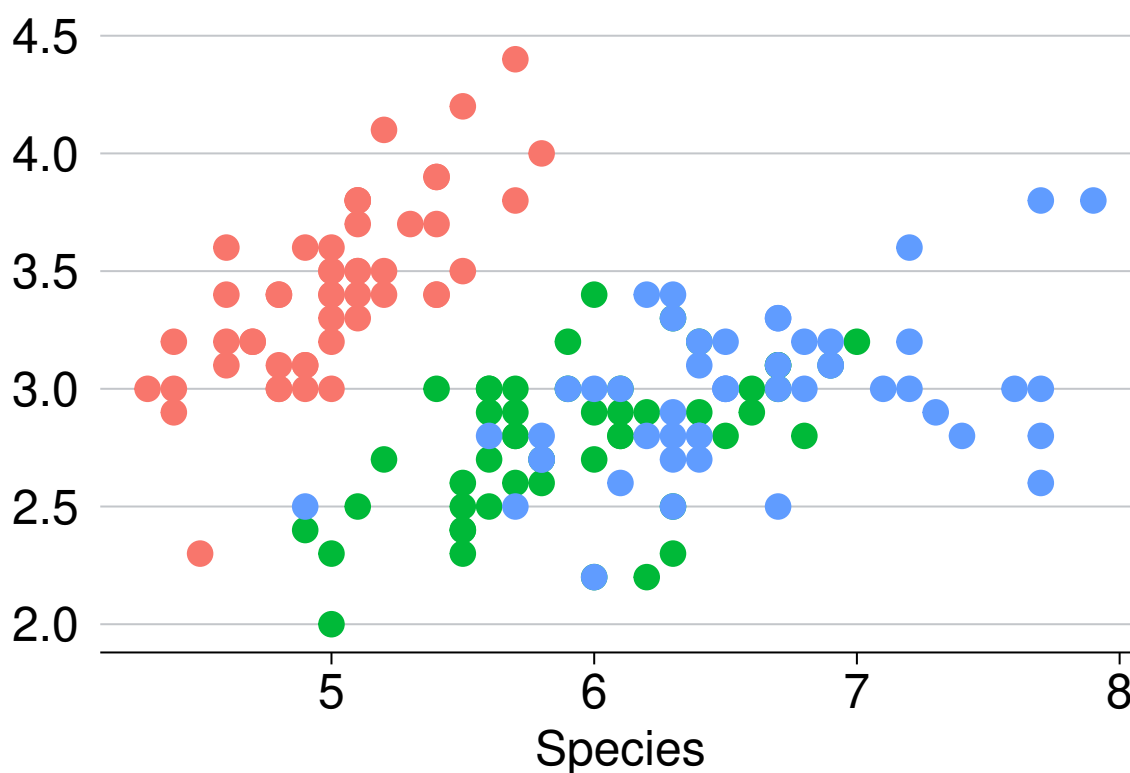


But it doesn't yet *look* like a Grattan chart. To adjust the *look* we adjust 'theme' elements, like `axis.ticks.x = element_line(colour = "black")` to adjust the axis tickmarks on the x axis; `panel.grid.major.x = element_blank()` to turn off vertical gridlines; and so on; and on; and on. We also need to adjust aesthetic colours to the Grattan palette; setting, for example, `fill = "#F68B33"`. The `grattantheme` package contains tools and shortcuts to simplify this process.

Formatting theme elements with `theme_grattan()`

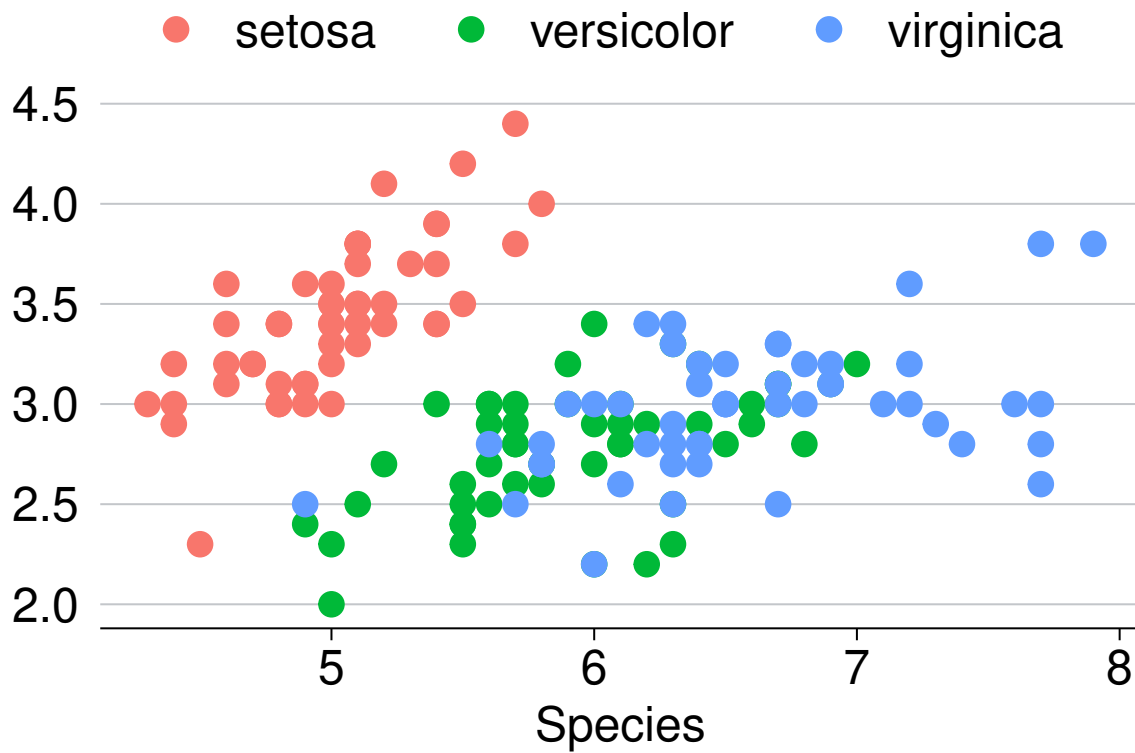
The function `theme_grattan()` contains all of the Grattan `theme` adjustments in one handy command. Combined with `grattan_colour_manual`, which easily changes colours of aesthetics, your R chart will be ready for a report or a slide in no time.

```
plot +  
  theme_grattan()
```



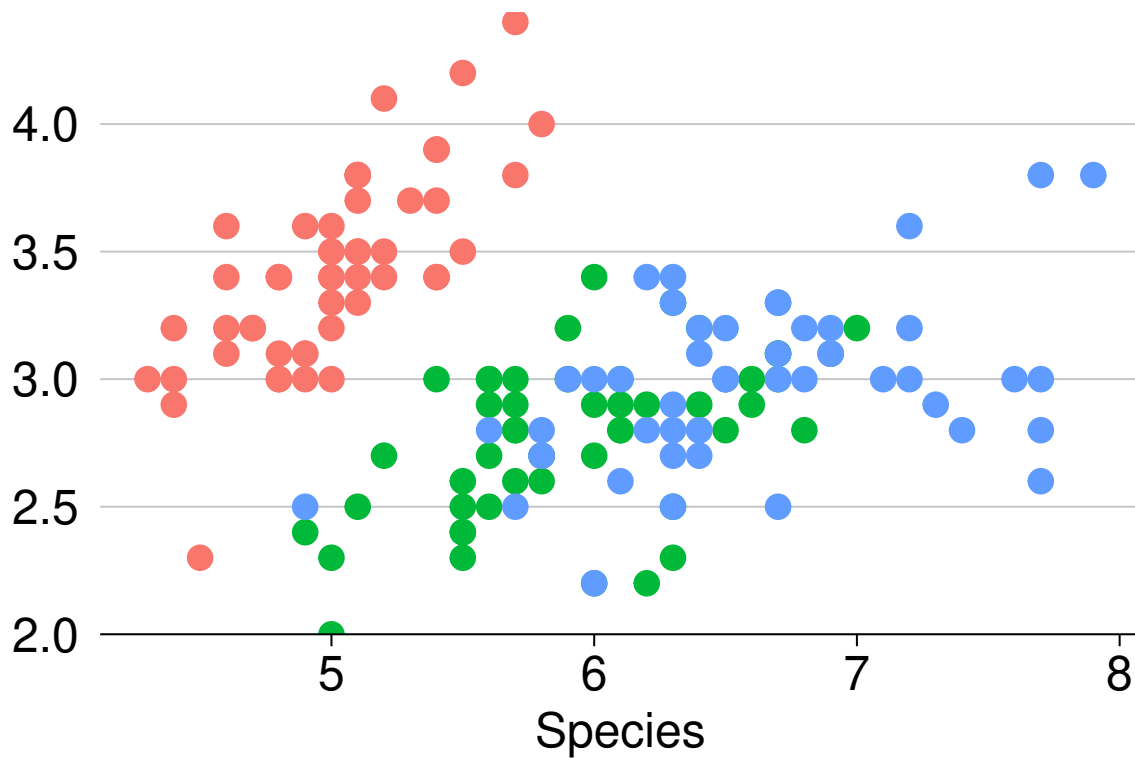
By default, `theme_grattan()` suppresses the legend to allow for clearer on-chart labelling. We can include the legend with the `legend` argument, which takes "off", "top", "bottom", "left" or "right":

```
plot +  
  theme_grattan(legend = "top")
```



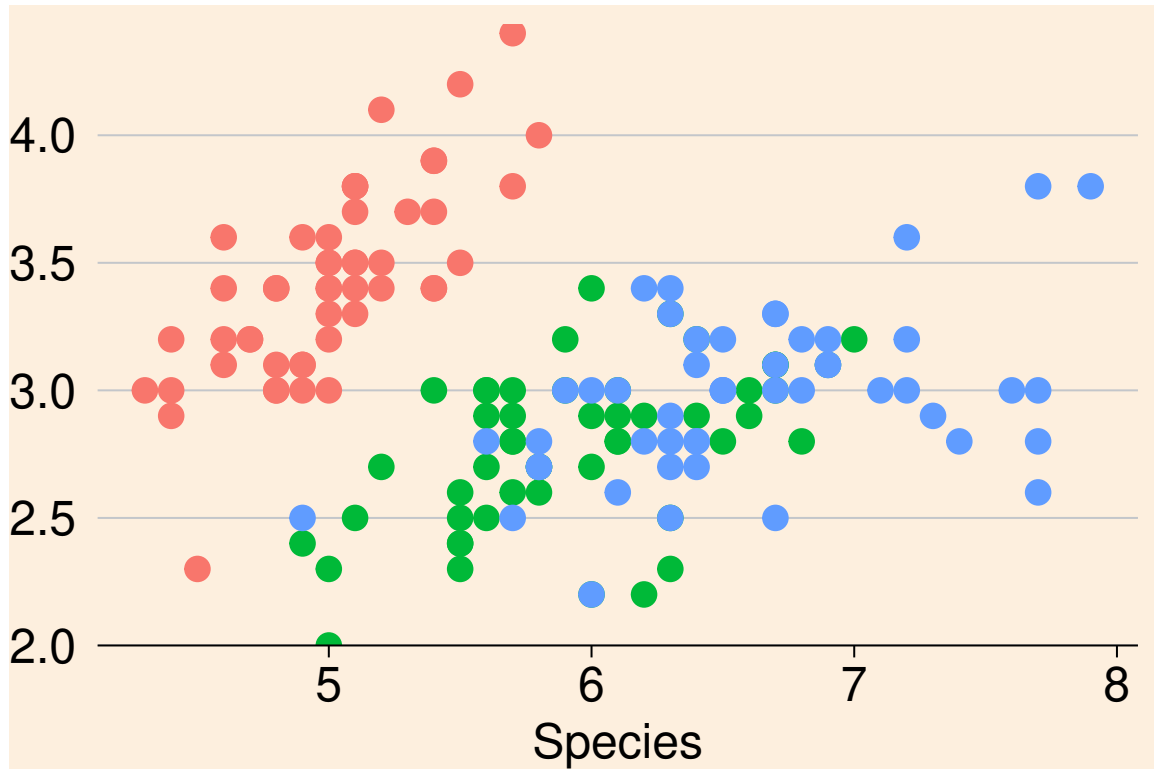
To align the y-axis with zero, change the y scale with `grattan_y_continuous()`:

```
plot +  
  theme_grattan() +  
  grattan_y_continuous()
```



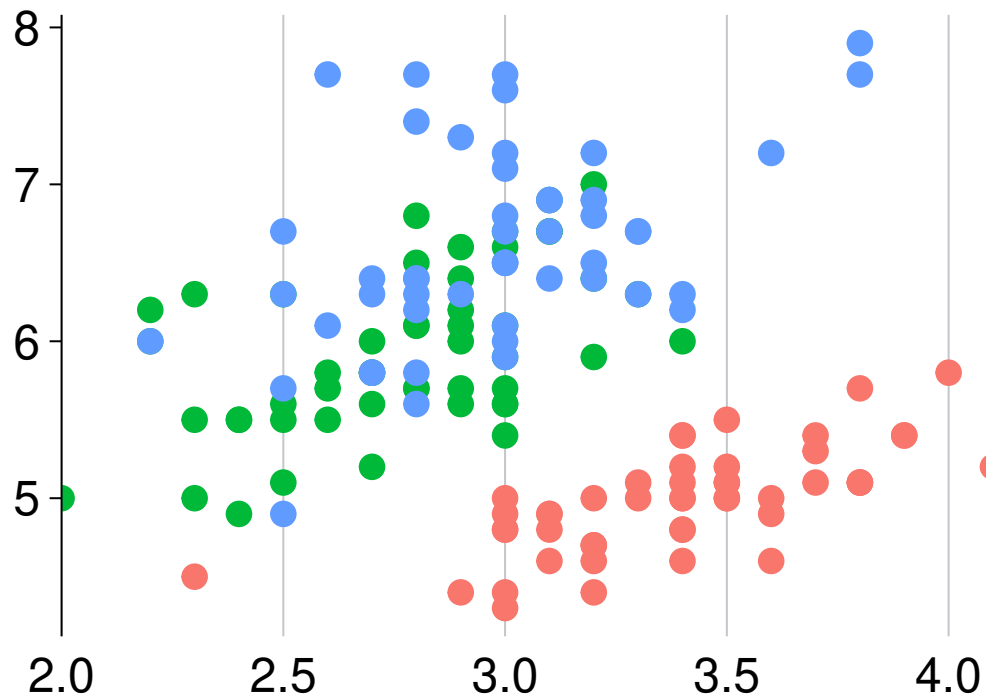
Sometimes we'll want a chart for a box in a report. We can change the background colour with the `background` argument:

```
plot +  
  theme_grattan(background = "box") +  
  grattan_y_continuous()
```



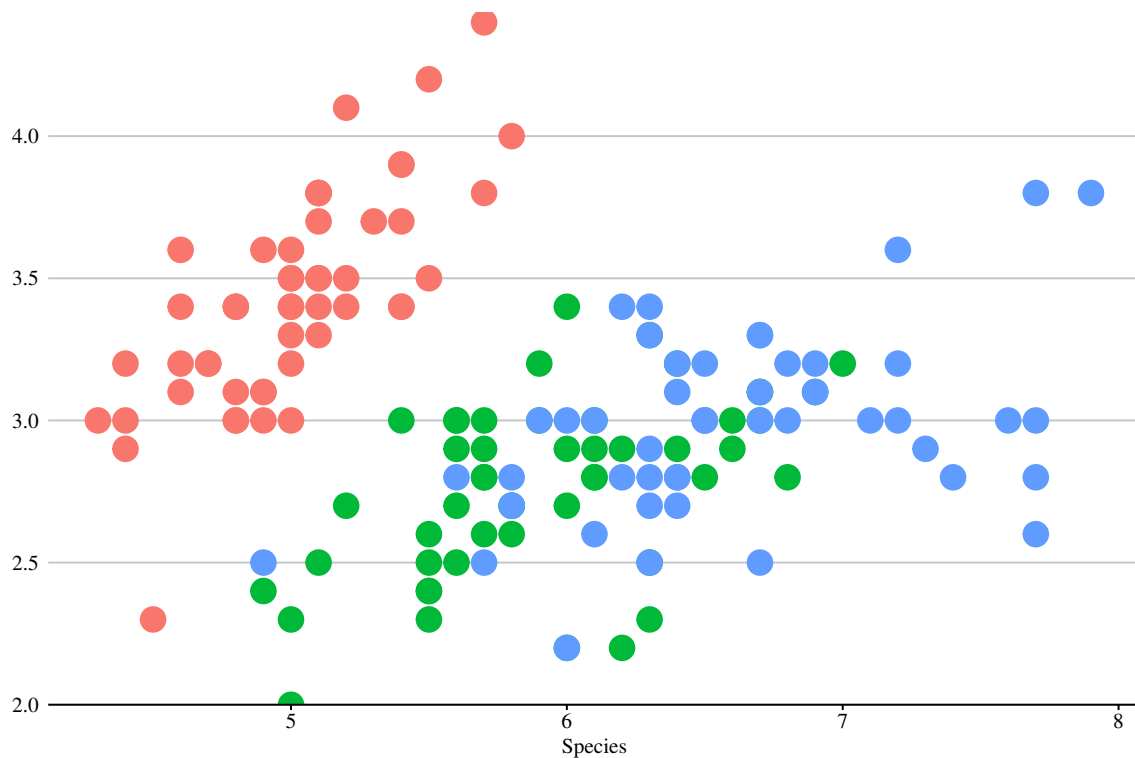
The standard Grattan rules for `x` and `y` axes flip if the chart is a horizontal bar chart. The `x` axis then follows the rules of the `y` axis, and vice-versa. If we are using a 'flipped' chart (implemented with `coord_flipped()`), we can tell `theme_grattan` this is the case using the argument `flipped` set to `TRUE`.

```
plot +  
  coord_flip() +  
  theme_grattan(flipped = TRUE) +  
  grattan_y_continuous()
```



The final adjustments we can specify with `theme_grattan` are the font size and font family. The defaults meet Grattan formatting requirements, but if we do need to change them we can:

```
plot +  
  theme_grattan(base_size = 8, base_family = "serif") +  
  grattan_y_continuous()
```



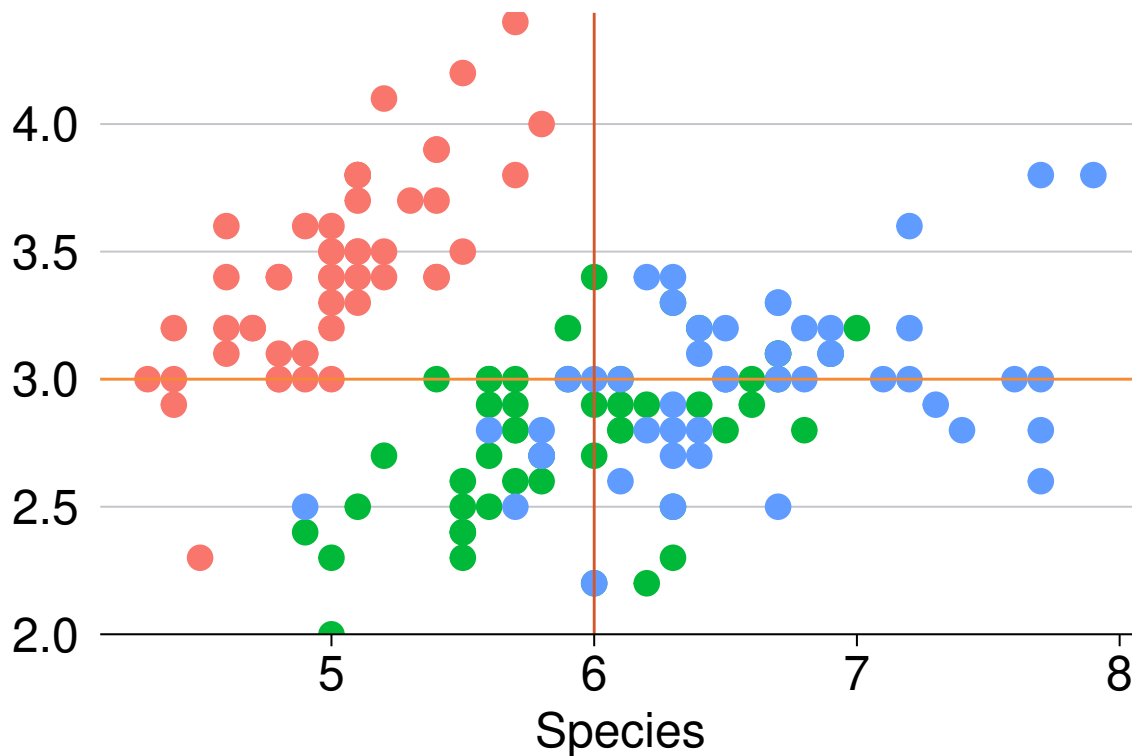
Using Grattan colours

Grattan's colours are loaded with `grattantheme`. The HEX codes for individual Grattan colours can be called using `grattan_[colourname]`, eg `grattan_lightorange`. Colours names are taken from the chart-guide and are:

	<code>grattan_lightyellow</code>		<code>grattan_red</code>
	<code>grattan_yellow</code>		<code>grattan_darkred</code>
	<code>grattan_lightorange</code>		<code>grattan_grey1</code>
	<code>grattan_darkorange</code>		<code>grattan_grey5</code>

We can call single colours:

```
plot +  
  geom_hline(yintercept = 3, colour = grattan_lightorange) +  
  geom_vline(xintercept = 6, colour = grattan_darkorange) +  
  theme_grattan() +  
  grattan_y_continuous()
```

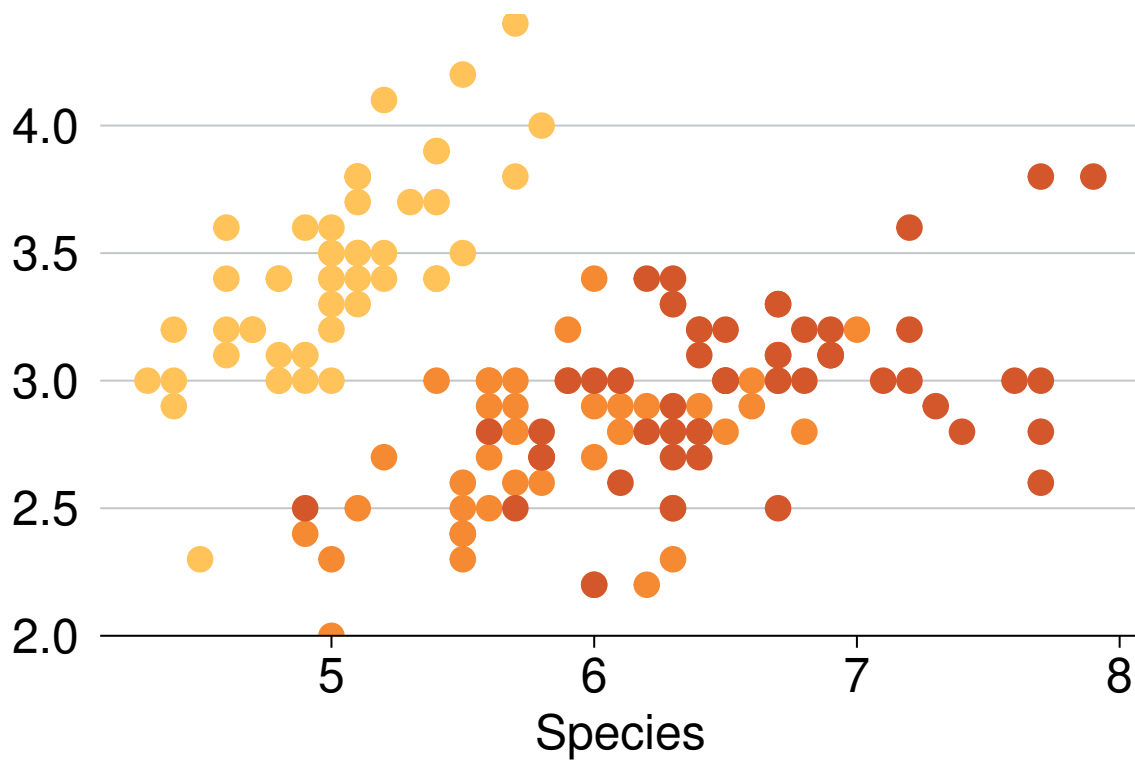


We can also use the `grattan_fill_manual()` or `grattan_colour_manual()` functions to change the colours of our fill or colour *aesthetics*. These can be used for discrete/catagorical data (the default) or continuous data.

Discrete colours

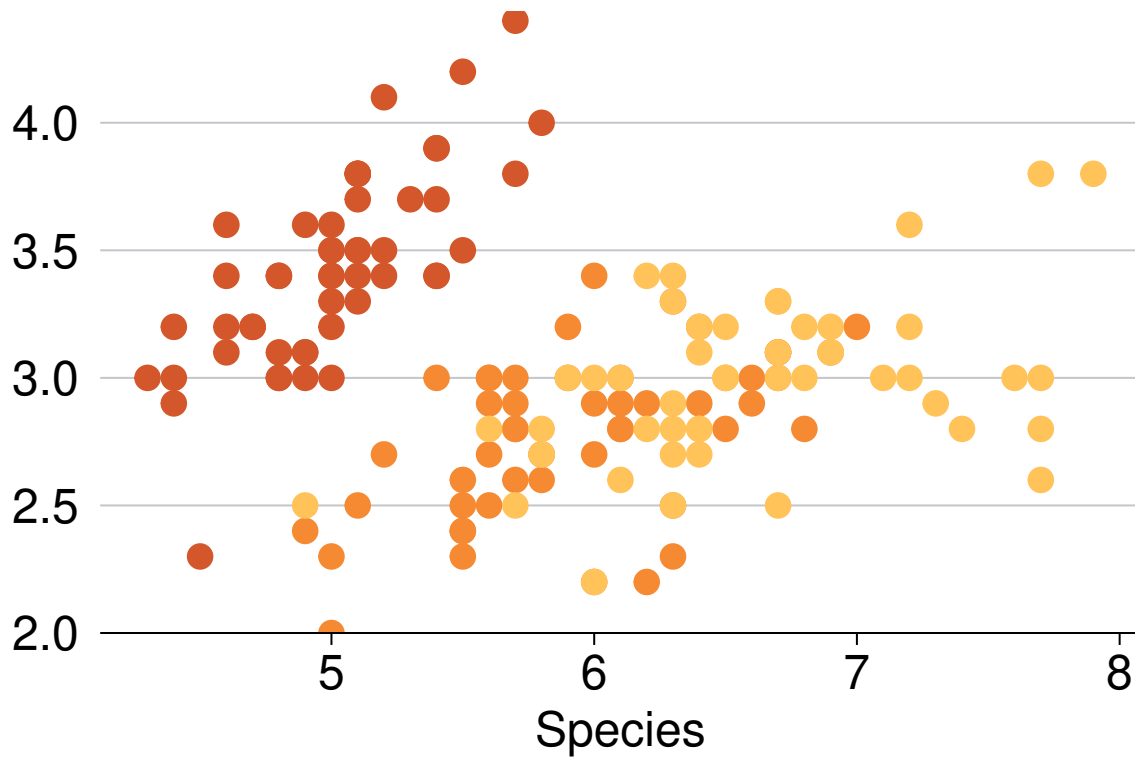
In our example, we have five different trees each represented by a colour, so we set the number of colours to five: `grattan_fill_manual(n = 5)`:

```
plot +  
  theme_grattan() +  
  grattan_y_continuous() +  
  grattan_colour_manual(3)
```



We can reverse the order of the fill colours using the `reverse` argument:

```
plot +  
  theme_grattan() +  
  grattan_y_continuous() +  
  grattan_colour_manual(3, reverse = TRUE)
```

Note that if you do not specify *enough* colours, will receive an error:

```
plot +
  theme_grattan() +
  grattan_y_continuous() +
  grattan_colour_manual(2)
#> Error: Insufficient values in manual scale. 3 needed but only 2 provided.
```

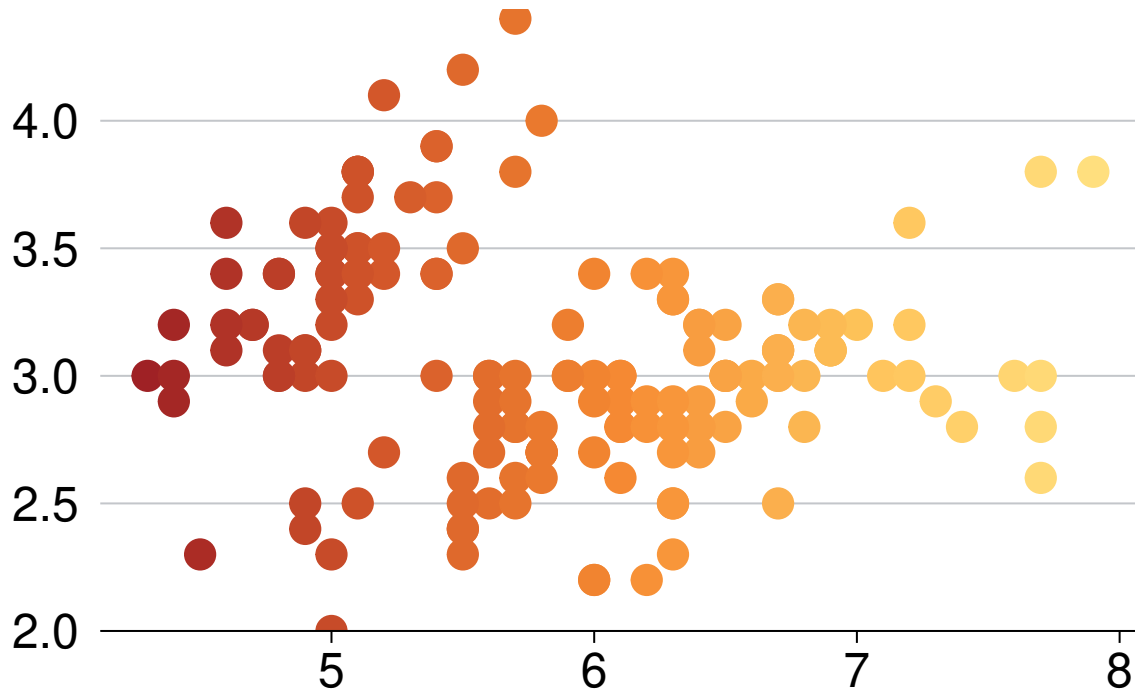
You won't receive an error if you specify *too many* colours. `ggplot` will choose the first `n` colours it needs and ignore the rest.

Continuous colours

`grattan_(colour|fill)_manual` includes an option for continuous colours: `discrete = FALSE`.

```
plot2 <- ggplot(iris,
  aes(x = Sepal.Length,
      y = Sepal.Width,
      colour = Sepal.Length)) +
  geom_point(size = 5) +
  grattan_y_continuous() +
  labs(x = "")

plot2 +
  theme_grattan() +
  grattan_colour_manual(discrete = FALSE)
```



We can change the set of colours the continuous palette is generated from using the `palette` argument:

```
p_full <- plot2 +
  theme_grattan() +
  grattan_colour_manual(discrete = FALSE, palette = "full")

p_full_f <- plot2 +
  theme_grattan() +
  grattan_colour_manual(discrete = FALSE, palette = "full_f")

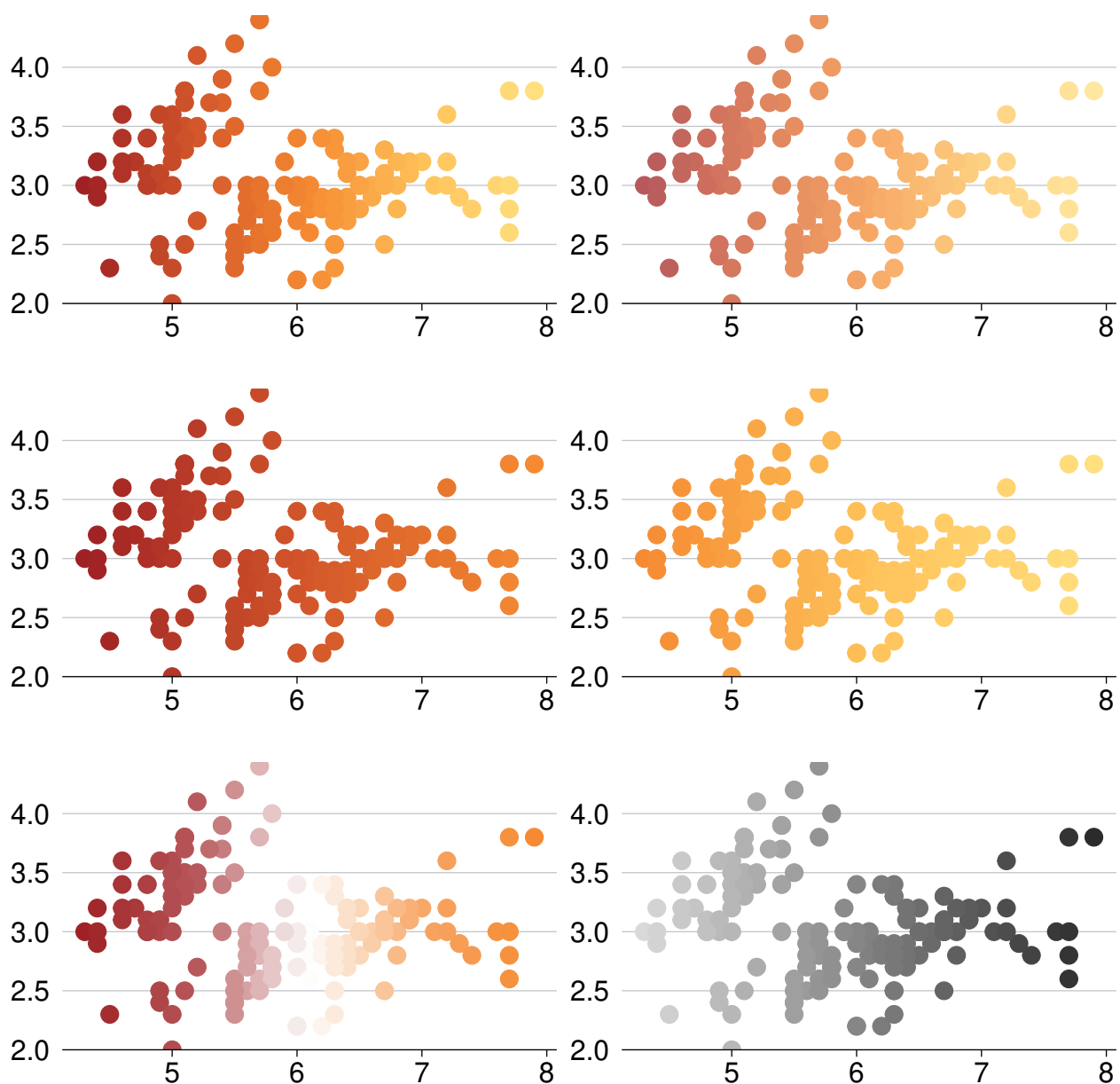
p_dark <- plot2 +
  theme_grattan() +
  grattan_colour_manual(discrete = FALSE, palette = "dark")

p_light <- plot2 +
  theme_grattan() +
  grattan_colour_manual(discrete = FALSE, palette = "light")

p_diverging <- plot2 +
  theme_grattan() +
  grattan_colour_manual(discrete = FALSE, palette = "diverging")

p_grey <- plot2 +
  theme_grattan() +
  grattan_colour_manual(discrete = FALSE, palette = "grey")

gridExtra::grid.arrange(p_full, p_full_f,
  p_dark, p_light,
  p_diverging, p_grey)
```



Saving plots with `grattan_save`

The `grattan_save()` function implements `ggsave()` with Grattan defaults for different occasions. We specify the occasion with the `type` argument that can take the arguments:

- "normal": a standard report chart size, and the default. [`height` = 14.5cm x `width` = 22.16cm]
- "wholecolumn": a taller whole-column chart for reports. [22.16 x 22.16]
- "fullpage": a full-page chart for reports. [22.16 x 44.32]
- "tiny": a rarely-used short chart for reports. [11.08 x 22.16]
- "fullslide" : to produce charts on a full slide with the Grattan logo—to be used in articles or blogposts—use `"type = fullslide"`. This option allows notes and sources to accompany the saved figure.

The argument `filename` is required. You must include an extension; `.pdf` is standard for Grattan charts for reports; `.png` is standard for the Grattan Blog.

Like `ggsave`, the `ggplot2` object to be saved will default to the last plot, but you can specify something else with `object`.

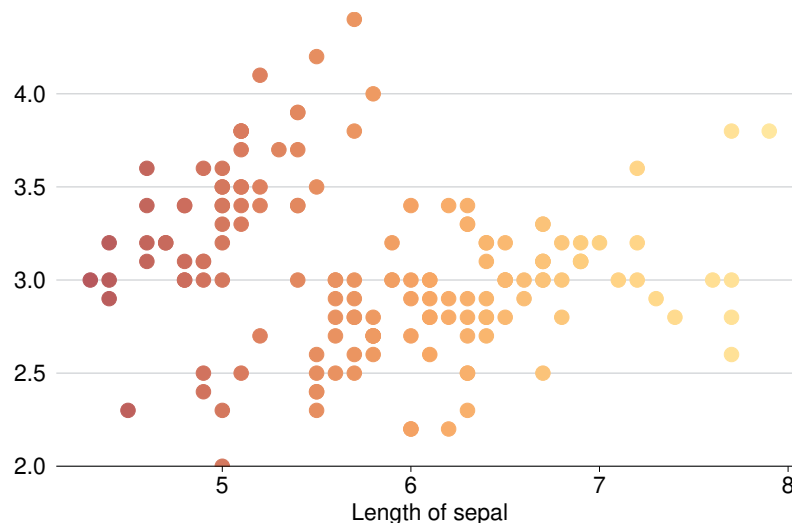
Now we can save our Grattan-formatted graph as a normally-formatted report chart:

```
# Create and store the plot:
plot_final <- p_full_f +
  labs(title = "Iris plants are rad!",
        subtitle = "Width of sepal",
        x = "Length of sepal",
        y = "",
        caption = "Notes: A classic dataset. Source: Fisher (1936)")

# Save the plot
grattan_save(filename = "iris.pdf",
             object = plot_final,
             type = "normal")

#> Note: normal charts remove titles, subtitles, or captions by default.
#> Set `force_labs` to TRUE to retain them, or use type = 'fullslide'
```

Which produces a chart that can fit into a report. Note that the title, subtitle and notes/sources have been removed.

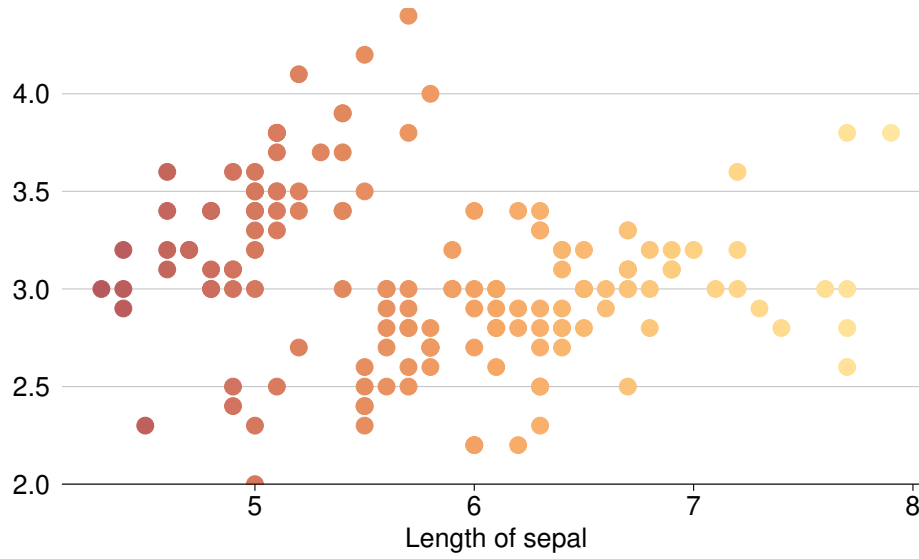


If we wanted to save a full-slide chart, complete with a title, subtitle, notes/sources and the Grattan slide format, we specify `type = "fullslide"`:

```
# Save the plot
grattan_save(filename = "iris_fullslide.pdf",
             object = plot_final,
             type = "fullslide")
```

Iris plants are rad!

Width of sepal



Notes: A classic dataset.
Source: Fisher (1936)