

# Exploring the Wide World of ggplot2 Extensions

Instructor notes

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```
library(palmerpenguins)
library(ggplot2)
```

## Intro

Go through introductory [slides](#) to start. Spend time looking through the ggplot extension gallery and ask if there are any particular packages that look useful to folks. Get people to share their discoveries.

## Demonstrations

Go through vignette or other short demo of several extension packages.

### **gganimate**

```
library(gganimate)
```

- Animate plots and save as gifs
- Takes some getting used to lingo
- Can control: what changes over time with `transition_*`, how the view of the axes changes with `view_*`, how much “memory” of the change there is, with `shadow_*`, and how data appears and disappears with `enter_*` and `exit_*`
- Use `?` with functions to figure out what label name is

- These penguin plots don't really make sense because the data isn't same individuals over the years; should set group in `transition_states`
- Mention `animate()` for more control
- Resources:
  - [Bat data slides](#)
  - [Penguins code source](#)

## esquisse

```
library(esquisse)
library(dplyr)
```

- “Esquisse” means rough first sketch in French
- `esquisse` [vignette](#)
- Point and click to generate ggplot plots from data
- Good for exploring datasets (look at data too)
- Still have to modify dataset outside of tool, e.g., recoding factors

```
penguins <- penguins %>% na.omit()
esquisser(penguins)
```

- Helpful things: color codes variables, suggests plot types
- Save plot as image file, or copy and paste code into console (uses `tidyverse`)
- Can add to Shiny app to let non-R users create ggplots of data

## plotly

```
library(plotly)
```

- `plotly` R package is interface to a JavaScript library for making interactive data visualizations
- Only important to know that because if you search for `plotly` help, sometimes you'll get JavaScript code examples.

- `plotly` *kind of* uses the grammar of graphics, but building a plot from scratch is tricky
- `plotly` functions mostly take lists as arguments, which makes it very difficult to figure out what default values are or even what arguments functions can take.
- `ggplotly()` **transforms** `ggplot` objects into interactive `plotly` plots. 90% of the time it gets you 90% of the way there

```
p <-
  ggplot(penguins, aes(x = bill_length_mm, y = bill_depth_mm, color = species)) +
  geom_point() +
  labs(x = "Bill length (mm)", y = "Bill depth (mm)", color = "Penguin Species") +
  theme_bw()

ggplotly(p)
```

By default, you get:

- Drag a box to zoom, double-click to zoom out
- Tooltips on hover
- Ability to hide and show points by clicking on the legend
- Toolbar with tools including ability to download a PNG of the plot

This works in R Markdown or Quarto documents (i.e. there's not a way to make these plots “stand-alone”—they're rendered as HTML)

## Types of plots

Works for most (all?) built-in geoms. Works for some plots made with `ggplot2` extensions also. Demo a few different geoms like `geom_boxplot()`, `geom_histogram()`, `geom_smooth()`

## Customize tooltip

You can customize the info displayed in the tooltip with `ggplotly`. Give it additional aesthetic **text** to include something *only* as a tooltip.

```

p <-
  ggplot(penguins, aes(x = bill_length_mm, y = bill_depth_mm, color = species)) +
  #Ignore unknown aesthetics warning
  geom_point(aes(text = sex)) +
  labs(x = "Bill length (mm)", y = "Bill depth (mm)", color = "Penguin Species") +
  theme_bw()

ggplotly(p, tooltip = c("text", "bill_depth_mm", "bill_length_mm"))

```

## Animations

You can use `ggplotly()` to make interactive animated plots by using the `frame` aesthetic

```

library(gapminder) #dataset
head(gapminder)
p2 <- ggplot(gapminder, aes(gdpPercap, lifeExp, color = continent)) +
  geom_point(aes(size = pop, frame = year, ids = country)) +
  scale_x_log10()

ggplotly(p2)

```

## ggrepel

The `ggrepel` package is helpful for directly labeling plots, especially when labels would otherwise overlap.

```

#make rownames into columns
dat <- mtcars |> tibble::rownames_to_column(var = "car")
dat
p3 <-
  ggplot(dat,
    aes(wt, mpg, label = car, colour = factor(cyl))) +
  geom_point()
#without ggrepel
p3 + geom_text()
p3 + geom_label()
library(ggrepel)
p3 + geom_text_repel()
p3 + geom_label_repel()

```

## Adjust appearance of labels

By default, little segments are drawn connecting labels to points only when labels are far enough away. This can be adjusted.

```
p3 + geom_label_repel(min.segment.length = 0)
```

## Overlapping labels

You see a warning about labels being removed due to overlaps. Make text smaller, plot larger, adjust `max.overlaps`, force labels further apart, make it try harder to find positions that don't overlap.

```
#don't usually need to do all of these things, but they're some options
p4 <-
  p3 +
    geom_label_repel(
      min.segment.length = 0,
      #smaller labels
      size = 3,
      label.padding = 0.15,
      #draw labels even if overlapping with things
      max.overlaps = 15, #default is 10, increase to get more labels drawn
      #force labels further from point
      force = 5,
      #ask it to try harder
      max.time = 2,
      max.iter = 15000,
      show.legend = FALSE
    )
p4
```

## Labels are random

Labels are positioned randomly, so they're different every time a plot is rendered

```
p4
p4
```

You can `set.seed()` at the top of your script to ensure reproducibility.

```
set.seed(123)
p4
set.seed(123)
p4
```

## Label just some points

You can label just select points

```
cars <- c("Volvo 142E", "Merc 230")
ggplot(dat, aes(wt, mpg, label = ifelse(car %in% cars, car, ""))) +
  geom_point(color = "red") +
  geom_label_repel(min.segment.length = 0, nudge_x = 0.5, nudge_y = 3)
```

## Wrap-up

Go through “where to find help” slides. Plug next workshop (our fall reproducibility series)

Ask if there are any other remaining `ggplot2`-related roadblocks people are experiencing and workshop it live.