

Using R at Grattan Institute

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Introduction

R is good and cool. Do you want to be good and cool? Use R!

Chapter 1

Using R at Grattan

```
library(tidyverse)
```

This document sets out good practices for structuring your R analysis at Grattan Institute. Having a clear, consistent structure for our analyses means that our work is more easily checked and revised, including by ourselves in the future. A small investment of time up front to set up your analysis will save time (your own and others') down the track.

This guide is designed for *everyone* using R at Grattan. It includes a combination of rules and guidelines.

You should also be aware of the Grattan Institute R Style Guide, which lives in the same place as this document.

Any complaints or comments about this guide can be sent to Will or Matt, respectively.

1.1 Why use R?

It's good and cool!

1.2 Using R projects for a fully reproducible workflow.

Finally adhering to the 'hit by a bus' rule.

Cover: 1. `setwd()` and machine-specific filepaths are bad 2. relative file paths are good 3. RStudio projects are an easy, reproducible way to set your wd

1.2.1 Filepaths

Filepaths should be relative to the working directory, and the working directory should be set by the project.

Good

```
hes <- read_csv("data/HES/hes1516.csv")
grattan_save("images/expenditure_by_income.pdf")
```

Bad

```
hes <- read_csv("/Users/mcowgill/Desktop/hes1516.csv")
hes <- read_csv("C:\\Users\\mcowgill\\Desktop\\hes1516.csv")
grattan_save("/Users/mcowgill/Desktop/images/expenditure_by_income.pdf")
```

1.2.2 Keep your scripts manageable

As a general rule of thumb, use one script per output. It should be clear what your script is trying to do (use comments!).

Consider breaking your analysis into pieces. For example:

- 01_import.R
- 02_tidy.R
- 03_model.R
- 04_visualise.R

Don't include interactive work (like `View(mydf)`, `str(mydf)`, `mean(mydf$variable)`, etc.) in your saved script.

1.2.3 Use subfolders of your project folder

Remember the hit-by-a-bus rule. It should be easy for any Grattan colleague to open your project folder and get up to speed with what it does. Putting all your files - raw data, scripts, output - in the one folder makes it harder to understand how your work fits together.

Use subfolders to clearly separate your code, raw data, and output.

1.3 Grattan coding style guide

Short summary of why

[Link to style guide](#)

1.4 What is the tidyverse and why do we use it?

Introduce following chapters

1.5 An introduction to RMarkdown

1.6 Resources in this package

- Starting a piece of analysis ‘cheat sheet’.
- Updated style guide.
- Written guide/slides.

Chapter 2

Data Visualisation

[intro]

2.1 Set-up and packages

This section uses the package `ggplot2` to visualise data, and `dplyr` functions to manipulate data. Both of these packages are loaded with `tidyverse`. The `scales` package is handy for nicely labelling your axes.

The `grattantheme` package is used to make charts look Grattan-y. The `absmapsdata` package is used to help make maps.

```
library(tidyverse)
library(grattantheme)
library(absmapsdata)
library(sf)
library(scales)
```

For most charts in this chapter, we'll use the `population_table` data summarised here. It contains the population in each state between 2013 and 2018:

```
population_table <- read_csv("data/population_sa4.csv") %>%
  mutate(state_long = state,
         state = strayr::strayr(state_long)) %>%
  filter(data_item == "Persons - Total (no.)") %>%
  mutate(pop = as.numeric(value),
         year = as.factor(year)) %>%
  group_by(year, state) %>%
  summarise(pop = sum(pop))
```

```
# Show the first six rows of the new dataset
head(population_table)
```

```
## # A tibble: 6 x 3
## # Groups:   year [1]
##   year state    pop
##   <fct> <chr>  <dbl>
## 1 2013 ACT     383257
## 2 2013 NSW    7404032
## 3 2013 NT      244684
## 4 2013 Qld    4652824
## 5 2013 SA     1671488
## 6 2013 Tas     512231
```

2.2 Concepts

The `ggplot2` package is based on the grammar of graphics. ...

The main ingredients to a `ggplot` chart:

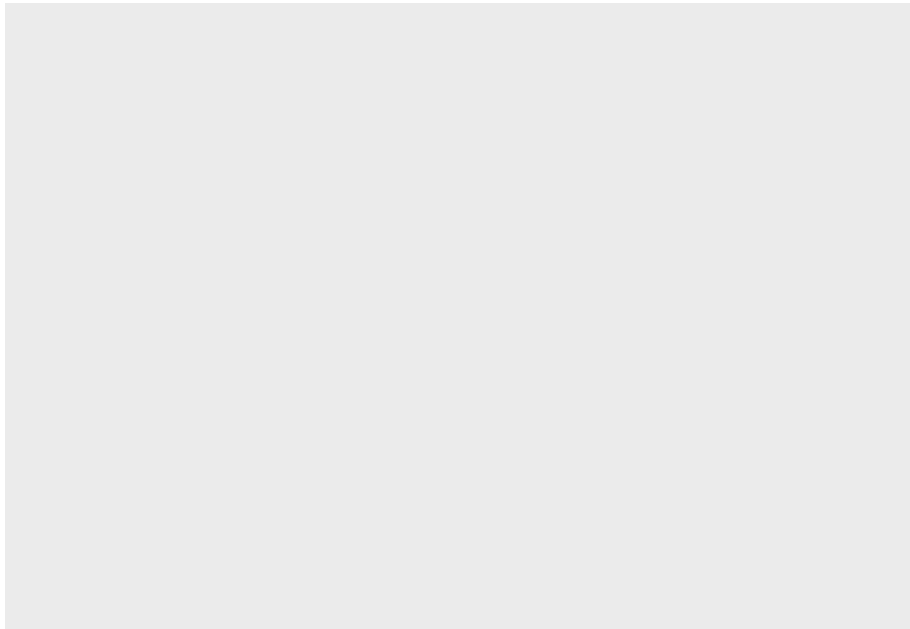
- **Data:** what data should be plotted. e.g. `data`
- **Aesthetics:** what variables should be linked to what chart elements. e.g. `aes(x = population, y = age)` to connect the `population` variable to the x axis, and the `age` variable to the y axis.
- **Geoms:** how the data should be plotted. e.g. `geom_point()` will produce a scatter plot, `geom_col` will produce a column chart.

Each plot you make will be made up of these three elements. The full list of standard geoms is listed in the `tidyverse` documentation.

```
ggplot(data = <DATA>) +
  <GEOM_FUNCTION> (
    mapping = aes(<MAPPINGS>),
    stat = <STAT>,
    position = <POSITION>
  ) +
  <COORDINATE_FUNCTION> +
  <FACET_FUNCTION>
```

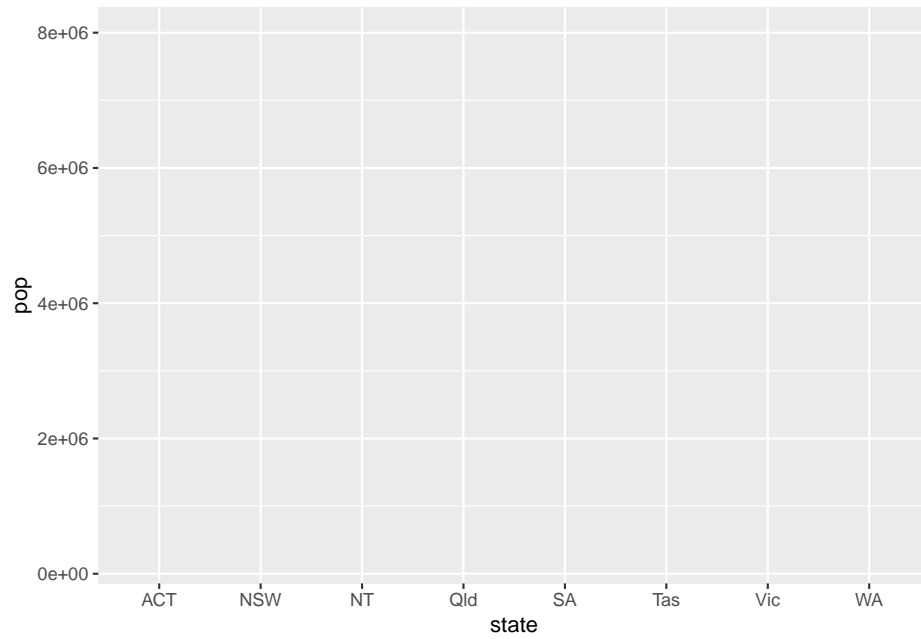
For example, you can plot a column chart by passing the `population_table` dataset into `ggplot()` (“make a chart with this data”). This produces an empty plot:

```
population_table %>%
  ggplot()
```



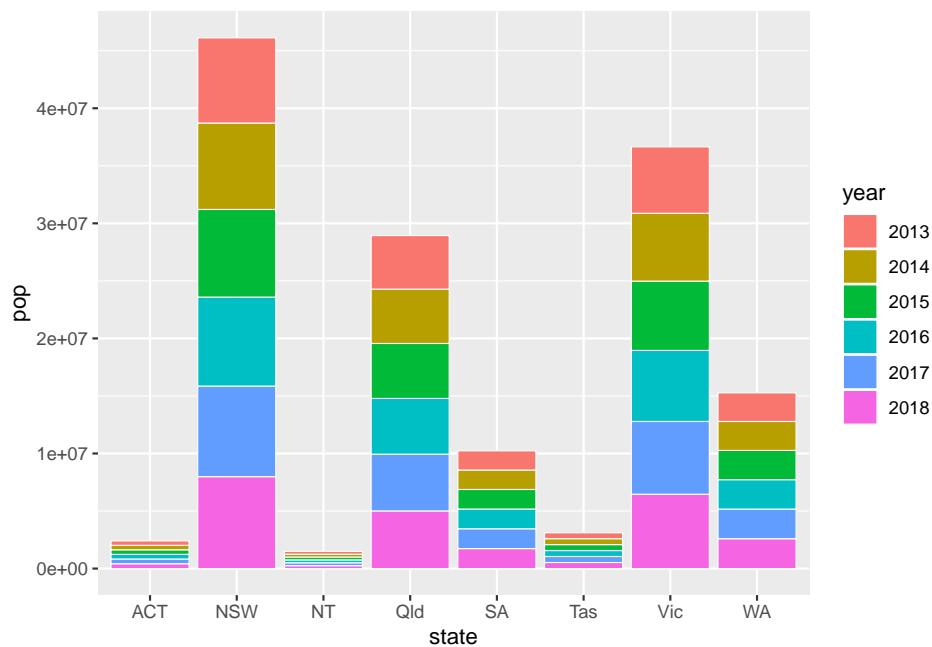
Next, set the `aes` (aesthetics) to `x = state` (“make the x-axis represent state”), `y = pop` (“the y-axis should represent population”), and `fill = year` (“the fill colour represents year”). Now `ggplot` knows where things should *go*:

```
population_table %>%  
  ggplot(aes(x = state,  
             y = pop,  
             fill = year))
```



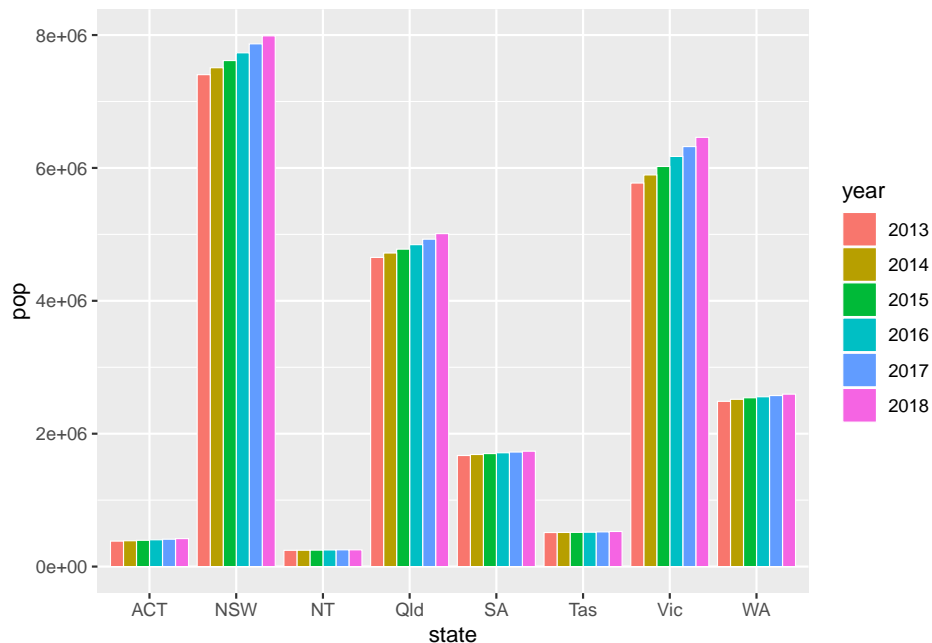
Now that `ggplot` knows where things should go, it needs to *how* to plot them. For this we use `geoms`. Tell it to plot a column chart by using `geom_col`:

```
population_table %>%  
  ggplot(aes(x = state,  
             y = pop,  
             fill = year)) +  
  geom_col()
```



Great! Although stacking populations is a bit silly. You can adjust the way `geoms` work with arguments. In this case, tell it to place the different categories next to each other rather than ontop of each other using `position = "dodge"`:

```
population_table %>%  
  ggplot(aes(x = state,  
             y = pop,  
             fill = year)) +  
  geom_col(position = "dodge")
```



That's nicer. The following sections in this chapter will build on this chart. The rest of the chapter will explore:

- Grattanising your charts and choosing colours
- Saving charts according to Grattan templates
- Making bar, line, scatter and distribution plots
- Making maps and interactive charts
- Adding chart labels

2.3 Making Grattan-y charts

The `grattantheme` package contains functions that help *Grattanise* your charts. It is hosted here: <https://github.com/mattcowgill/grattantheme>

You can install it with `devtools::install_github` from the package:

```
install.packages("devtools")
remotes::install_github("mattcowgill/grattantheme")
```

The key functions of `grattantheme` are:

- `theme_grattan`: set size, font and colour defaults that adhere to the Grattan style guide.
- `grattan_y_continuous`: sets the right defaults for a continuous y-axis.
- `grattan_colour_continuous`: pulls colours from the Grattan colour palette for colour aesthetics.

- `grattan_fill_continuous`: pulls colours from the Grattan colour palette for fill aesthetics.
- `grattan_save`: a save function that exports charts in correct report or presentation dimensions.

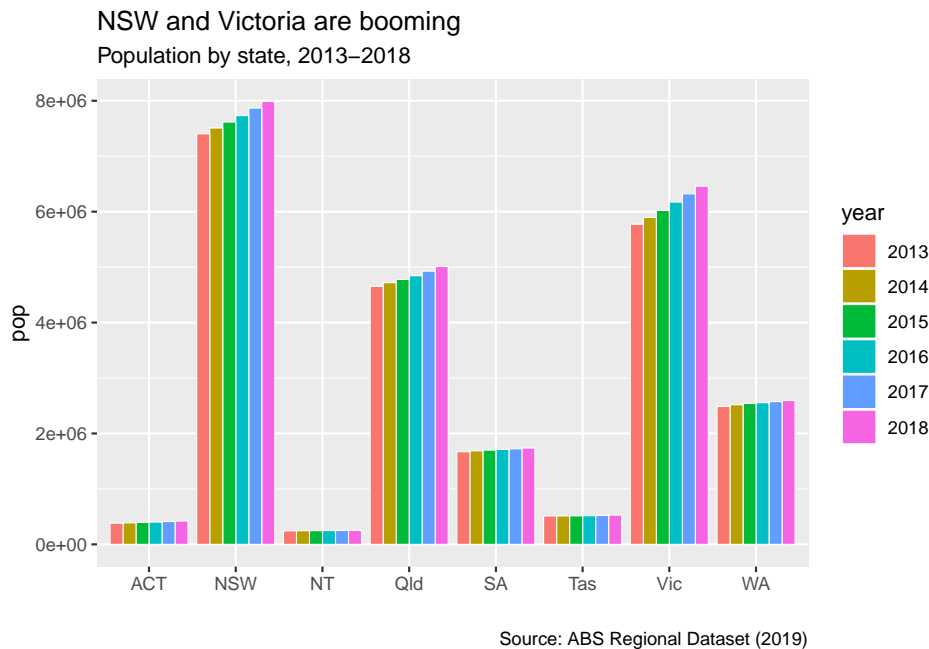
This section will run through some examples of *Grattanising* charts. The `ggplot` functions are explored in more detail in the next section.

2.3.1 Making Grattan charts

Start with a column chart, similar to the one made above:

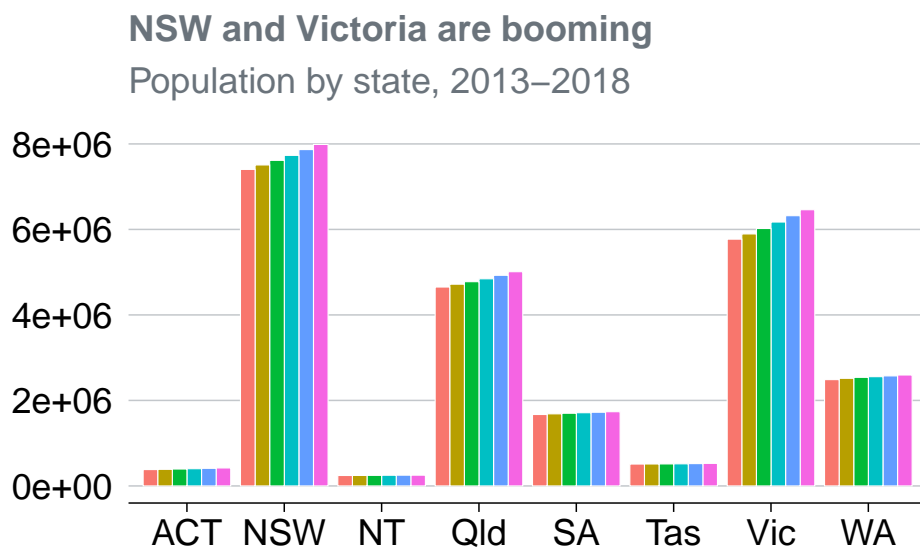
```
base_chart <- population_table %>%
  ggplot(aes(x = state,
             y = pop,
             fill = year)) +
  geom_col(position = "dodge") +
  labs(x = "",
       title = "NSW and Victoria are booming",
       subtitle = "Population by state, 2013-2018",
       caption = "Source: ABS Regional Dataset (2019)")
```

base_chart



Let's make it Grattany. First, add `theme_grattan` to your plot:

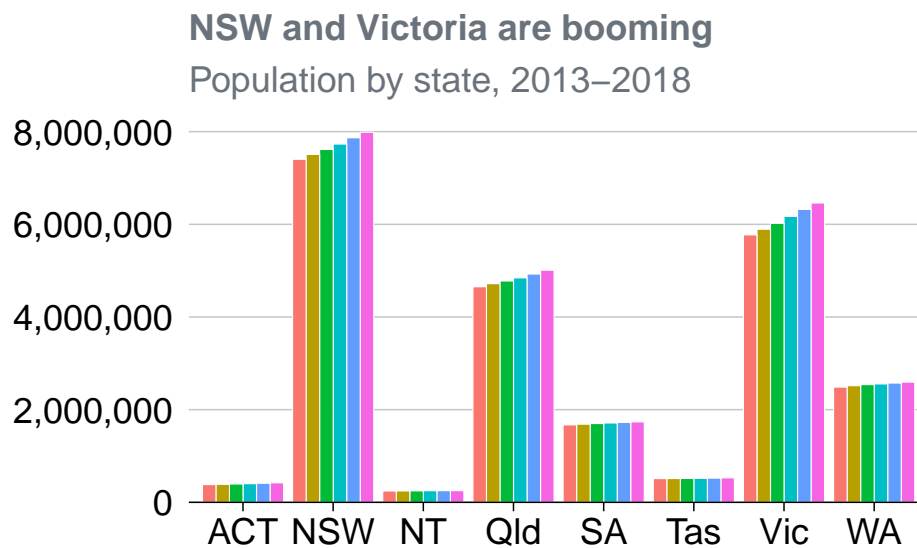
```
base_chart +  
  theme_grattan()
```



Source: ABS Regional Dataset (2019)

Then `grattan_y_continuous` to align the x-axis with zero. This function takes the same arguments as `scale_y_continuous`, so you can add `labels = comma()` to reformat the y-axis labels:

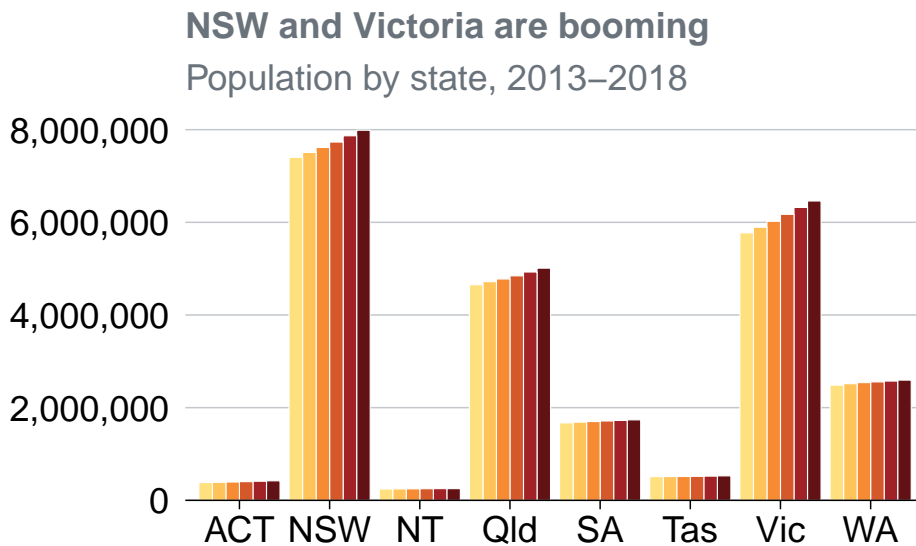
```
base_chart +  
  theme_grattan() +  
  grattan_y_continuous(labels = comma)
```



Source: ABS Regional Dataset (2019)

To define fill colours, use `grattan_fill_manual` with the number of colours you need (six, in this case):

```
pop_chart <- base_chart +  
  theme_grattan() +  
  grattan_y_continuous(labels = comma) +  
  grattan_fill_manual(6)  
  
pop_chart
```



Source: ABS Regional Dataset (2019)

Nice chart! Now you can save it and share it with the world.

2.3.2 Saving Grattan charts

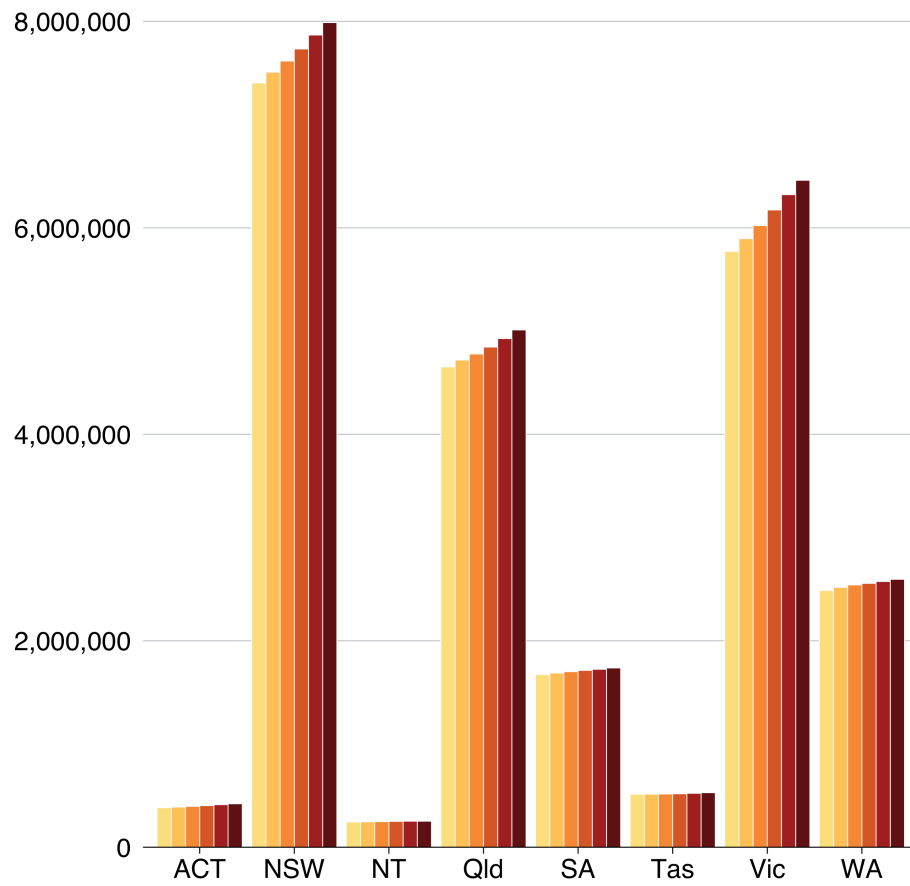
The `grattan_save` function saves your charts according to Grattan templates. It takes these arguments:

- **filename:** the path, name and file-type of your saved chart. eg: "atlas/population_chart.pdf".
- **object:** the R object that you want to save. eg: `pop_chart`. If left blank, it grabs the last chart that was displayed.
- **type:** the Grattan template to be used. This is one of:
 - "normal" The default. Use for normal Grattan report charts, or to paste into a 4:3 Powerpoint slide. Width: 22.2cm, height: 14.5cm.
 - "normal_169" Only useful for pasting into a 16:9 format Grattan Powerpoint slide. Width: 30cm, height: 14.5cm.
 - "tiny" Fills the width of a column in a Grattan report, but is shorter than usual. Width: 22.2cm, height: 11.1cm.
 - "wholecolumn" Takes up a whole column in a Grattan report. Width: 22.2cm, height: 22.2cm.
 - "fullpage" Fills a whole page of a Grattan report. Width: 44.3cm, height: 22.2cm.
 - "fullslide" Creates an image that looks like a 4:3 Grattan Powerpoint slide, complete with logo. Width: 25.4cm, height: 19.0cm.

- "fullslide_169" Creates an image that looks like a 16:9 Grattan Powerpoint slide, complete with logo. Use this to drop into standard presentations. Width: 33.9cm, height: 19.0cm
- "blog" Creates a 4:3 image that looks like a Grattan Powerpoint slide, but with less border whitespace than 'fullslide'."
- "fullslide_44" Creates an image that looks like a 4:4 Grattan Powerpoint slide. This may be useful for taller charts for the Grattan blog; not useful for any other purpose. Width: 25.4cm, height: 25.4cm.
- Set `type = "all"` to save your chart in all available sizes.
- `height`: override the height set by `type`. This can be useful for really long charts in blogposts.
- `save_data`: exports a csv file containing the data used in the chart.
- `force_labs`: override the removal of labels for a particular `type`. eg `force_labs = TRUE` will keep the y-axis label.

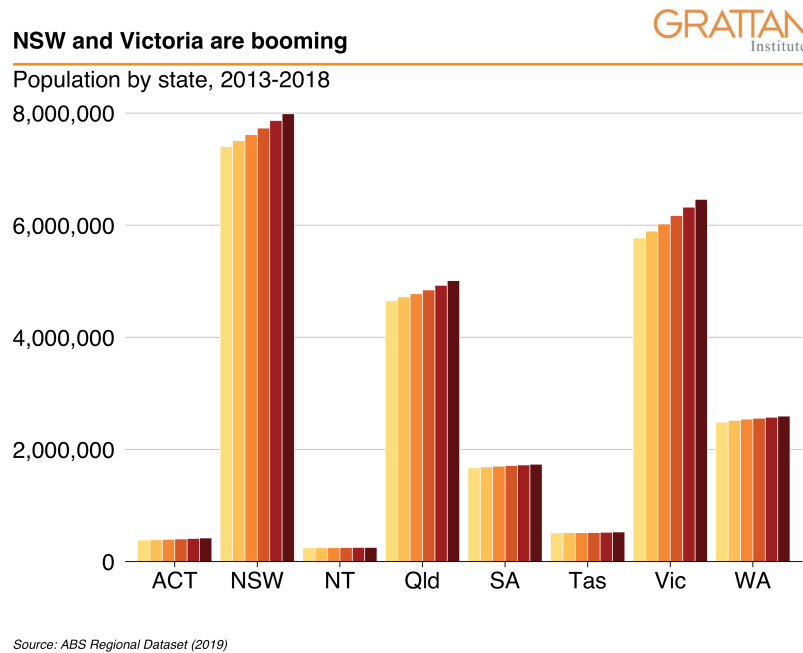
To save the `pop_chart` plot created above as a whole-column chart for a **report**:

```
grattan_save("atlas/population_chart_report.pdf", pop_chart, type = "wholecolumn")
```



To save it as a **presentation** slide instead, use `type = "fullslide"`:

```
grattan_save("atlas/population_chart_presentation.pdf", pop_chart, type = "fullslide")
```

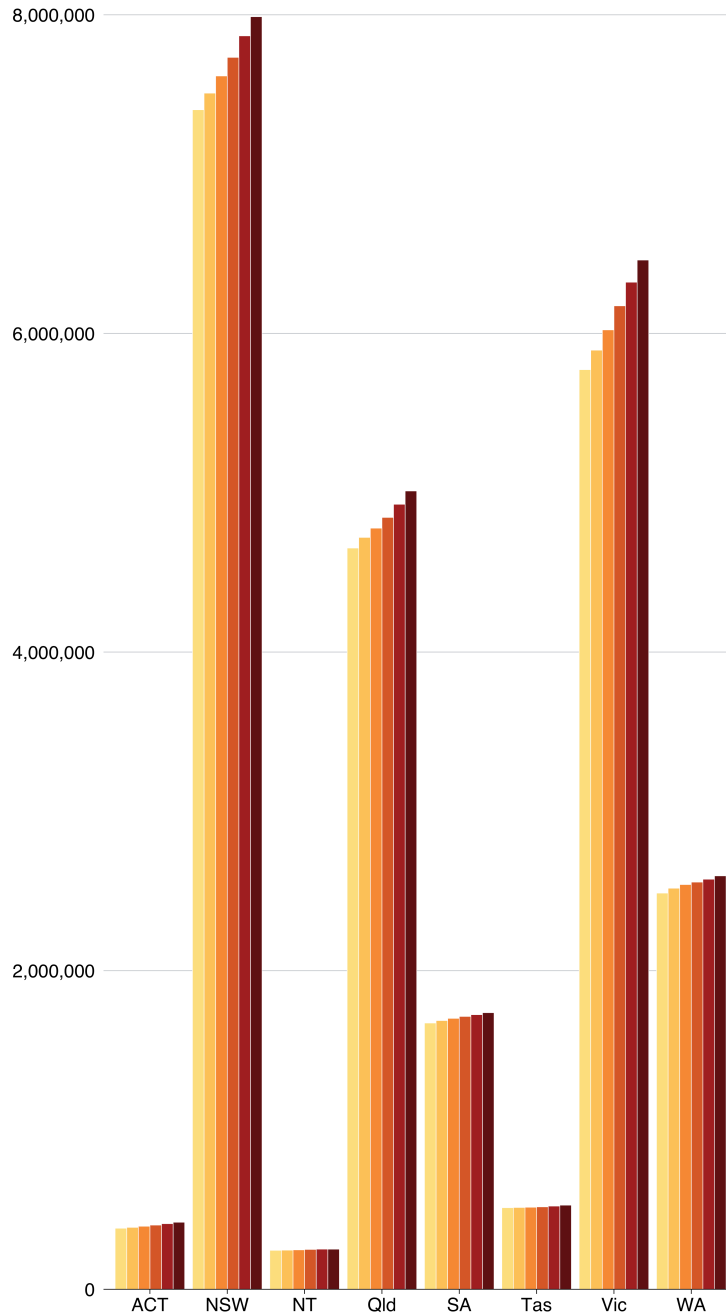


Or, if you want to emphasise the point in a *really tall* chart for a **blogpost**, you can use `type = "blog"` and adjust the `height` to be 50cm. Also note that because this is for the blog, you should save it as a `png` file:

```
grattan_save("atlas/population_chart_blog.png", pop_chart,  
             type = "blog", height = 50)
```

NSW and Victoria are boomingGRATTAN
Institute

Population by state, 2013-2018



Source: ABS Regional Dataset (2019)

And that's it! The following sections will go into more detail about different chart types in R, but you'll mostly use the same basic `grattantheme` formatting you've used here.

2.4 Chart cookbook

This section takes you through a few often-used chart types.

2.4.1 Bar charts

Bar charts are made with `geom_bar` or `geom_col`. Creating a bar chart will look something like this:

```
ggplot(data = <data>) +
  geom_bar(aes(x = <xvar>, y = <yvar>),
    stat = <STAT>,
    position = <POSITION>
  )
```

It has two key arguments: `stat` and `position`.

First, `stat` defines what kind of *operation* the function will do on the dataset before plotting. Some options are:

- "count", the default: count the number of observations in a particular group, and plot that number. This is useful when you're using microdata. When this is the case, there is no need for a `y` aesthetic.
- "sum": sum the values of the `y` aesthetic.
- "identity": directly report the values of the `y` aesthetic. This is how Powerpoint and Excel charts work.

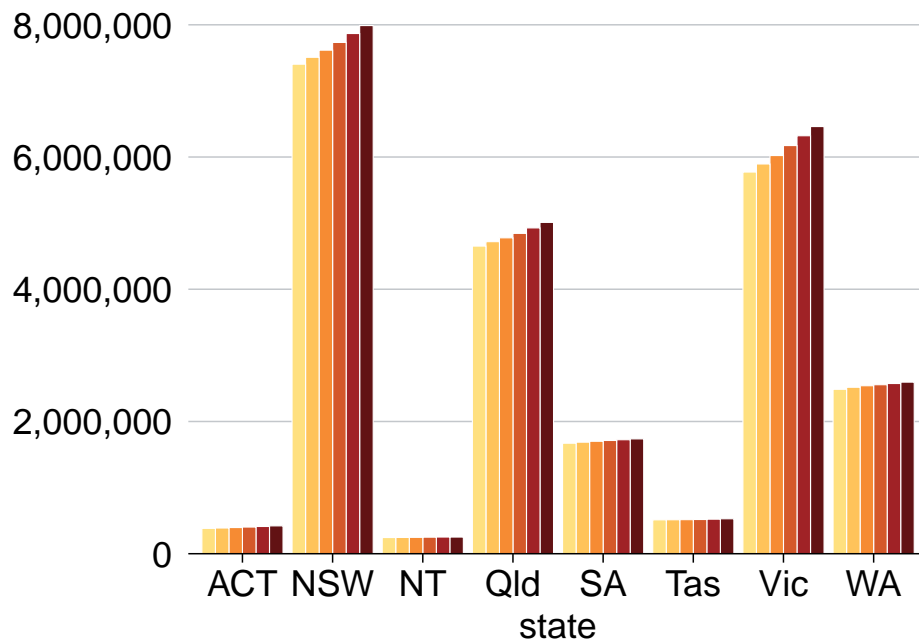
You can use `geom_col` instead, as a shortcut for `geom_bar(stat = "identity")`.

Second, `position`, dictates how multiple bars occupying the same x-axis position will be positioned. The options are:

- "stack", the default: bars in the same group are stacked atop one another.
- "dodge": bars in the same group are positioned next to one another.
- "fill": bars in the same group are stacked and all fill to 100 per cent.

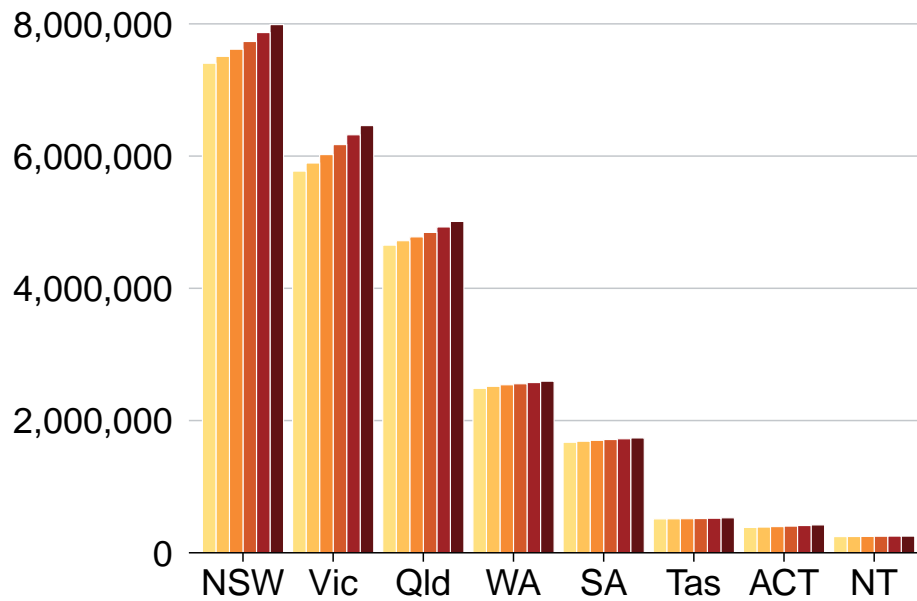
```
population_table %>%
  ggplot(aes(x = state,
    y = pop,
    fill = year)) +
  geom_bar(stat = "identity",
    position = "dodge") +
  theme_grattan() +
```

```
grattan_y_continuous(labels = comma) +
grattan_fill_manual(6)
```



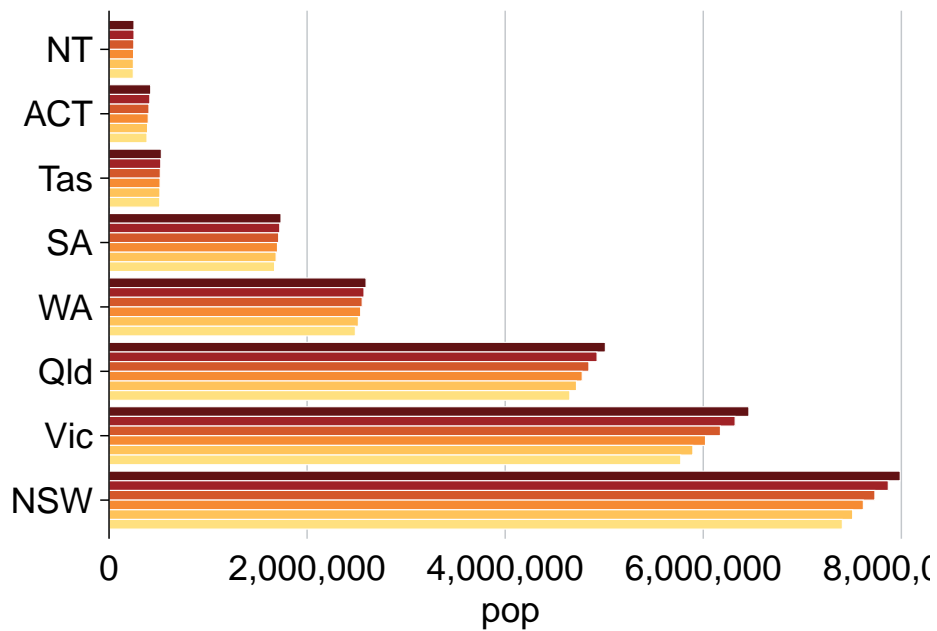
You can also **order** the groups in your chart by a variable. If you want to order states by population, use **reorder** inside **aes**:

```
population_table %>%
  ggplot(aes(x = reorder(state, -pop), # reorder state by negative population
             y = pop,
             fill = year)) +
  geom_bar(stat = "identity",
           position = "dodge") +
  theme_grattan() +
  grattan_y_continuous(labels = comma) +
  grattan_fill_manual(6) +
  labs(x = "")
```



To flip the chart – a useful move when you have long labels – add `coord_flipped` (ie ‘flip coordinates’) and tell `theme_grattan` that the plot is flipped using `flipped = TRUE`.

```
population_table %>%
  ggplot(aes(x = reorder(state, -pop),
              y = pop,
              fill = year)) +
  geom_bar(stat = "identity",
           position = "dodge") +
  coord_flip() + # flip the coordinates
  theme_grattan(flipped = TRUE) + # tell theme_grattan
  grattan_y_continuous(labels = comma) +
  grattan_fill_manual(6) +
  labs(x = "")
```

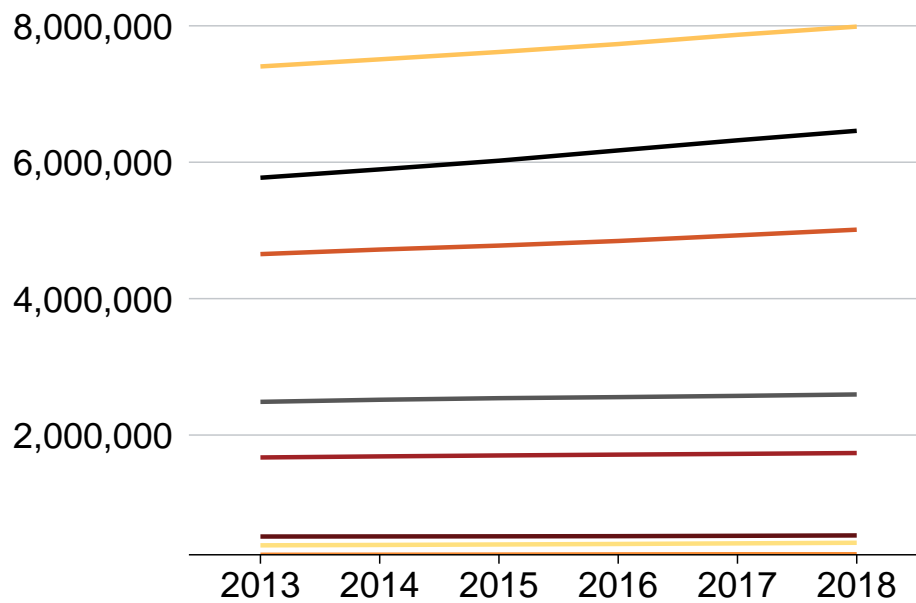


2.4.2 Line charts

A line chart has one key aesthetic: `group`. This tells `ggplot` how to connect individual lines.

```
population_table %>%
  ggplot(aes(x = year,
             y = pop,
             colour = state,
             group = state)) +
  geom_line() +
  theme_grattan() +
  grattan_y_continuous(labels = comma) +
  grattan_colour_manual(8) +
  labs(x = "")
```

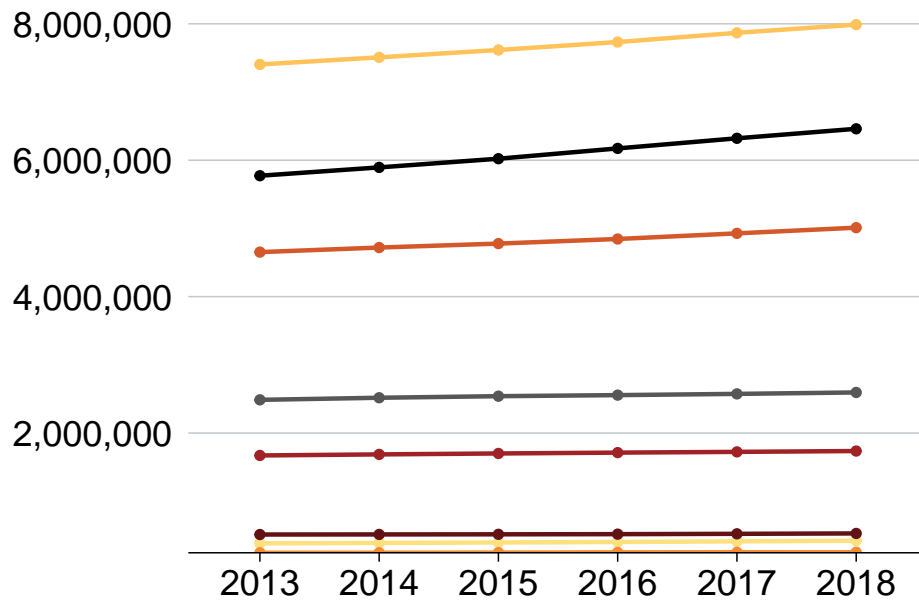
```
## Warning in grattantheme::grattan_pal(n = n, reverse = reverse): Using more
## than six colours is not recommended.
```



You can also add dots for each year by layering `geom_point` on top of `geom_line`:

```
population_table %>%
  ggplot(aes(x = year,
             y = pop,
             colour = state,
             group = state)) +
  geom_line() +
  geom_point(size = 2) +
  theme_grattan() +
  grattan_y_continuous(labels = comma) +
  grattan_colour_manual(8) +
  labs(x = "'')
```

```
## Warning in grattantheme::grattan_pal(n = n, reverse = reverse): Using more
## than six colours is not recommended.
```



If you wanted to show each state individually, you could **facet** your chart so that a separate plot was produced for each state:

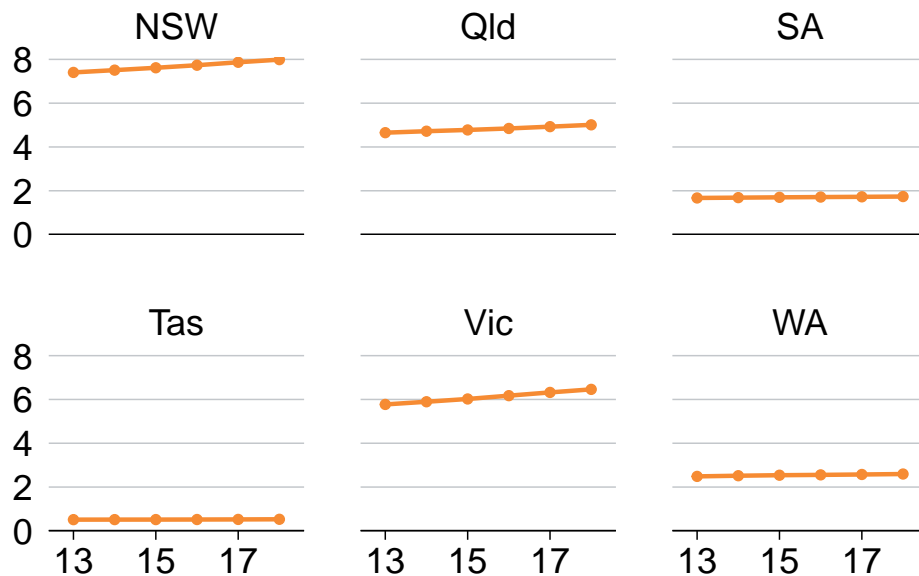
```
population_table %>%
  filter(state != "ACT",
         state != "NT") %>%
  ggplot(aes(x = year,
             y = pop,
             group = state)) +
  geom_line() +
  geom_point(size = 2) +
  theme_grattan() +
  grattan_y_continuous() +
  facet_wrap(state ~ .) +
  labs(x = "")
```



To tidy this up, we can:

1. shorten the years to be “13”, “14”, etc instead of “2013”, “2014”, etc (via the x aesthetic)
2. shorten the y-axis labels to “millions” (via the y aesthetic)
3. add a black horizontal line at the bottom of each facet
4. give the facets a bit of room by adjusting `panel.spacing`
5. define our own x-axis label breaks to just show 13, 15 and 17

```
population_table %>%
  filter(state != "ACT",
         state != "NT") %>%
  ggplot(aes(x = substr(year, 3, 4), # 1: just take the last two characters
             y = pop / 1e6, # 2: divide population by one million
             group = state)) +
  geom_line() +
  geom_point(size = 2) +
  geom_hline(yintercept = 0) + # 3: add horizontal line at the bottom
  theme_grattan() +
  theme(panel.spacing = unit(10, "mm")) + # 4: add panel spacing
  grattan_y_continuous(labels = comma) +
  scale_x_discrete(breaks = c("13", "15", "17")) + # 5: define our own label breaks
  facet_wrap(state ~ .) +
  labs(x = "")
```



2.4.3 Scatter plots

Scatter plots require *x* and *y* aesthetics. These can then be coloured and faceted.

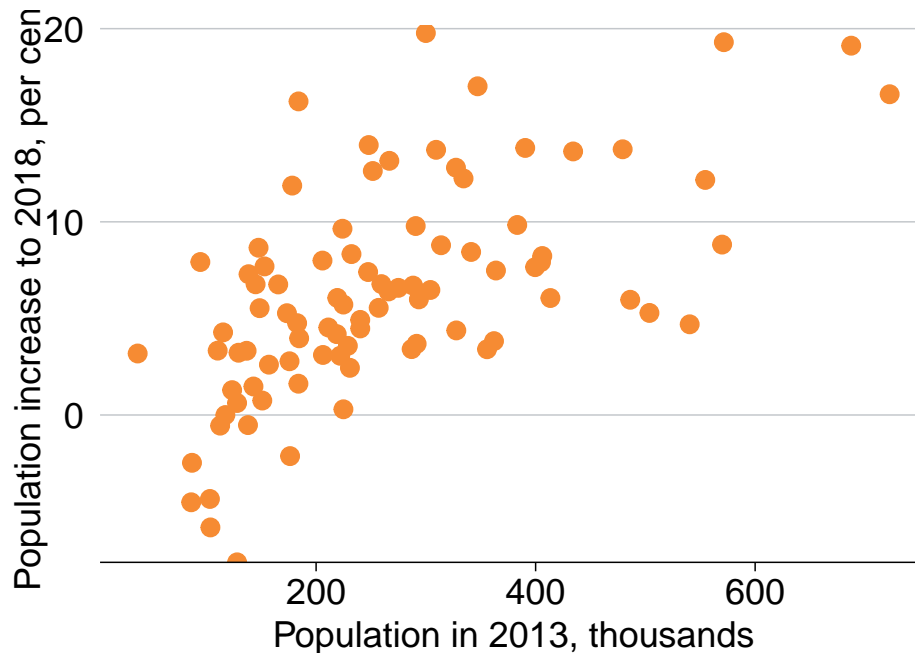
First, create a dataset that we'll use for scatter plots. Take the `population_table` dataset and transform it to have one variable for population in 2013, and another for population in 2018:

```
population_diff <- read_csv("data/population_sa4.csv") %>%
  mutate(state_long = state,
         state = strayr::strayr(state_long),
         pop = as.numeric(value),
         year = as.factor(glue::glue("y{year}"))) %>%
  filter(year %in% c("y2013", "y2018"),
         data_item == "Persons - Total (no.)",
         sa4_name != "Other Territories") %>%
  group_by(year, state, sa4_name) %>%
  summarise(pop = sum(pop)) %>%
  spread(year, pop) %>%
  mutate(pop_change = 100 * (y2018 / y2013 - 1))
```

```
population_diff %>%
  ggplot(aes(x = y2013/1000,
            y = pop_change)) +
  geom_point(size = 4) +
  theme_grattan() +
```

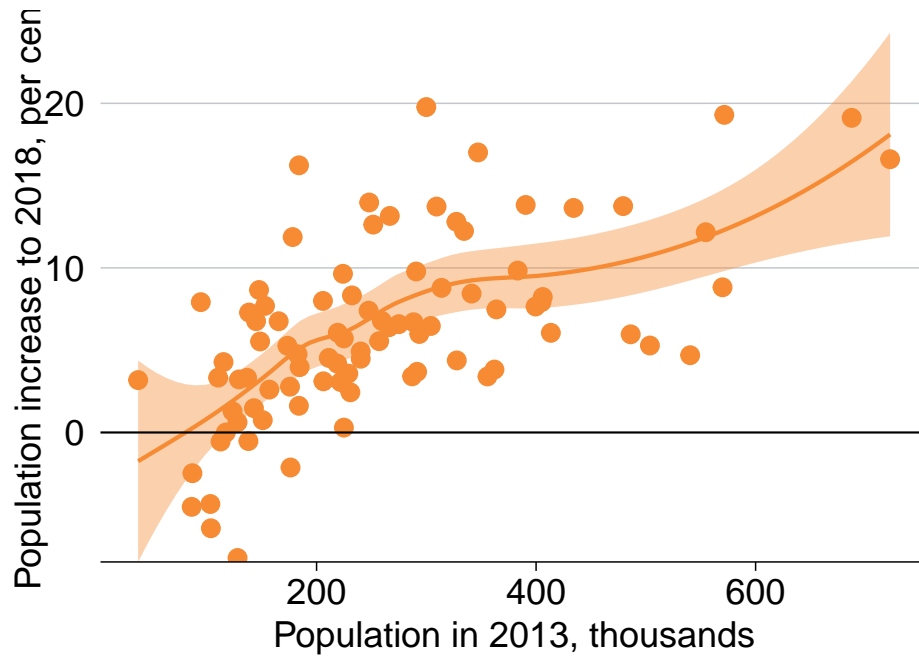


```
theme(axis.title.y = element_text(angle = 90)) +
grattan_y_continuous() +
labs(y = "Population increase to 2018, per cent",
     x = "Population in 2013, thousands")
```



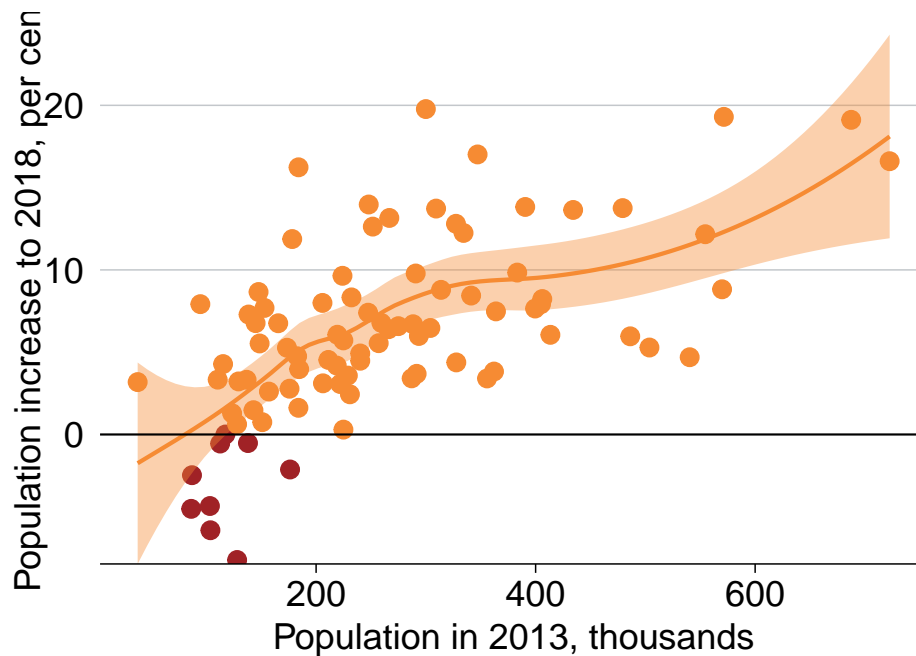
It looks like the areas with the largest population grew the most between 2013 and 2018. To explore the relationship further, you can add a line-of-best-fit with `geom_smooth`:

```
population_diff %>%
  ggplot(aes(x = y2013/1000, # display the x-axis as thousands
            y = pop_change)) +
  geom_point(size = 4) +
  geom_smooth() +
  geom_hline(yintercept = 0) +
  theme_grattan() +
  theme(axis.title.y = element_text(angle = 90)) +
  grattan_y_continuous() +
  labs(y = "Population increase to 2018, per cent",
       x = "Population in 2013, thousands")
```



You could colour-code positive and negative changes from within the `geom_point` aesthetic. Making a change there won't pass through to the `geom_smooth` aesthetic, so your line-of-best-fit will apply to all data points.

```
population_diff %>%
  ggplot(aes(x = y2013/1000, # display the x-axis as thousands
             y = pop_change)) +
  geom_point(aes(colour = pop_change < 0),
             size = 4) +
  geom_smooth() +
  geom_hline(yintercept = 0) +
  theme_grattan() +
  theme(axis.title.y = element_text(angle = 90)) +
  grattan_y_continuous() +
  grattan_colour_manual(2) +
  labs(y = "Population increase to 2018, per cent",
       x = "Population in 2013, thousands")
```



Like the charts above, you could facet this by state to see if there were any interesting patterns. We'll filter out ACT and NT because they only have one and two data points (SA4s) in them, respectively.

```
population_diff %>%
  filter(state != "ACT",
         state != "NT") %>%
  ggplot(aes(x = y2013/1000, # display the x-axis as thousands
             y = pop_change)) +
  geom_point(aes(colour = pop_change < 0),
            size = 2) +
  geom_smooth() +
  geom_hline(yintercept = 0) +
  theme_grattan() +
  theme(axis.title.y = element_text(angle = 90)) +
  grattan_y_continuous() +
  grattan_colour_manual(2) +
  labs(y = "Population increase to 2018, per cent",
       x = "Population in 2013, thousands") +
  facet_wrap(state ~ .)
```



2.4.4 Distributions

```
geom_histogram geom_density
ggribes::
```

2.4.5 Maps

2.4.6 sf objects

[what is]

2.4.7 Using absmappedata

The `absmappedata` contains compressed, and tidied `sf` objects containing geometric information about ABS data structures. The included objects are:

- Statistical Area 1 2011: `sa12011`
- Statistical Area 1 2016: `sa12016`
- Statistical Area 2 2011: `sa22011`
- Statistical Area 2 2016: `sa22016`
- Statistical Area 3 2011: `sa32011`
- Statistical Area 3 2016: `sa32016`

- Statistical Area 4 2011: `sa42011`
- Statistical Area 4 2016: `sa42016`
- Greater Capital Cities 2011: `gcc2011`
- Greater Capital Cities 2016: `gcc2016`
- Remoteness Areas 2011: `ra2011`
- Remoteness Areas 2016: `ra2016`
- State 2011: `state2011`
- State 2016: `state2016`
- Commonwealth Electoral Divisions 2018: `ced2018`
- State Electoral Divisions 2018: `sed2018`
- Local Government Areas 2016: `lga2016`
- Local Government Areas 2018: `lga2018`

You can install the package from Github. You will also need the `sf` package installed to handle the `sf` objects.

```
devtools::install_github("wfmackey/absmapsdata")
library(absmapsdata)

install.packages("sf")
library(sf)
```

2.4.7.1 Making choropleth maps

Choropleth maps break an area into ‘bits’, and colours each ‘bit’ according to a variable.

SA4 is the largest non-state statistical area in the ABS ASGS standard.

You can join the `sf` objects from `absmapsdata` to your dataset using `left_join`. The variable names might be different – eg `sa4_name` compared to `sa4_name_2016` – so use the `by` function to match them.

```
map_data <- population_diff %>%
  left_join(sa42016, by = c("sa4_name" = "sa4_name_2016"))

head(map_data %>%
  select(sa4_name, geometry))

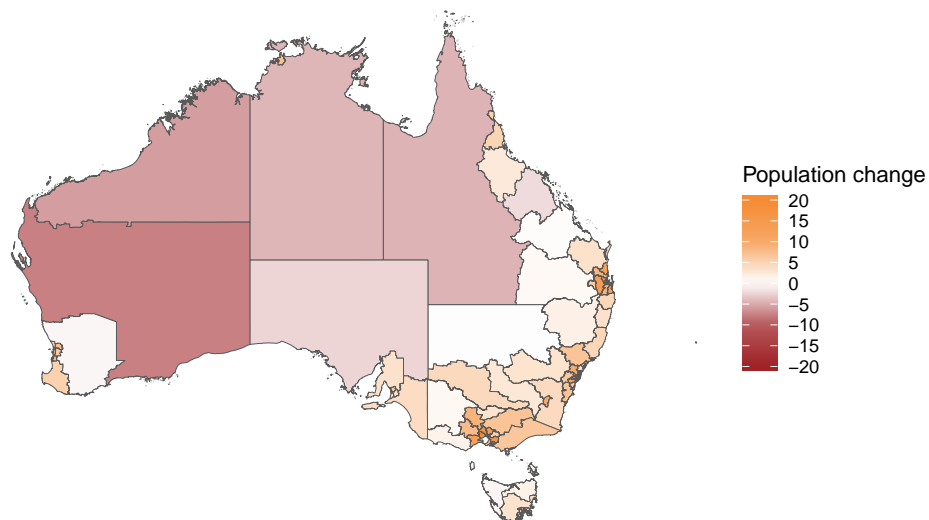
## # A tibble: 6 x 3
## # Groups:   state [2]
##   state sa4_name geometry
##   <chr> <chr>      <MULTIPOLYGON [°]>
## 1 ACT   Australian Capita~ (((148.8041 -35.71402, 148.8018 -35.7121, 148.7~
## 2 NSW   Capital Region    (((150.3113 -35.66588, 150.3126 -35.66814, 150.~
## 3 NSW   Central Coast     (((151.315 -33.55582, 151.3159 -33.55503, 151.3~
## 4 NSW   Central West      (((150.6107 -33.06614, 150.6117 -33.07051, 150.~
```

```
## 5 NSW Coffs Harbour - G~ (((153.2785 -29.91874, 153.2773 -29.92067, 153.~
## 6 NSW Far West and Orana (((150.1106 -31.74613, 150.1103 -31.74892, 150.~
```

You then plot a map like you would any other `ggplot`: provide your data, choose your `aes` and your `geom`. For maps with `sf` objects, the key **aesthetic** is `geometry = geometry`, and the **geom** is `geom_sf`.

```
map <- map_data %>%
  ggplot(aes(geometry = geometry,
             fill = pop_change)) +
  geom_sf(lwd = 0) +
  theme_void() +
  grattan_fill_manual(discrete = FALSE,
                     palette = "diverging",
                     limits = c(-20, 20),
                     breaks = seq(-20, 20, 5)) +
  labs(fill = "Population change")
```

```
map
```



2.5 Creating simple interactive graphs with plotly

```
plotly::ggplotly()
```

2.6 bin: generate data used (before prior sections are constructed)

```
library(tidyverse)
library(janitor)
library(absmapsdata)

data <- read_csv("data/ABS_REGIONAL_ASGS2016_02082019164509969.csv") %>%
  clean_names() %>%
  select(data_code = measure,
         data_item,
         asgs = regiontype,
         sa4_code_2016 = asgs_2016,
         sa4_name_2016 = region,
         year = time,
         value) %>%
  mutate(sa4_code_2016 = as.character(sa4_code_2016)) %>%
  left_join(sa42016 %>% select(sa4_code_2016, state_name_2016)) %>%
  rename(state = state_name_2016,
         sa4_code = sa4_code_2016,
         sa4_name = sa4_name_2016)

write_csv(data, "data/population_sa4.csv")
```


Chapter 3

Reading data

3.1 Importing data

3.1.1 Reading CSV files

3.1.1.1 `read_csv()`

The `read_csv()` function from the `tidyverse` is quicker and smarter than `read.csv` in base R.

Pitfalls: 1. `read_csv` is quicker because it surveys a sample of the data

We can also compress `.csv` files into `.zip` files and read them *directly* using `read_csv()`:

```
read_csv("data/my_data.zip")
```

This is useful for two reasons:

1. The data takes up less room on your computer; and
2. The original data, which shouldn't ever be directly edited, is protected and cannot be directly edited.

3.1.1.2 `data.table::fread()`

The `fread` function from `data.table` is quicker than both `read.csv` and `read_csv`.

3.1.2 `readxl::read_excel()`

3.1.3 `rio`

3.1.4 `readabs`

3.2 Reading common files:

- TableBuilder CSVSTRINGS
- HES household file
- SIH
- LSAY and derivatives

See data directory for a list of microdata available to Grattan.

3.3 Appropriately renaming variables

As shown in the style guide

Add `rename_abs` function to a common Grattan package?

3.4 Getting to tidy data

`pivot_long()` and `pivot_wide()` *Make sure these are stable btw*

Chapter 4

Different data types

4.1 Tidy data

Other data structures

4.2 Dates with `lubridate::`

The `lubridate::` package

4.3 Strings with `stringr::`

- Replacing values
- Matching values
- Separating columns

4.4 Factors with `forcats::`

- Dangers with factors

Chapter 5

Data transformation

5.1 The pipe

5.2 Key dplyr functions:

All have the same syntax structure, which enable pipe-chains.

5.3 Filter with `filter()`

5.4 Arrange with `arrange()`

5.5 Select variables with `select()`

5.6 Group data with `group_by()`

5.7 Edit and add new variables with `mutate()`

5.7.1 Cases when you should use `case_when()`

5.8 Summarise data with `summarise()`

5.9 Joining datasets with `*_join()`

Chapter 6

Analysis

Chapter 7

Creating functions

7.1 It can be useful to make your own function

Why on earth would you create your own function?

7.2 Defining simple functions

7.3 More complex functions

7.4 Sets of functions

7.5 Using `purrr::map`

7.6 Sharing your useful functions with Grattan

Chapter 8

Version control

8.1 Version control is important and intimidating

Version control is great!

8.2 Github

We use Github to version-control and share reports in LaTeX, so you're already a bit set-up.

8.3 Git

Using Git within R Studio...