

# R COURSE

## Data visualization

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# R - Session 03

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- Graph types
- Grammar of graphics
- Playing with ggplot2
- Multiple graphs
- ggplot2 syntax

# Intro to Data vizualisation

# Installation and Resources

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## Packages

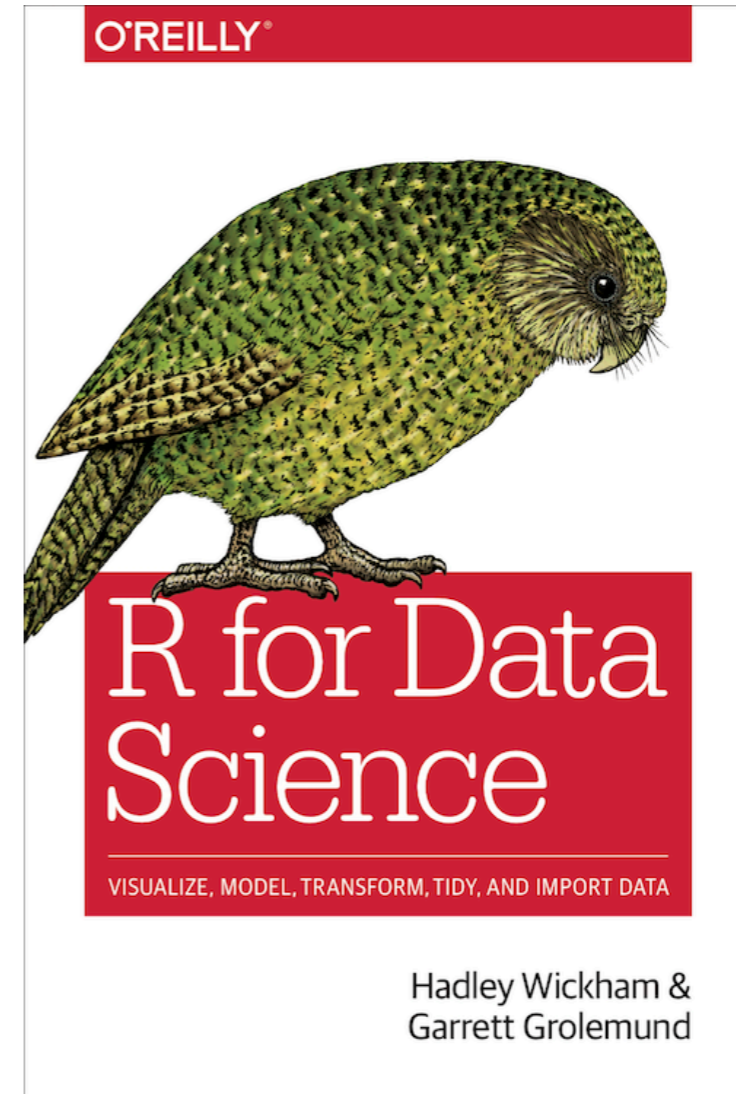
- `ggplot2`
- `patchwork`

## Reading

- [Chapter 28 of R for data science](#)

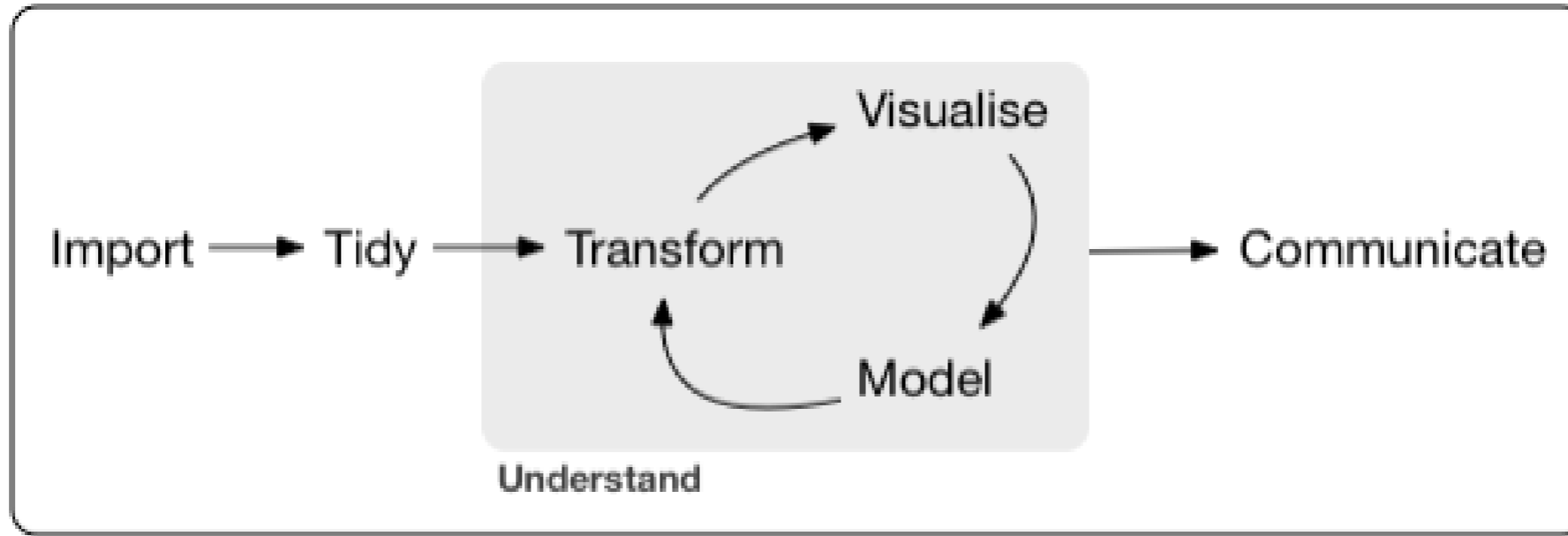
## Resources

- [Fundamental of data visualization](#)
- [Data visualization: practical introduction](#)



# Workflow

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## Graph purposes

- **Analysis graphs**

- design to see patterns, trends
- aid the process of data description
- interpretation

- **Presentation graphs**

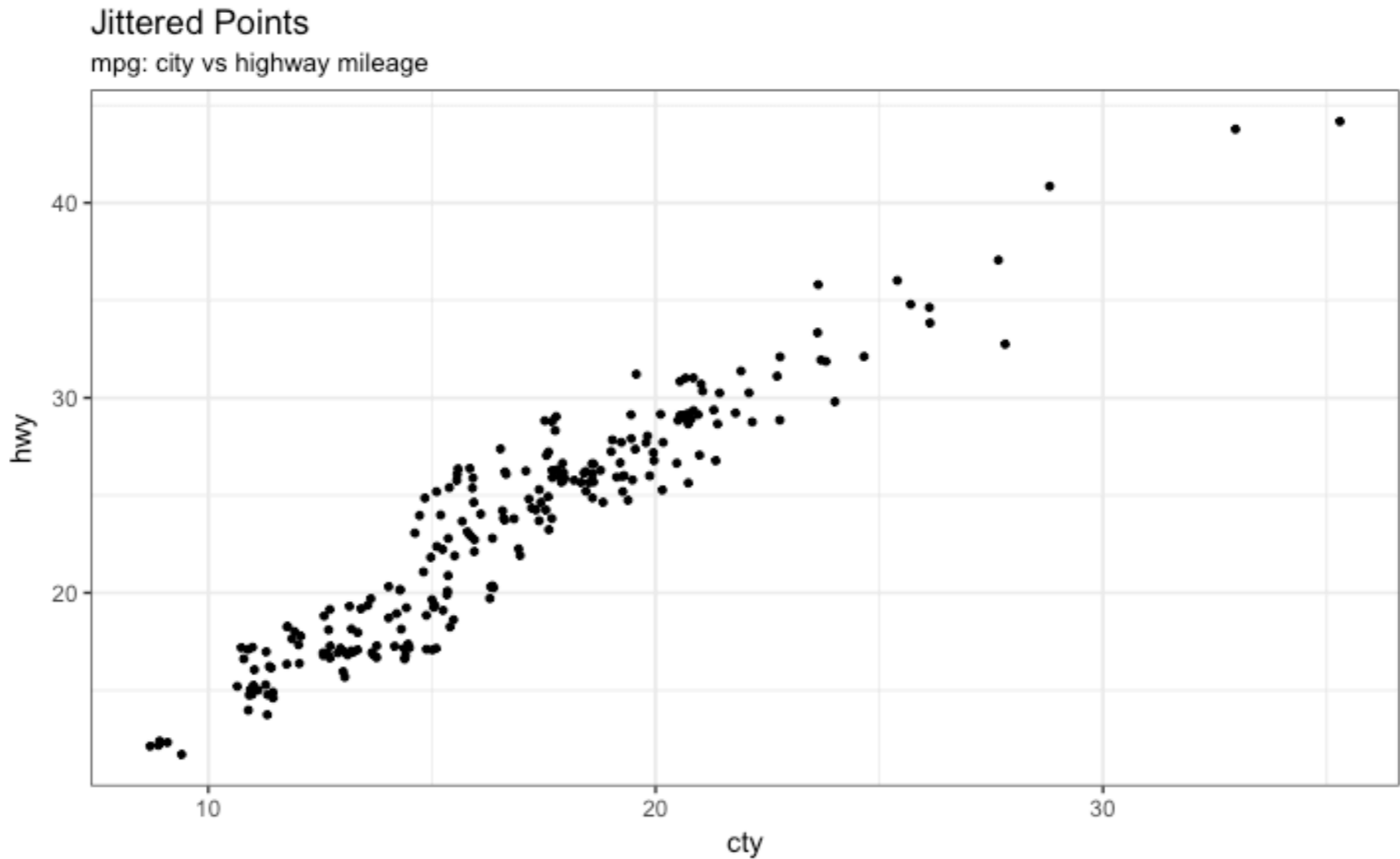
- design to attract attention
- make a point
- illustrate a conclusion

Source: [Michael Friendly](#)

# Graph types

# Jitter

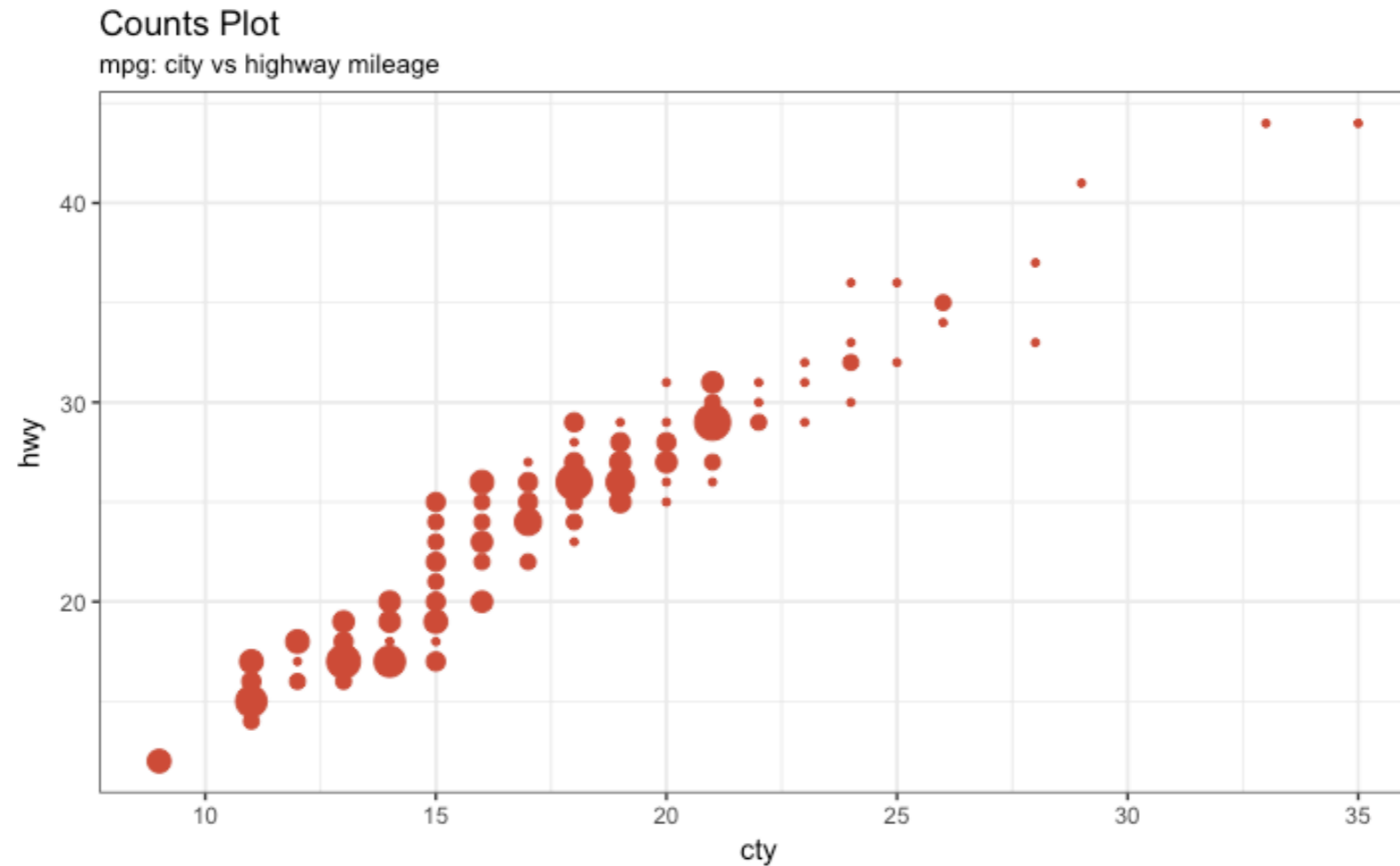
- Two variables numerical



# Bubble

---

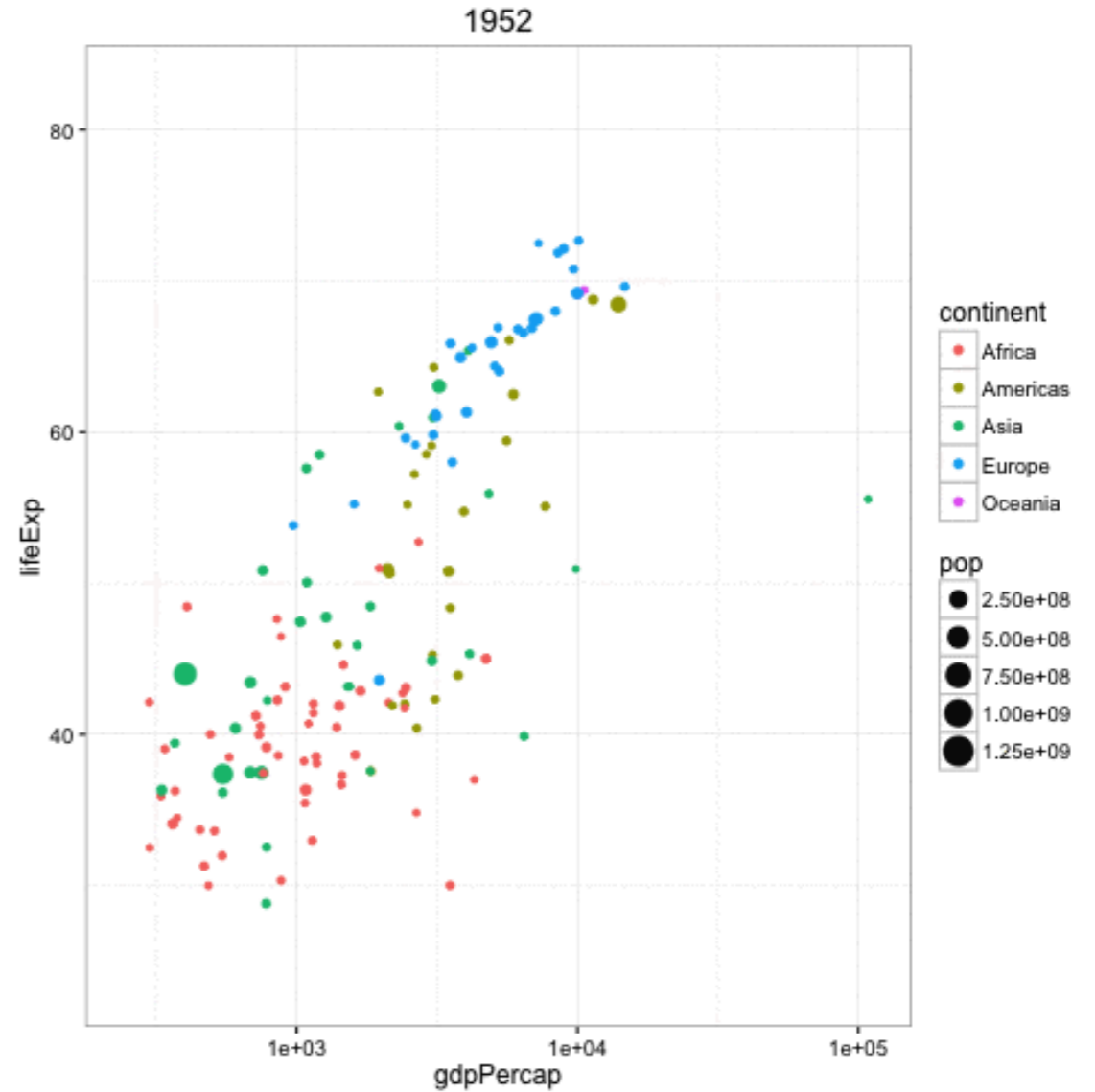
- Two variables numerical
- **Add another variable numerical**





# Animate

- Two variables numerical
- One variable numerical
- One variable categorical
- **Animate another variable**

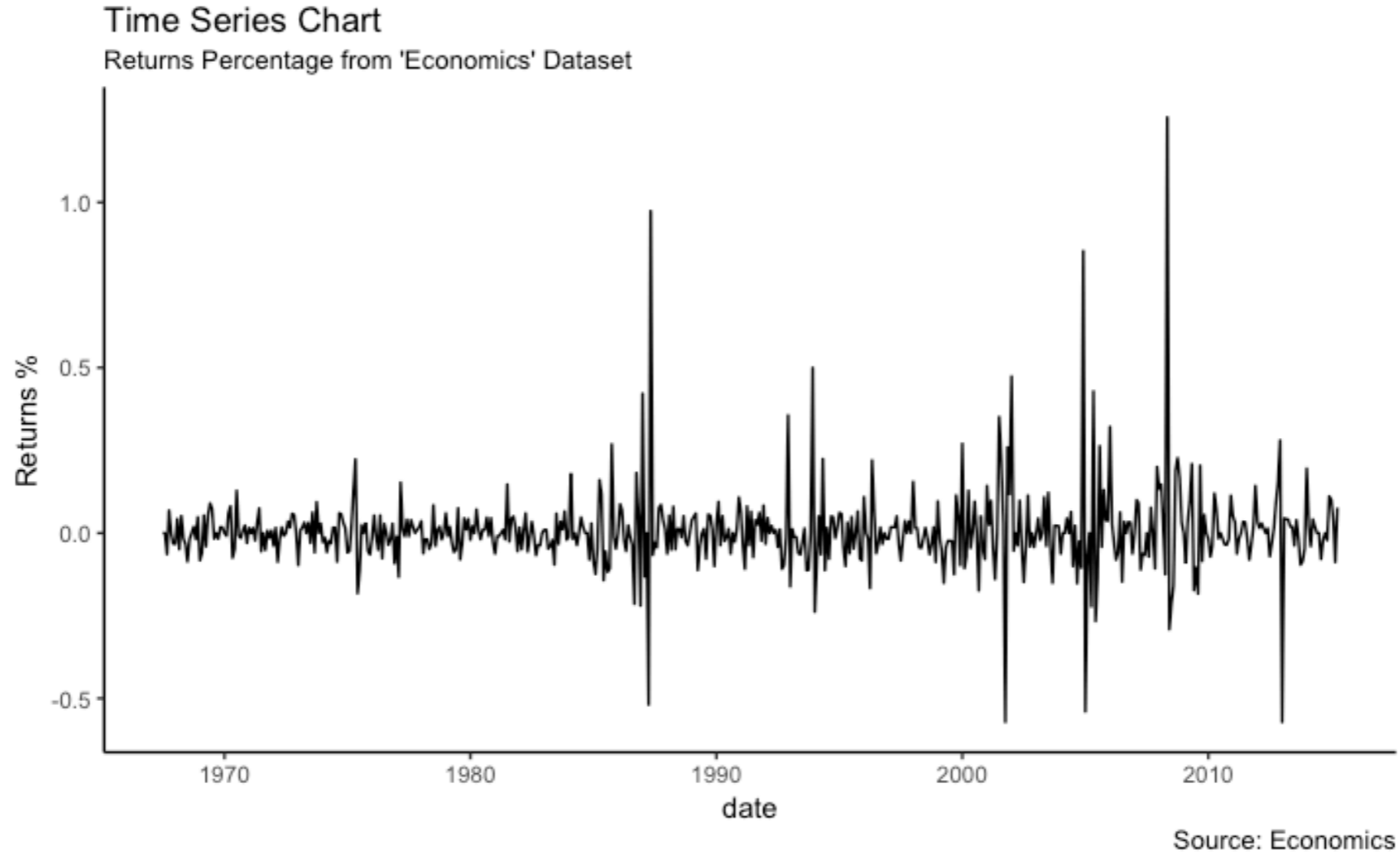




# Times series

---

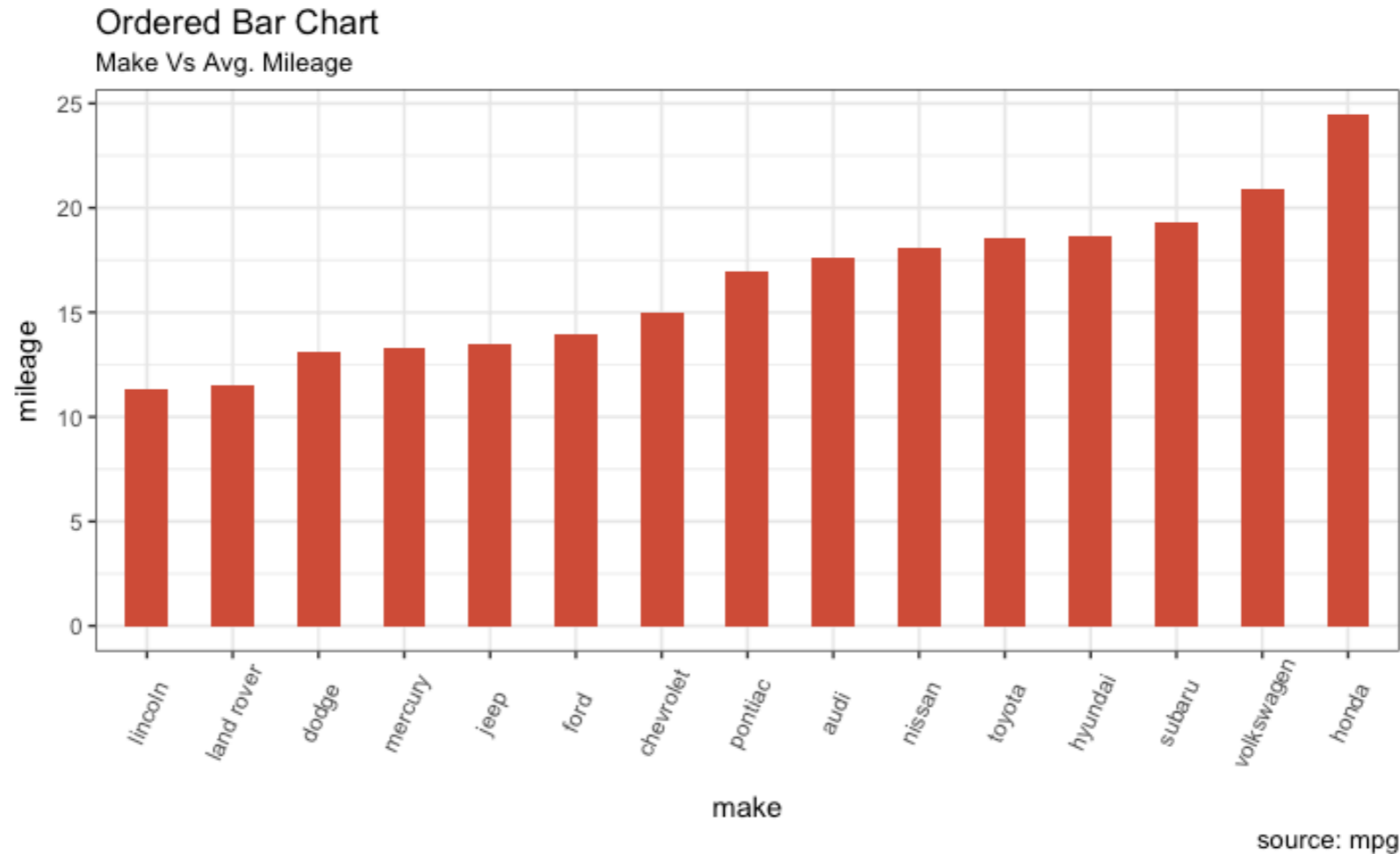
- Line graph



# Bargraphs

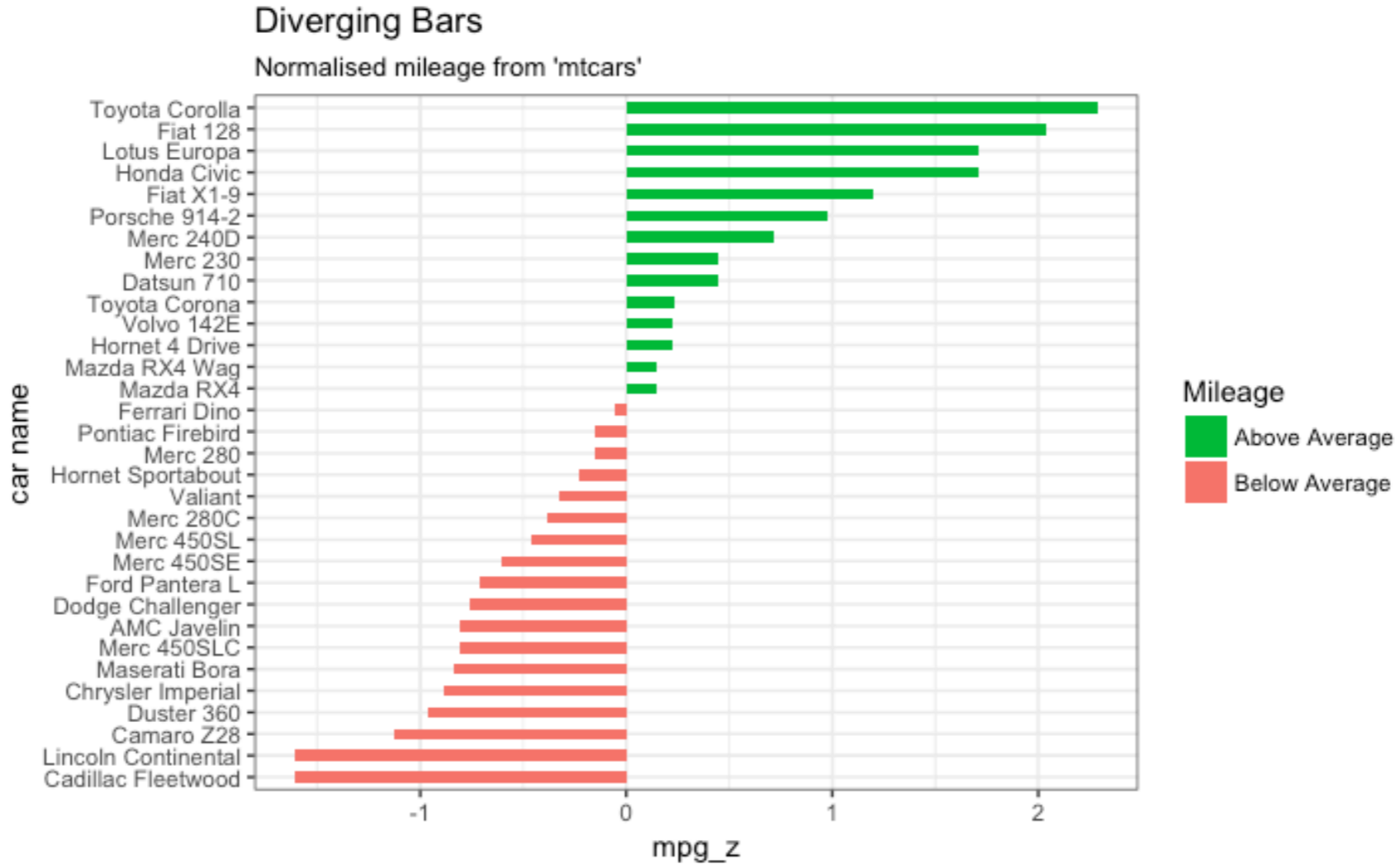
---

- One variable categorical
- One variable numerical



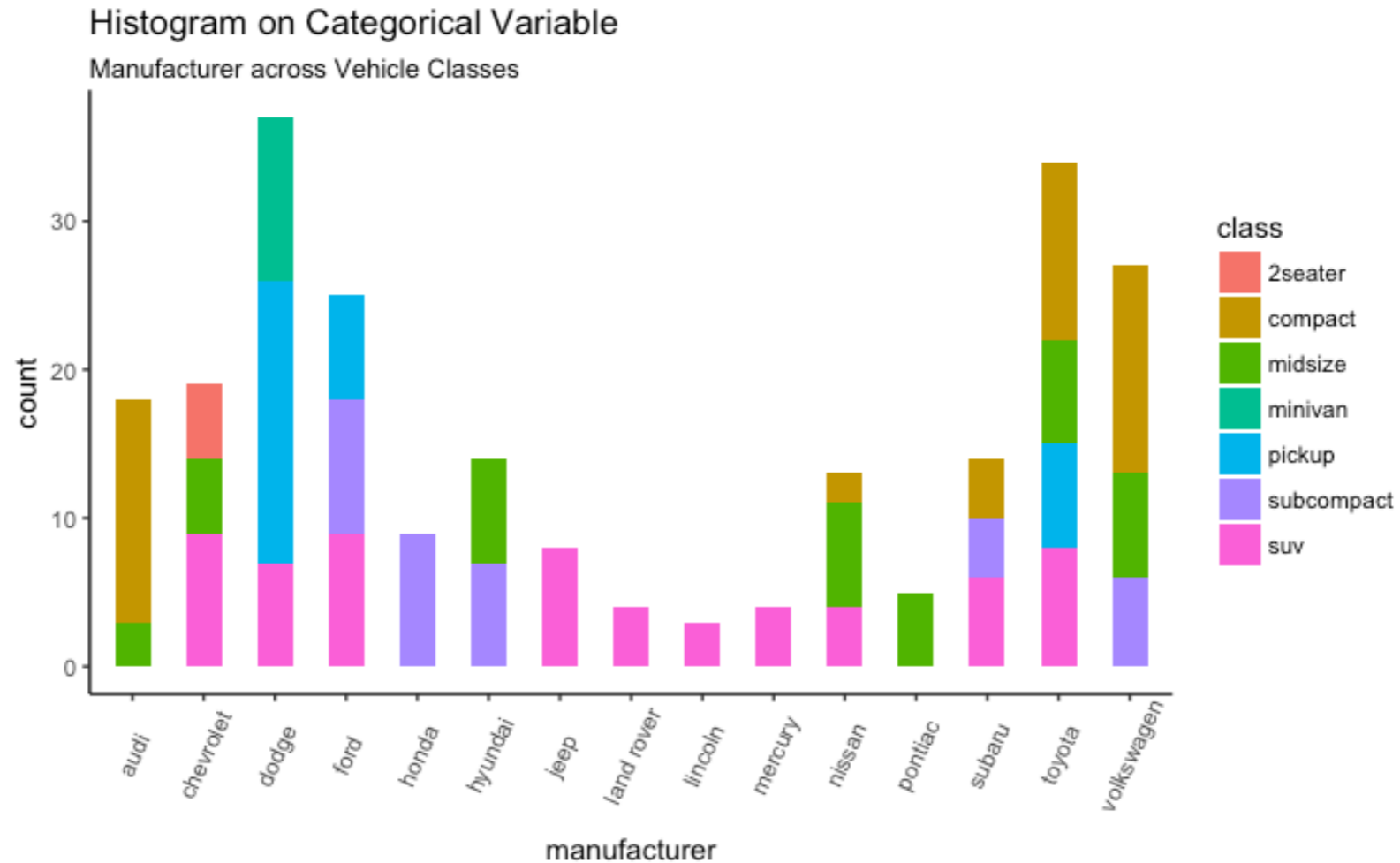
# Bargraphs

- Rotate



# Bargraphs

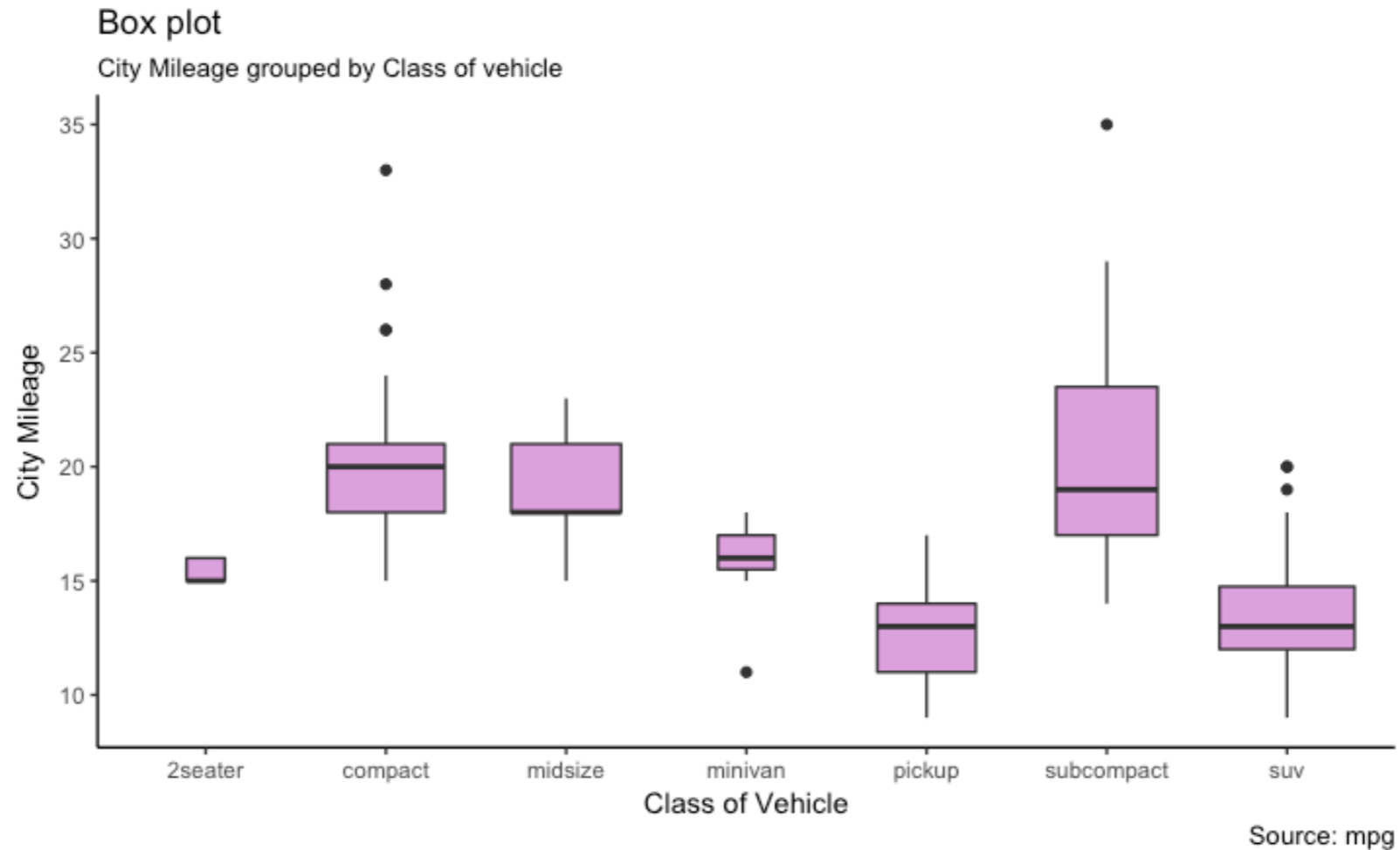
- Two variable categorical
- One variable numerical



# Boxplots

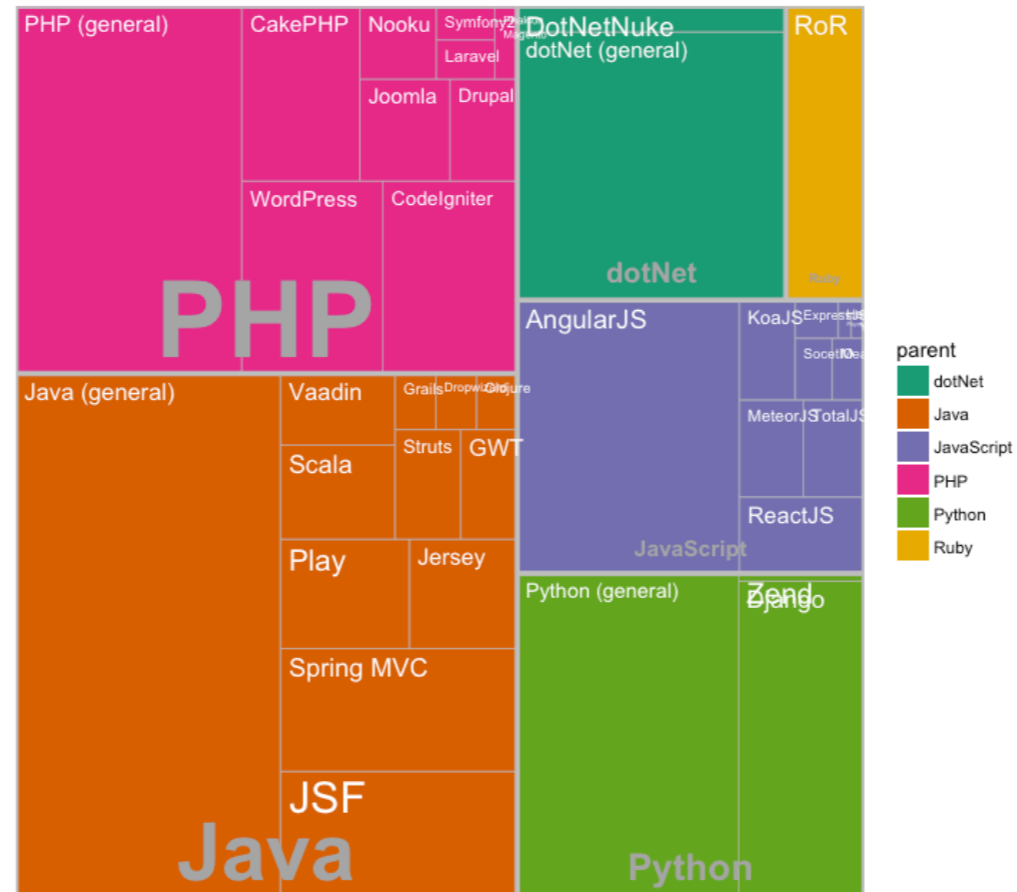
---

- One variable categorical
- One variable numerical but with many values



# Treemaps

- One variable categorical
- One variable numerical
- Much better than pie charts



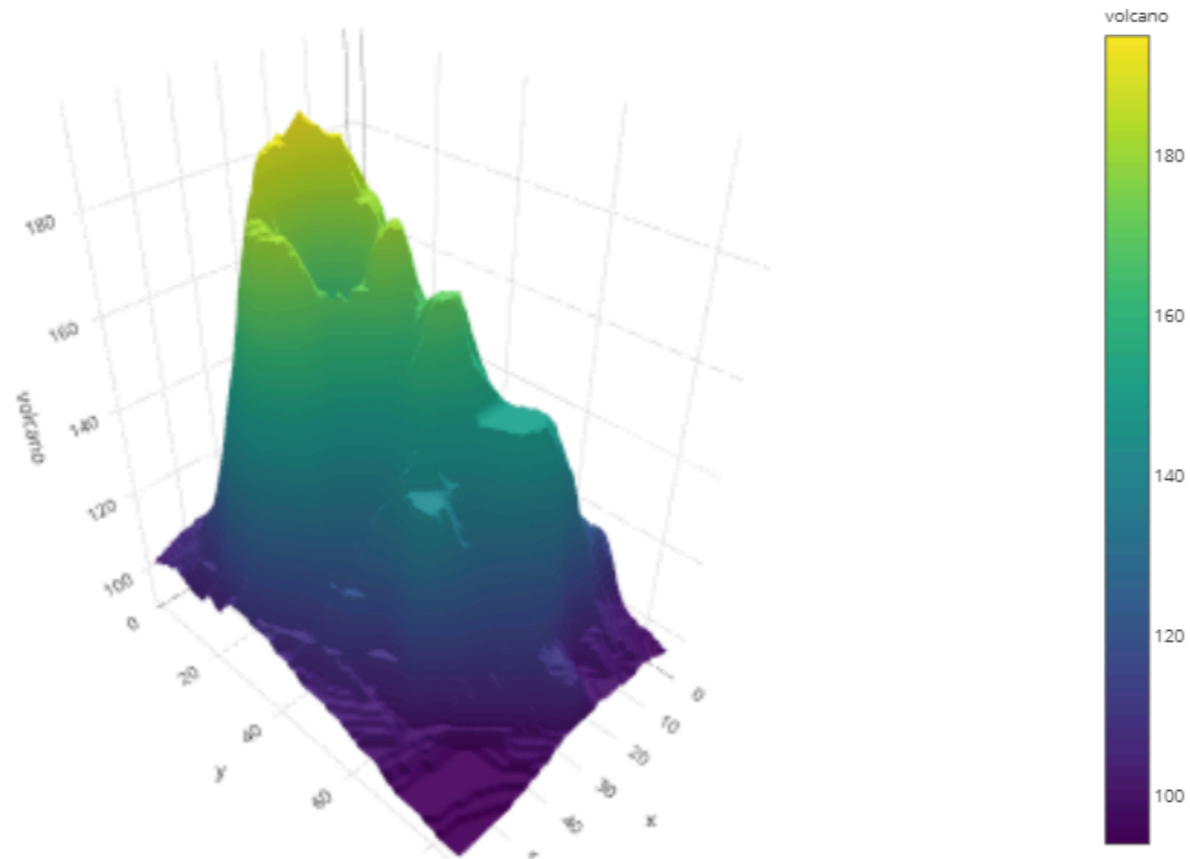
R - data visualization



# 3D

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- Three variable numerical
- Avoid unless it is a simple shape

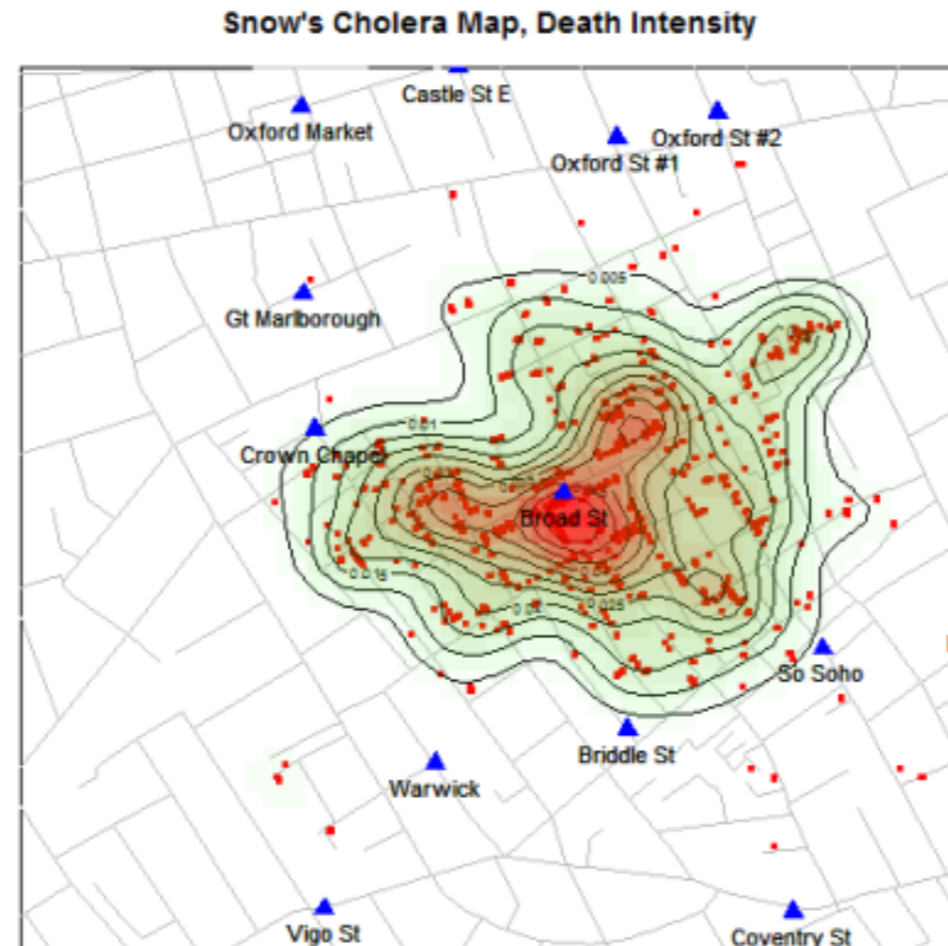


R - data visualization

# Contours

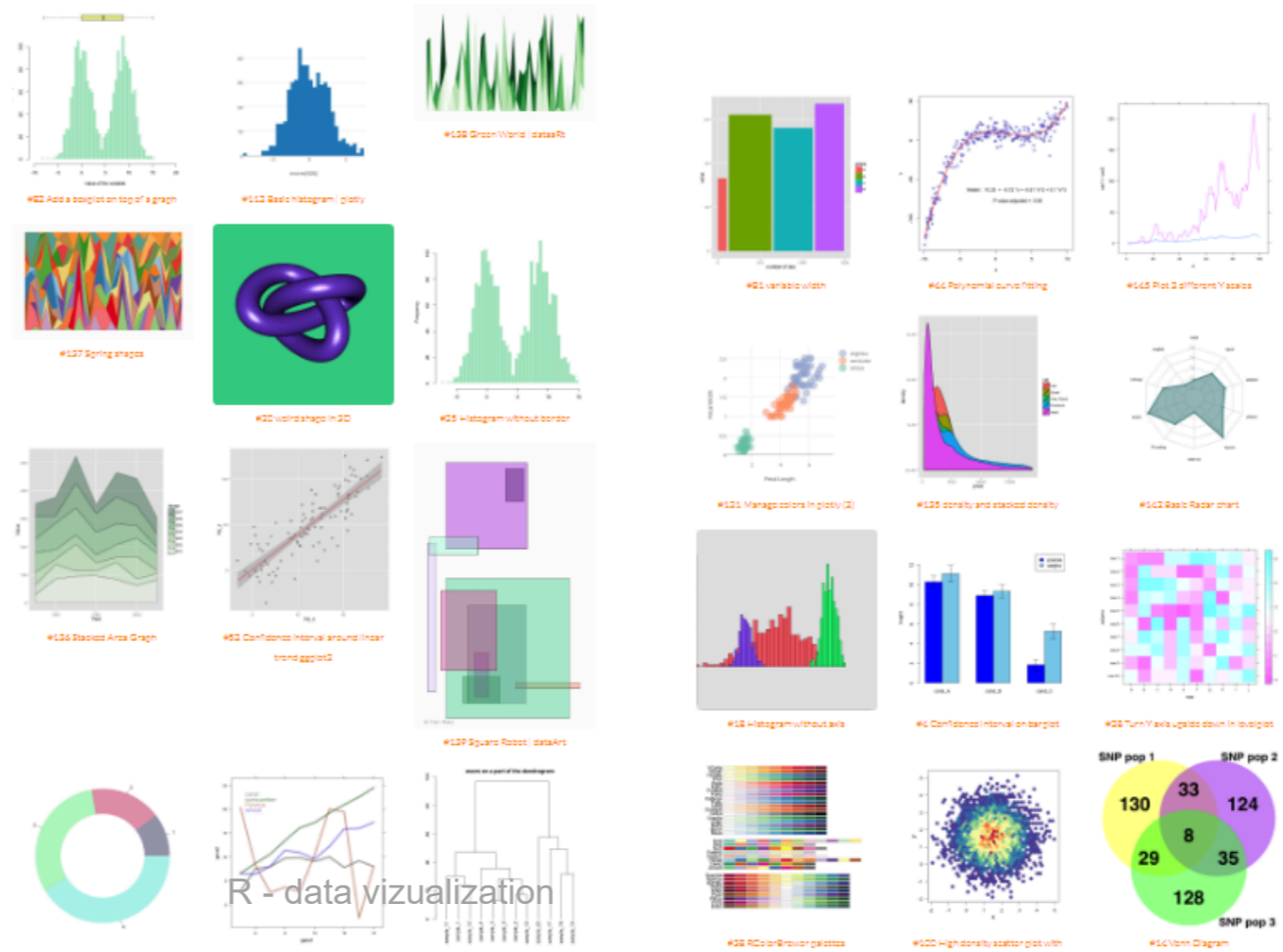
---

- Three variable numerical
- Better than 3D



# Many...

- Choose as a function of what you want to analyze or the story you want to tell
- <https://www.r-graph-gallery.com/all-graphs/>



R - data visualization

# ggplot2

# ggplot2: Build a data MASTERPIECE



# Initialize

## Load necessary libraries

```
library("readxl") # Import the data from Excel file
library("dplyr") # filter and reformat data frames
library("ggplot2") # graphics
library("patchwork") # arrange graphics
```

## Read the data

```
samples <- readxl::read_excel("data/CARBOM data.xlsx",
                             sheet = "Samples_boat") %>%
  tidyr::fill(station)
```

sample number	transect	station	date	time	depth	level	latitude	longitude	picoeuks	nanoeuks	phosphates	nitrates	temperature	salinity
10	1	81	2013-11-13	1899-12-31 01:00:00	140	Deep	-27.42	-44.72	3278	1232	0.20	0.26	17.3	35.9
11	1	85	2013-11-13	1899-12-31 13:30:00	110	Deep	-26.80	-45.30	16312	1615	0.29	0.22	21.3	36.5
120	2	96	2013-11-18	1899-12-31 23:50:00	5	Surf	-27.39	-47.82	1150	75	0.43	0.19	23.1	33.5
121	2	96	2013-11-18	1899-12-31 23:50:00	30	Deep	-27.39	-47.82	1737	218	0.43	0.23	22.6	33.7
122	2	96	2013-11-18	1899-12-31 23:50:00	50	Deep	-27.39	-47.82	853	234	0.56	0.21	20.3	35.9
125	2	98	2013-11-18	1899-12-31 05:00:00	5	Surf	-27.59	-47.39	3086	1300	0.29	0.25	23.1	35.7
126	2	98	2013-11-18	1899-12-31 05:00:00	50	Deep	-27.59	-47.39	1217	782	0.25	0.20	23.7	37.2
127	2	98	2013-11-18	1899-12-31 05:00:00	85	Deep	-27.59	-47.39	3420	226	0.25	0.47	22.9	37.0
13	1	86	2013-11-13	1899-12-31 17:00:00	105	Deep	-26.33	-45.41	6366	1007	0.34	0.15	20.9	36.3
140	2	101	2013-11-18	1899-12-31 12:00:00	5	Surf	-27.79	-46.96	500	366	0.29	0.14	23.5	36.5

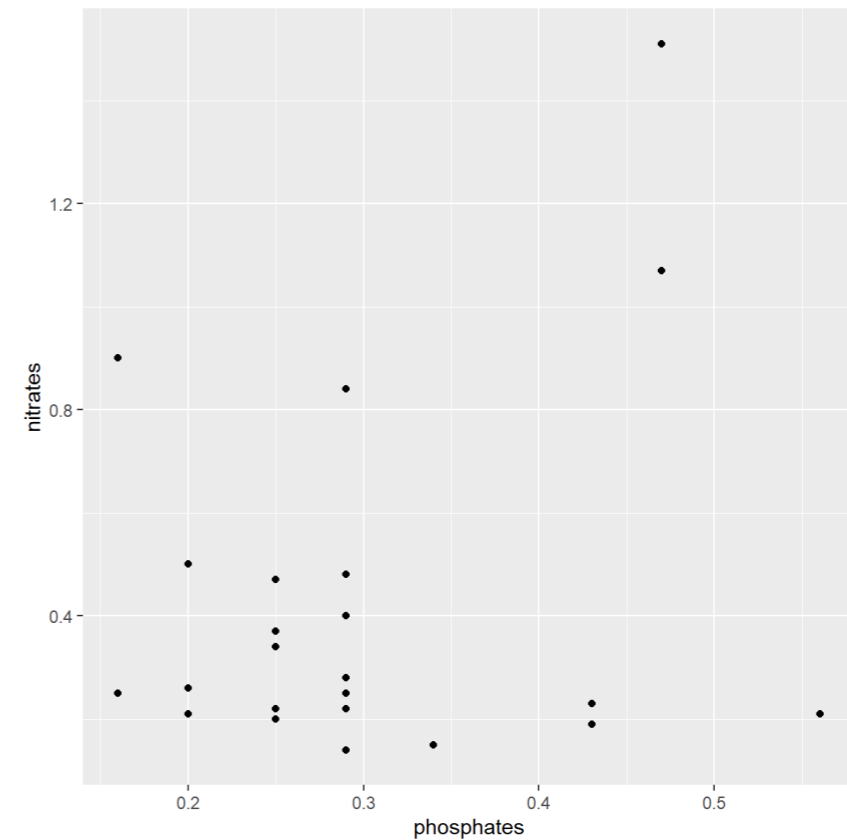
# A simple plot

---

- Choose the data set
- Choose the geometric representation
- Choose the **aesthetics** : x,y, color, shape etc...

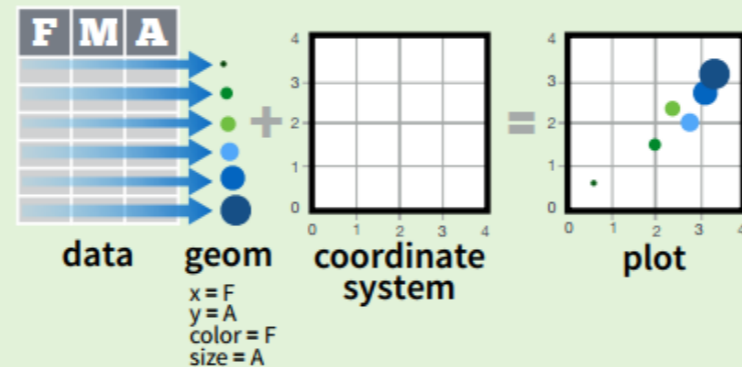
```
# All functions are from ggplot2
# package unless specified

ggplot(data=samples) +
  geom_point(mapping = aes(x=phosphates,
                           y=nitrates))
```



# The grammar of graphics

To display values, map variables in the data to visual properties of the geom (**aesthetics**) like **size**, **color**, and **x** and **y** locations.



```
ggplot(data=samples) +  
  geom_point(mapping = aes(x=phosphates,  
                           y=nitrates))
```

Every graph can be described as a combination of independent building blocks:

- **data**: a data frame: quantitative, categorical; local or data base query
- **aesthetic** mapping of variables into visual properties: size, color, x, y
- **geometric** objects (“geom”): points, lines, areas, arrows, ...
- **coordinate** system (“coord”): Cartesian, log, polar, map

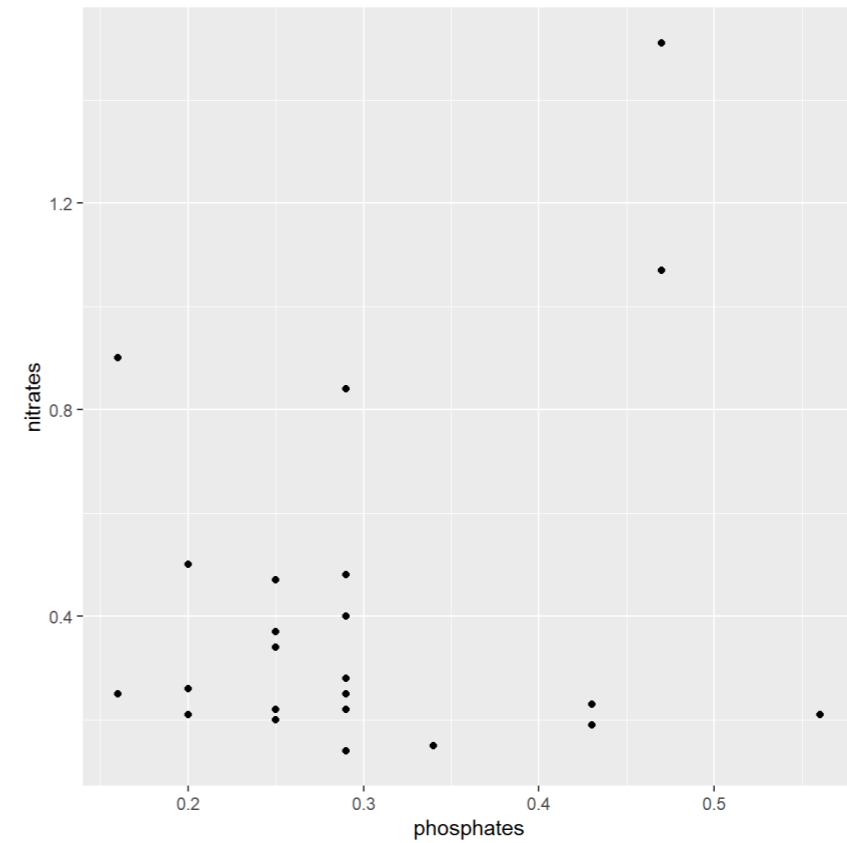


# Alternatively

---

- Move mapping into ggplot function

```
ggplot(data=samples,  
       mapping = aes(x=phosphates,  
                     y=nitrates)) +  
geom_point()
```

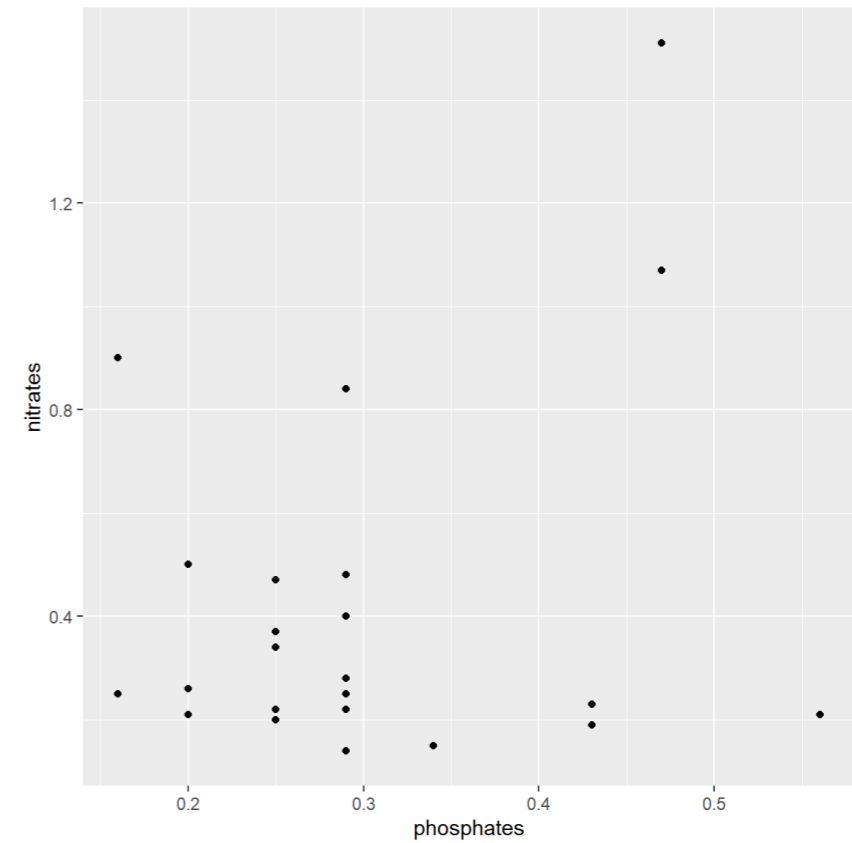


# Alternatively

---

- Remove function arguments

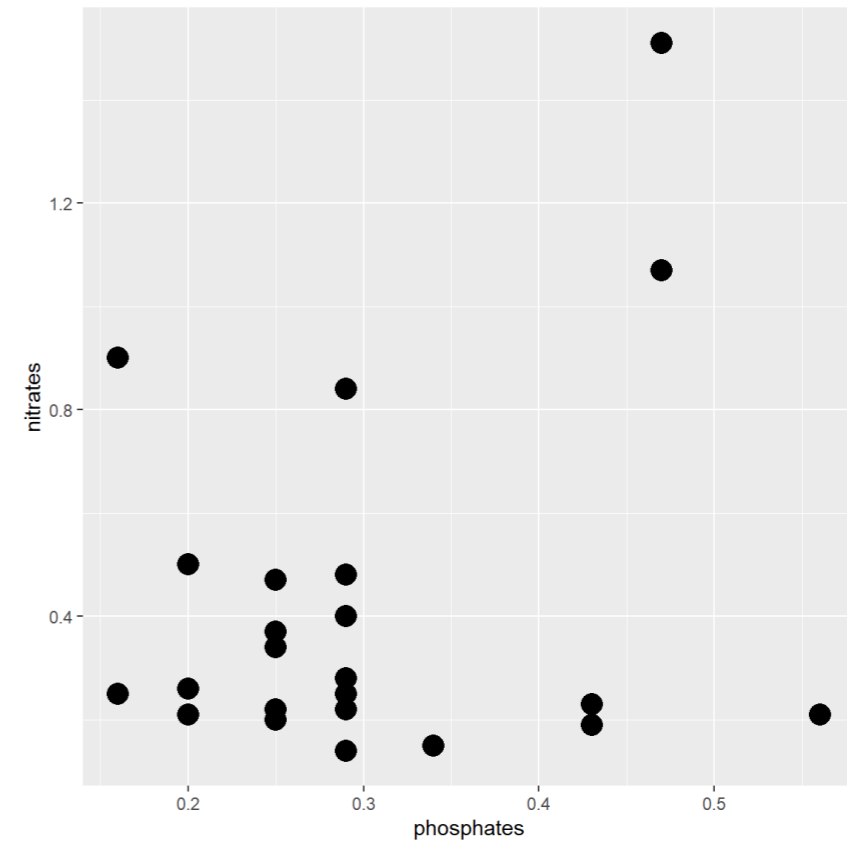
```
ggplot(samples,  
  aes(x=phosphates,  
      y=nitrates)) +  
  geom_point()
```



# Makes dots bigger

- Add: **size=5** outside of the aesthetics function

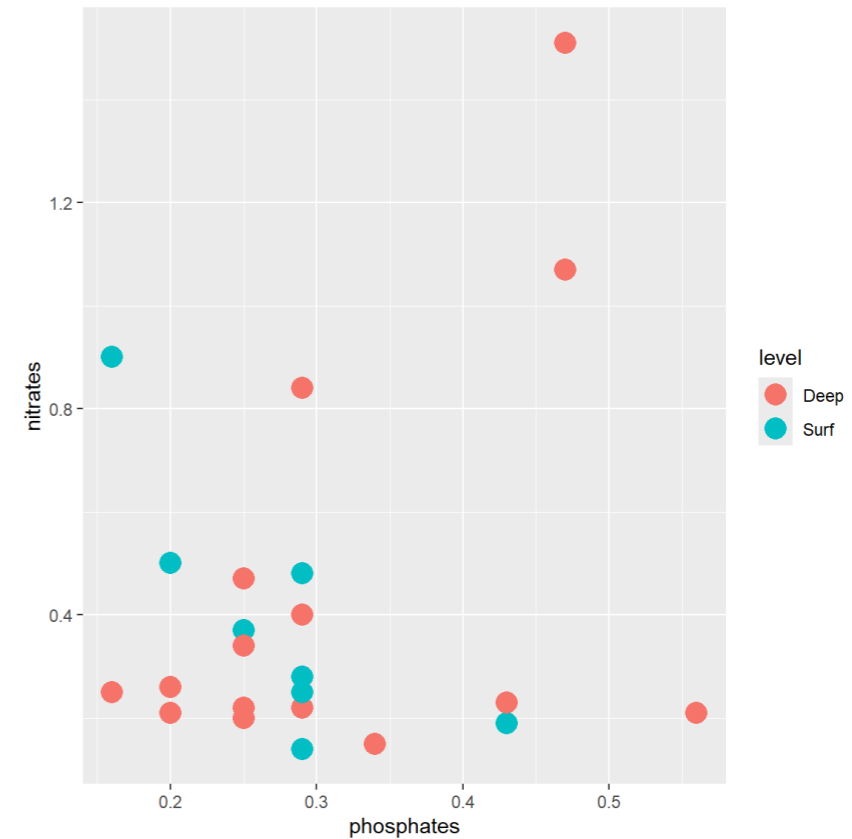
```
ggplot(samples,  
  aes(x=phosphates,  
      y=nitrates)) +  
  geom_point(size=5)
```



# Color according to depth level (discrete)

- The mapping aesthetics must be an argument of the aes function
- `geom_point(color=level, size=5)` will generate an error...

```
ggplot(samples,  
  aes(x=phosphates,  
      y=nitrates,  
      color=level)) +  
  geom_point(size=5)
```



# Color according to depth level (discrete)

---

- The mapping aesthetics must be an argument of the aes function
- `geom_point(color=level, size=5)` will generate an error...

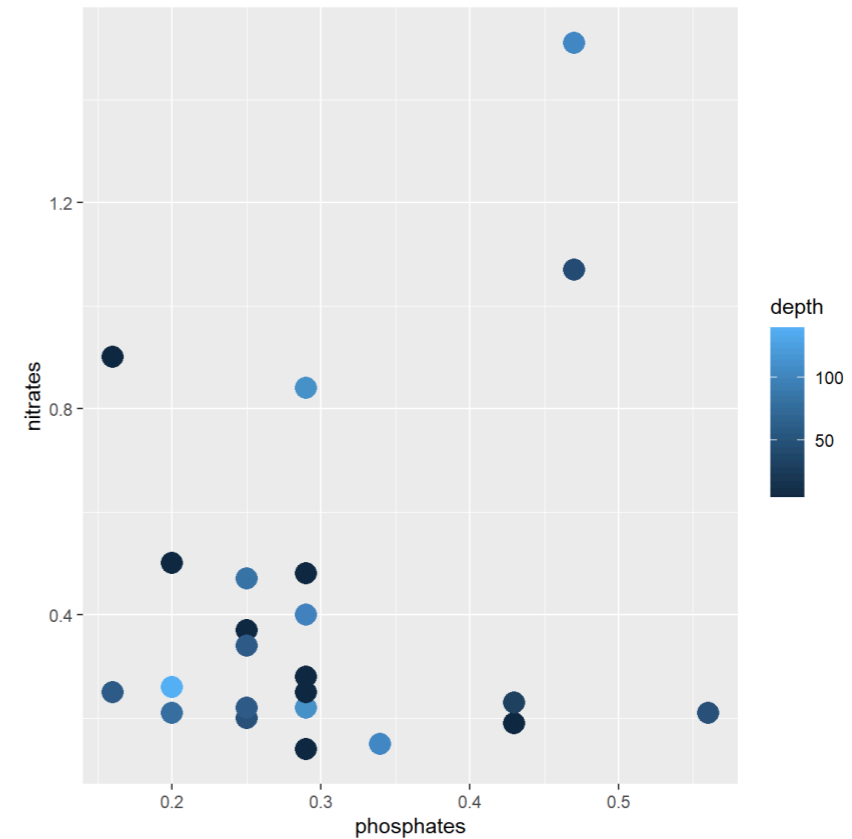
```
ggplot(samples,  
  aes(x=phosphates,  
      y=nitrates)) +  
  geom_point(color=level, size=5)
```

```
Error in eval(expr, envir, enclos): object 'level'  
not found
```

# Color according to depth (continuous)

- The mapping aesthetics must be an argument of the aes function
- Add: **color=depth**

```
ggplot(samples,  
  aes(x=phosphates,  
      y=nitrates,  
      color=depth)) +  
  geom_point(size=5)
```



# Symbol according to transect (continuous)

---

- Add: `shape=transect`

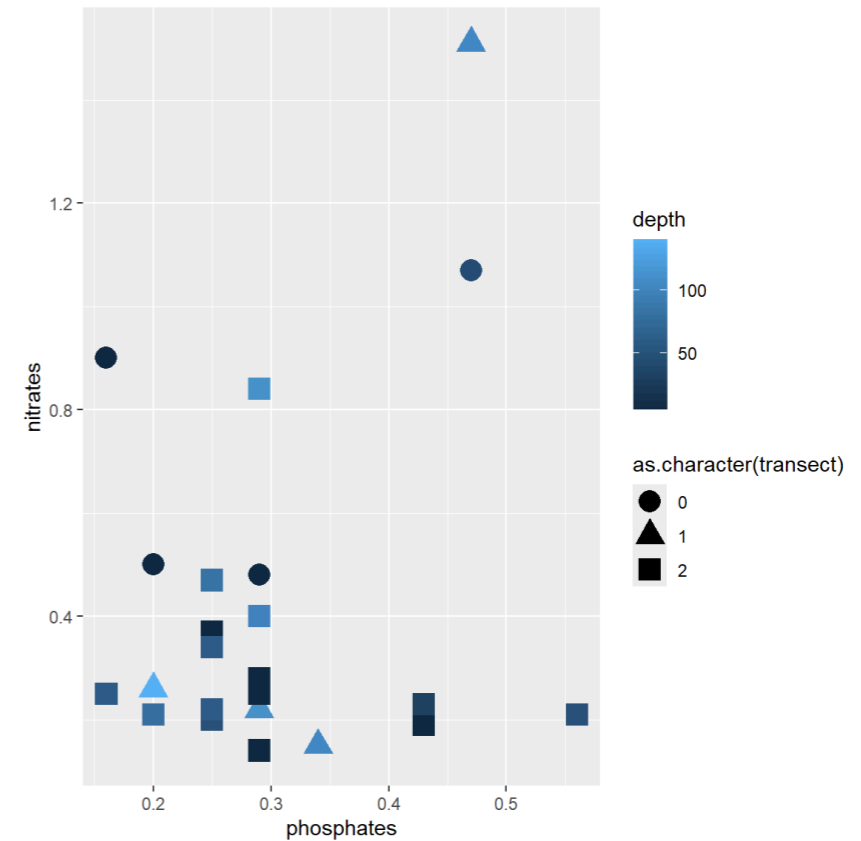
```
ggplot(samples,  
  aes(x=phosphates,  
      y=nitrates,  
      color=depth,  
      shape=transect)) +  
  geom_point(size=5)
```

```
Error in `geom_point()`:  
! Problem while computing aesthetics.  
i Error occurred in the 1st layer.  
Caused by error in `scale_f()`:  
! A continuous variable cannot be mapped to the  
shape aesthetic.  
i Choose a different aesthetic or use  
`scale_shape_binned()`.
```

# Symbol according to transect (continuous)

- Add: `shape=as.character(transect)`

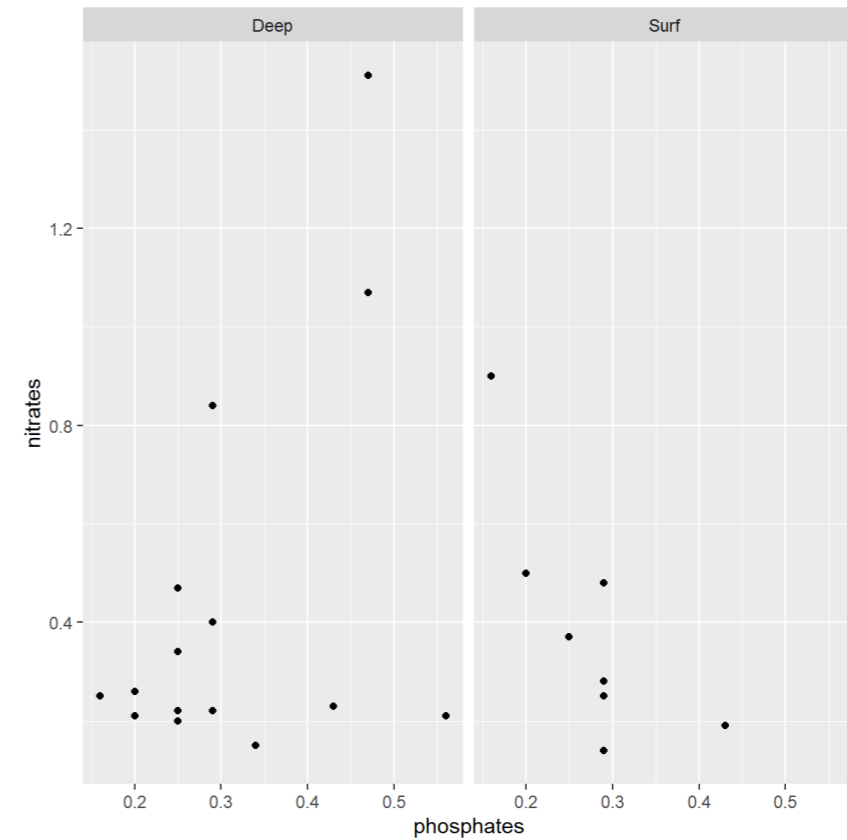
```
ggplot(samples,
  aes(x=phosphates,
      y=nitrates,
      color=depth,
      shape=as.character(transect))) +
  geom_point(size=5)
```





# Panels depending on one variable

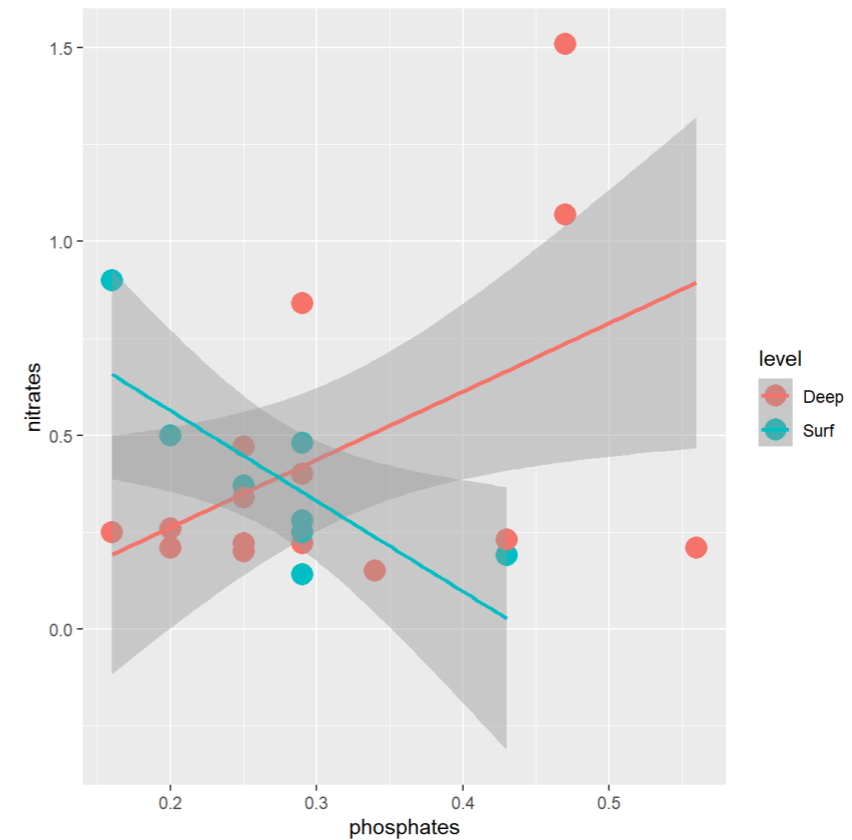
```
ggplot(samples,  
        aes(x=phosphates,  
            y=nitrates)) +  
  geom_point() +  
  facet_wrap(~ level)
```



# Adding a regression line

- Add: `geom_smooth()`
- You can choose the type of smoothing “lm” is for linear model

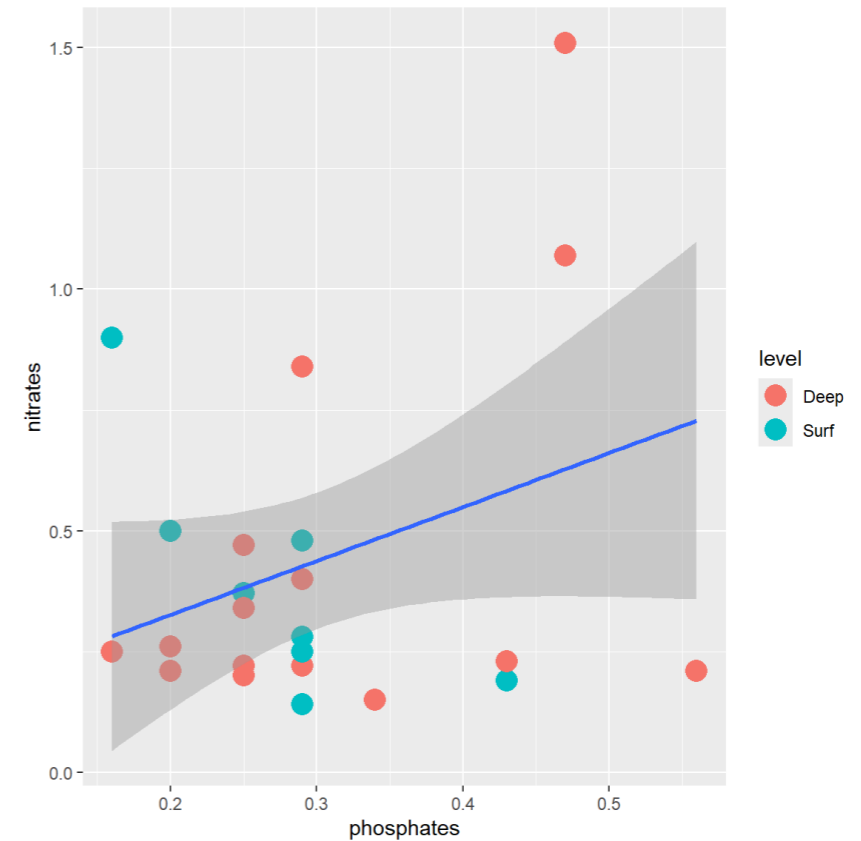
```
ggplot(samples,
  aes(x=phosphates,
      y=nitrates,
      color=level)) +
  geom_point(size=5) +
  geom_smooth(mapping = aes(x=phosphates,
                           y=nitrates),
             method="lm")
```



# Adding a regression line

- If the mapping is in the ggplot function is for all the geom....

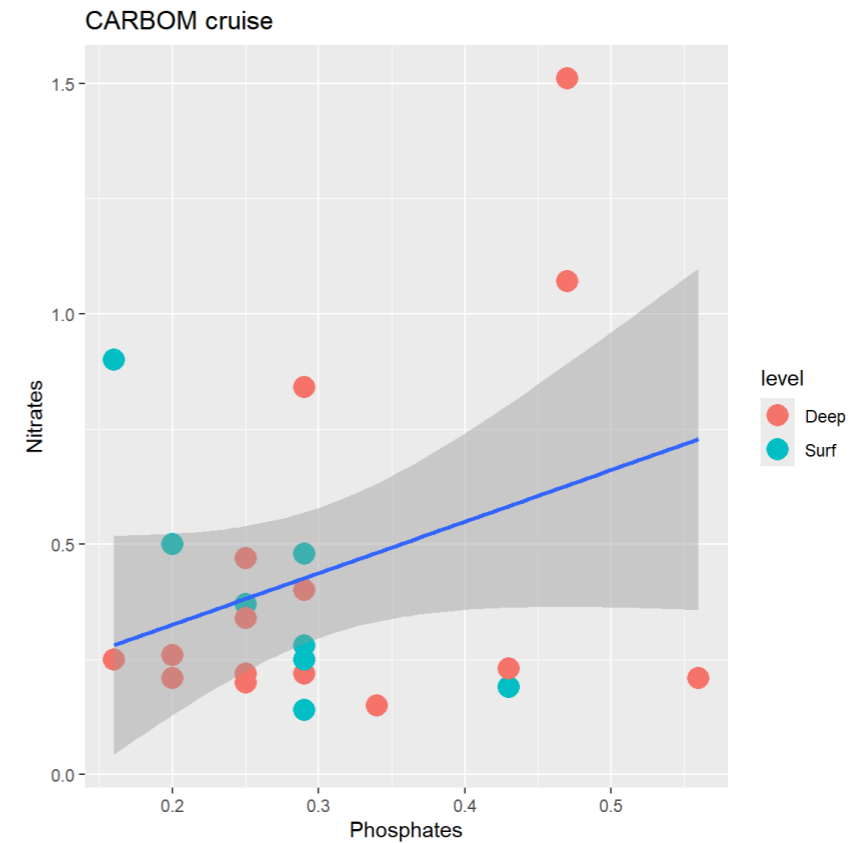
```
ggplot(samples,
  aes(x=phosphates,
      y=nitrates)) +
  geom_point(aes(color=level),
            size=5) +
  geom_smooth(mapping = aes(x=phosphates,
                           y=nitrates),
             method="lm")
```



# Finalizing the graph

- Adding labels and legends

```
ggplot(samples) +  
  geom_point(mapping = aes(x=phosphates,  
                           y=nitrates,  
                           color=level),  
            size=5) +  
  geom_smooth(mapping = aes(x=phosphates,  
                           y=nitrates),  
            method="lm") +  
  xlab("Phosphates") +  
  ylab("Nitrates") +  
  ggtitle("CARBOM cruise")
```

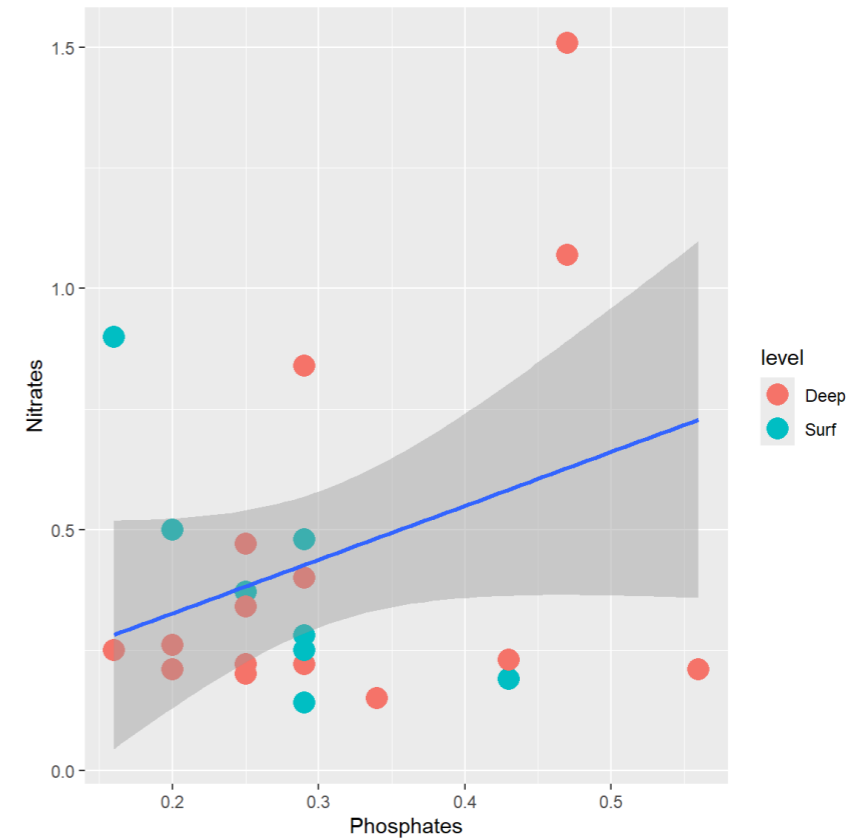


# Multigraphs (patchwork package)

# First graph

```
g1 <- ggplot(samples) +  
  geom_point(mapping = aes(x=phosphates,  
                           y=nitrates,color=  
                           level), size=5)  
  geom_smooth(mapping = aes(x=phosphates,  
                           y=nitrates),  
             method="lm") +  
  xlab("Phosphates") +  
  ylab("Nitrates")
```

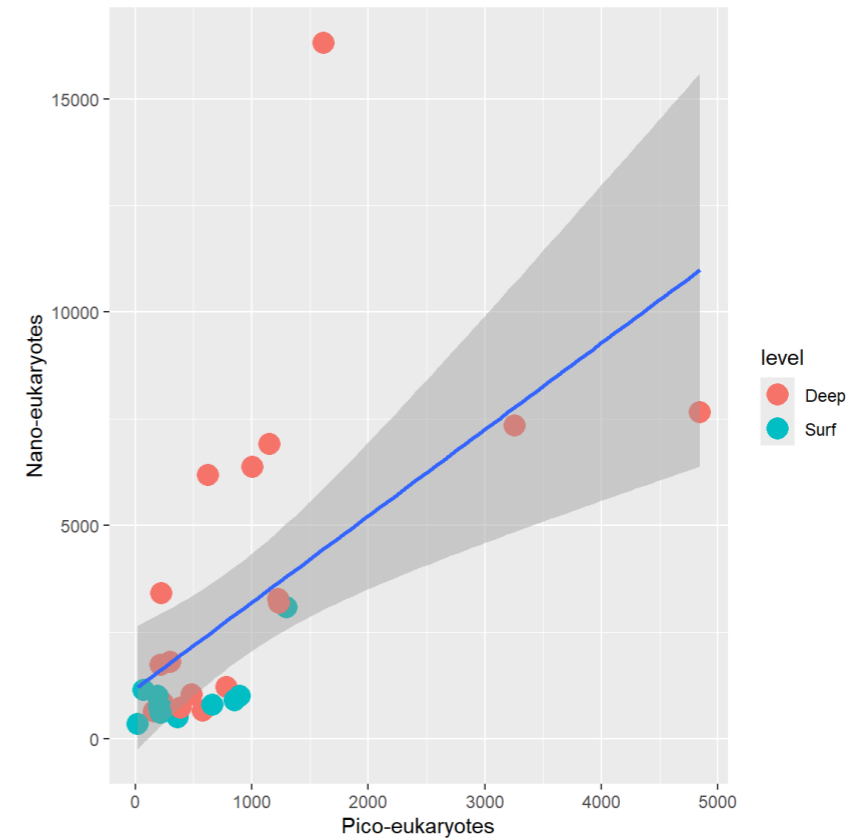
g1



# Second graph

```
g2<- ggplot(samples) +  
  geom_point(mapping = aes(x=nanoeuks,  
                           y=picoeuks,  
                           color=level),  
            size=5) +  
  geom_smooth(mapping = aes(nanoeuks,  
                           y=picoeuks),  
            method="lm") +  
  xlab("Pico-eukaryotes") +  
  ylab("Nano-eukaryotes")
```

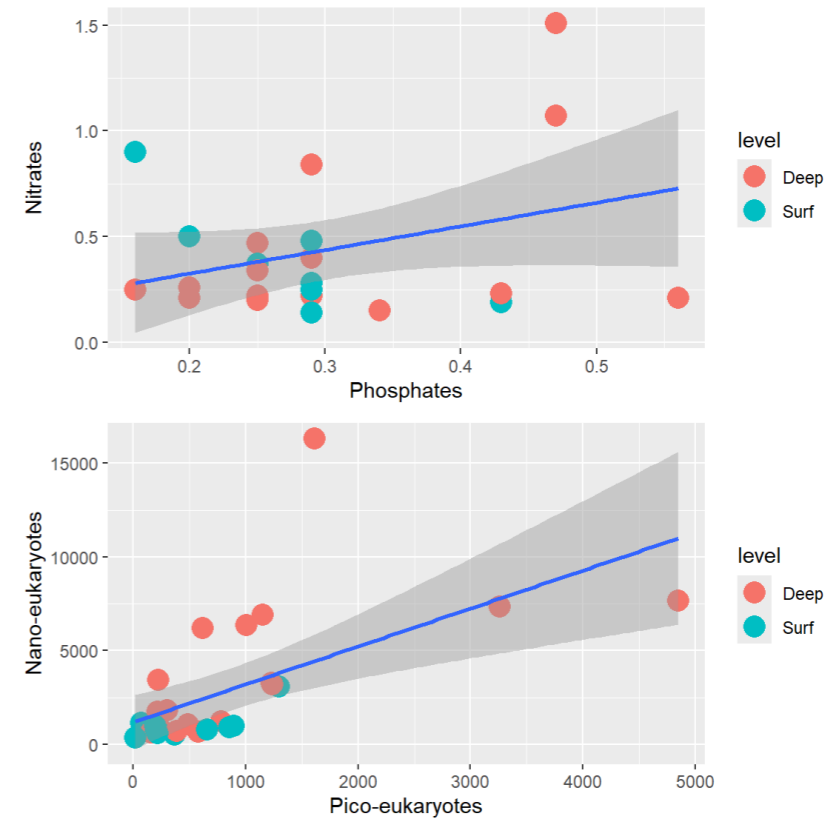
g2



# Package patchwork

- Other packages:
  - gridExtra
  - cowplot

```
library(patchwork)
(g1 / g2)
```

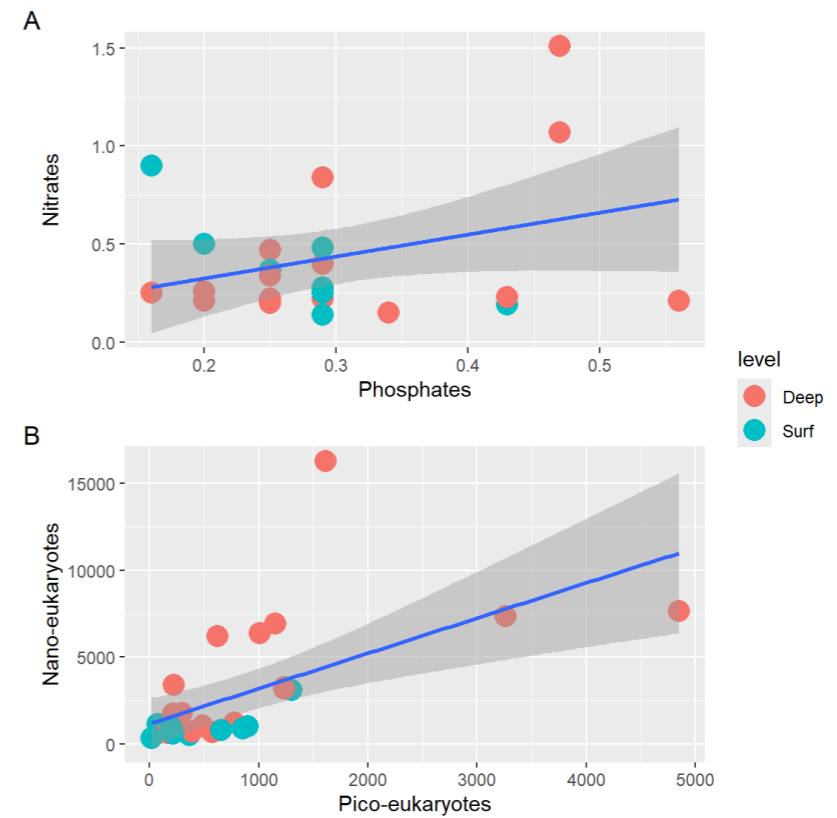




# Package patchwork

- Adding annotation
- Collecting legends

```
g1 / g2 +  
  plot_annotation(tag_levels = 'A') +  
  plot_layout(guides = 'collect')
```



# Package esquisse

esquisse 2.0.1.9000 Reference Articles Changelog

Search for



## Get started with esquisse

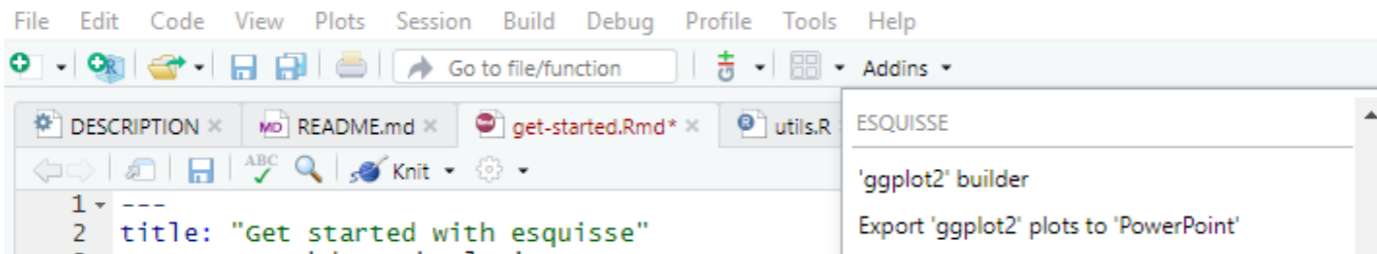


Source: [vignettes/get-started.Rmd](#)

```
library(esquisse)
```

## Launch the addin

In RStudio, you can use the *Addins* menu :



Or in the R console :

```
esquisser()
```

To use a `data.frame` by default, if using the *Addins* menu highlight with the cursor a

### On this page

Launch the addin

Import data into  
{esquisse}

Create a plot

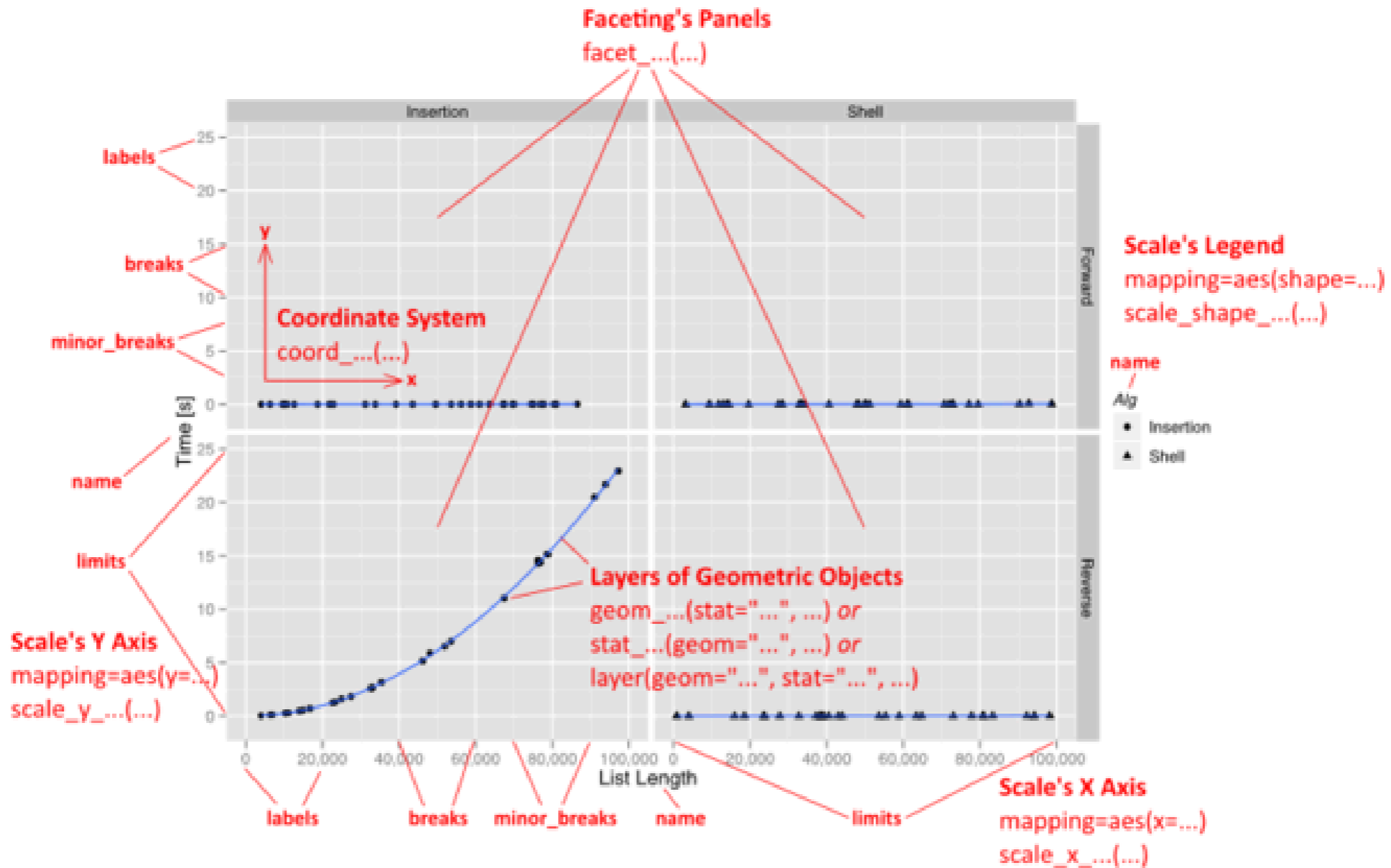
Controls

Export

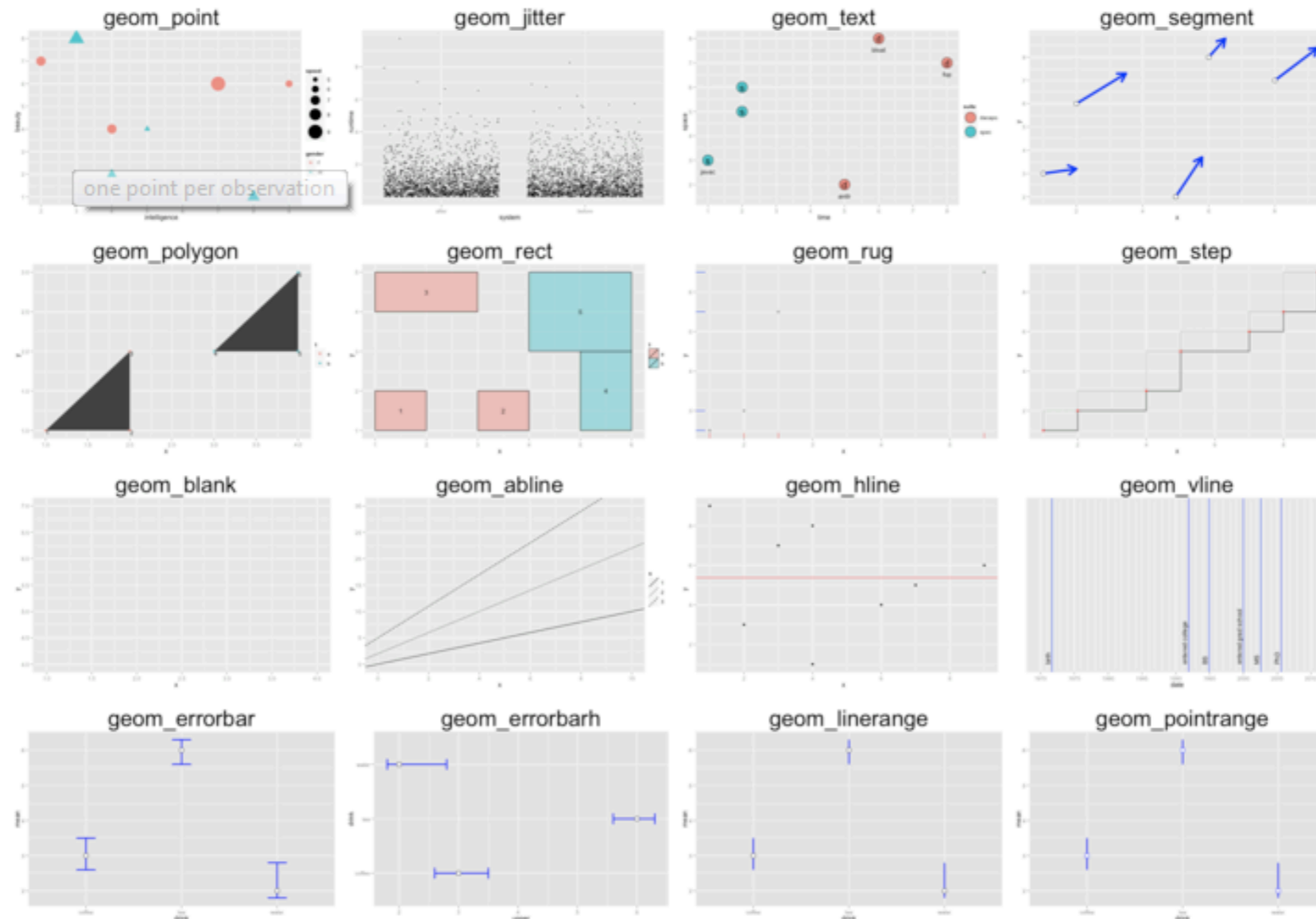
Addin options

# ggplot2 syntax

# Anatomy of a plot



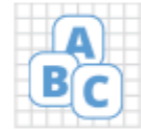
# Geometries



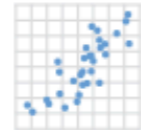
# Continuous x and y

## continuous x , continuous y

```
e <- ggplot(mpg, aes(cty, hwy))
```



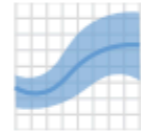
**e + geom\_label**(aes(label = cty), nudge\_x = 1, nudge\_y = 1, check\_overlap = TRUE) x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust



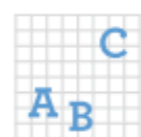
**e + geom\_jitter**(height = 2, width = 2) x, y, alpha, color, fill, shape, size



**e + geom\_point**(), x, y, alpha, color, fill, shape, size, stroke



**e + geom\_smooth**(method = lm), x, y, alpha, color, fill, group, linetype, size, weight



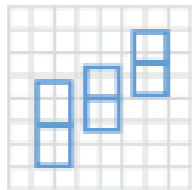
**e + geom\_text**(aes(label = cty), nudge\_x = 1, nudge\_y = 1, check\_overlap = TRUE), x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust

# Plotting error

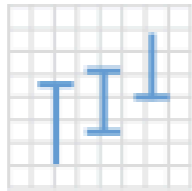
---

## visualizing error

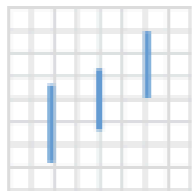
```
df <- data.frame(grp = c("A", "B"), fit = 4:5, se = 1:2)
j <- ggplot(df, aes(grp, fit, ymin = fit-se, ymax = fit+se))
```



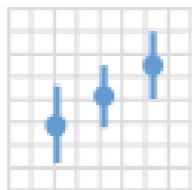
**j + geom\_crossbar(fatten = 2)**  
x, y, ymax, ymin, alpha, color, fill, group, linetype, size



**j + geom\_errorbar()**, x, ymax, ymin, alpha, color, group, linetype, size, width (also **geom\_errorbarh()**)



**j + geom\_linerange()**  
x, ymin, ymax, alpha, color, group, linetype, size



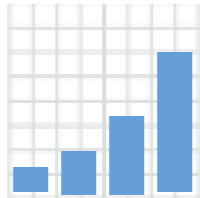
**j + geom\_pointrange()**  
x, y, ymin, ymax, alpha, color, fill, group, linetype, shape, size

# Discrete x - Continuous y

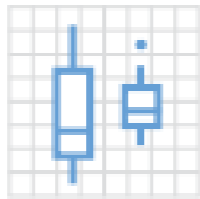
---

## discrete x , continuous y

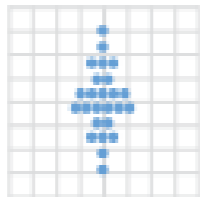
```
f <- ggplot(mpg, aes(class, hwy))
```



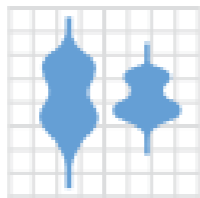
**f + geom\_col()**, x, y, alpha, color, fill, group, linetype, size



**f + geom\_boxplot()**, x, y, lower, middle, upper, ymax, ymin, alpha, color, fill, group, linetype, shape, size, weight



**f + geom\_dotplot(binaxis = "y", stackdir = "center")**, x, y, alpha, color, fill, group



**f + geom\_violin(scale = "area")**, x, y, alpha, color, fill, group, linetype, size, weight



# Continuous x

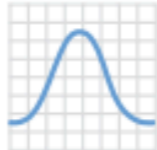
---

## ONE VARIABLE continuous

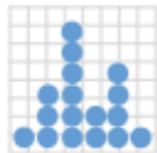
```
c <- ggplot(mpg, aes(hwy)); c2 <- ggplot(mpg)
```



```
c + geom_density(kernel = "gaussian")
```



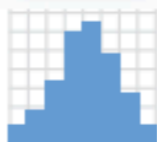
```
c + geom_density(kernel = "gaussian")  
x, y, alpha, color, fill, group, linetype, size, weight
```



```
c + geom_dotplot()  
x, y, alpha, color, fill
```



```
c + geom_histogram(binwidth = 5)
```



```
c + geom_histogram(binwidth = 5) x, y, alpha,  
color, fill, linetype, size, weight
```



```
c + geom_line()
```

# 3D

## THREE VARIABLES

```
seals$z <- with(seals, sqrt(delta_long^2 + delta_lat^2)) l <- ggplot(seals, aes(long, lat))
```



**l + geom\_contour(aes(z = z))**

x, y, z, alpha, colour, group, linetype,  
size, weight



**l + geom\_raster(aes(fill = z), hjust=0.5, vjust=0.5,  
interpolate=FALSE)**  
x, y, alpha, fill



**l + geom\_tile(aes(fill = z)), x, y, alpha, color, fill,  
linetype, size, width**

# Modifying axis and scales

## GENERAL PURPOSE SCALES

Use with most aesthetics

**scale\_\*\_continuous()** - map cont' values to visual ones

**scale\_\*\_discrete()** - map discrete values to visual ones

**scale\_\*\_identity()** - use data values as visual ones

**scale\_\*\_manual(values = c())** - map discrete values to manually chosen visual ones

**scale\_\*\_date(date\_labels = "%m/%d", date\_breaks = "2 weeks")** - treat data values as dates.

**scale\_\*\_datetime()** - treat data x values as date times. Use same arguments as `scale_x_date()`. See `?strptime` for label formats.

## X & Y LOCATION SCALES

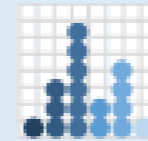
Use with x or y aesthetics (x shown here)

**scale\_x\_log10()** - Plot x on log10 scale

**scale\_x\_reverse()** - Reverse direction of x axis

**scale\_x\_sqrt()** - Plot x on square root scale

## COLOR AND FILL SCALES (CONTINUOUS)



**o <- c + geom\_dotplot(aes(fill = ..x..))**

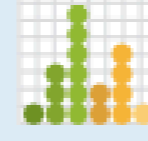
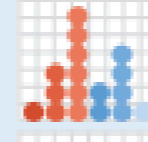
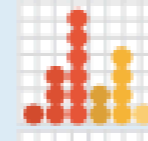
**o + scale\_fill\_distiller(palette = "Blues")**

**o + scale\_fill\_gradient(low="red", high="yellow")**

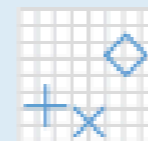
**o + scale\_fill\_gradient2(low="red", high="blue", mid = "white", midpoint = 25)**

**o + scale\_fill\_gradientn(colours=topo.colors(6))**

Also: `rainbow()`, `heat.colors()`, `terrain.colors()`, `cm.colors()`, `RColorBrewer::brewer.pal()`



## SHAPE AND SIZE SCALES



**p <- e + geom\_point(aes(shape = fl, size = cyl))**

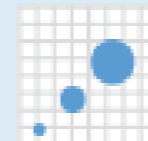
**p + scale\_shape() + scale\_size()**

**p + scale\_shape\_manual(values = c(3:7))**

0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25  
□ ○ △ + × ◇ ▽ ⊠ \* ⊕ ⊗ ⊘ ⊙ ⊚ ⊛ ⊜ ⊝ ⊞ ⊟ ⊠ ⊡ ⊢ ⊣ ⊤ ⊥ ⊦ ⊧ ⊨ ⊩ ⊪ ⊫ ⊬ ⊭ ⊮ ⊯ ⊰ ⊱ ⊲ ⊳ ⊴ ⊵ ⊶ ⊷ ⊸ ⊹ ⊺ ⊻ ⊼ ⊽ ⊾ ⊿ ⊿

**p + scale\_radius(range = c(1,6))**

**p + scale\_size\_area(max\_size = 6)**



# Palettes

- Package `tmaptools`
  - Function : `palette_explorer()`
- Package `paletteer`
  - More than 1000 palettes

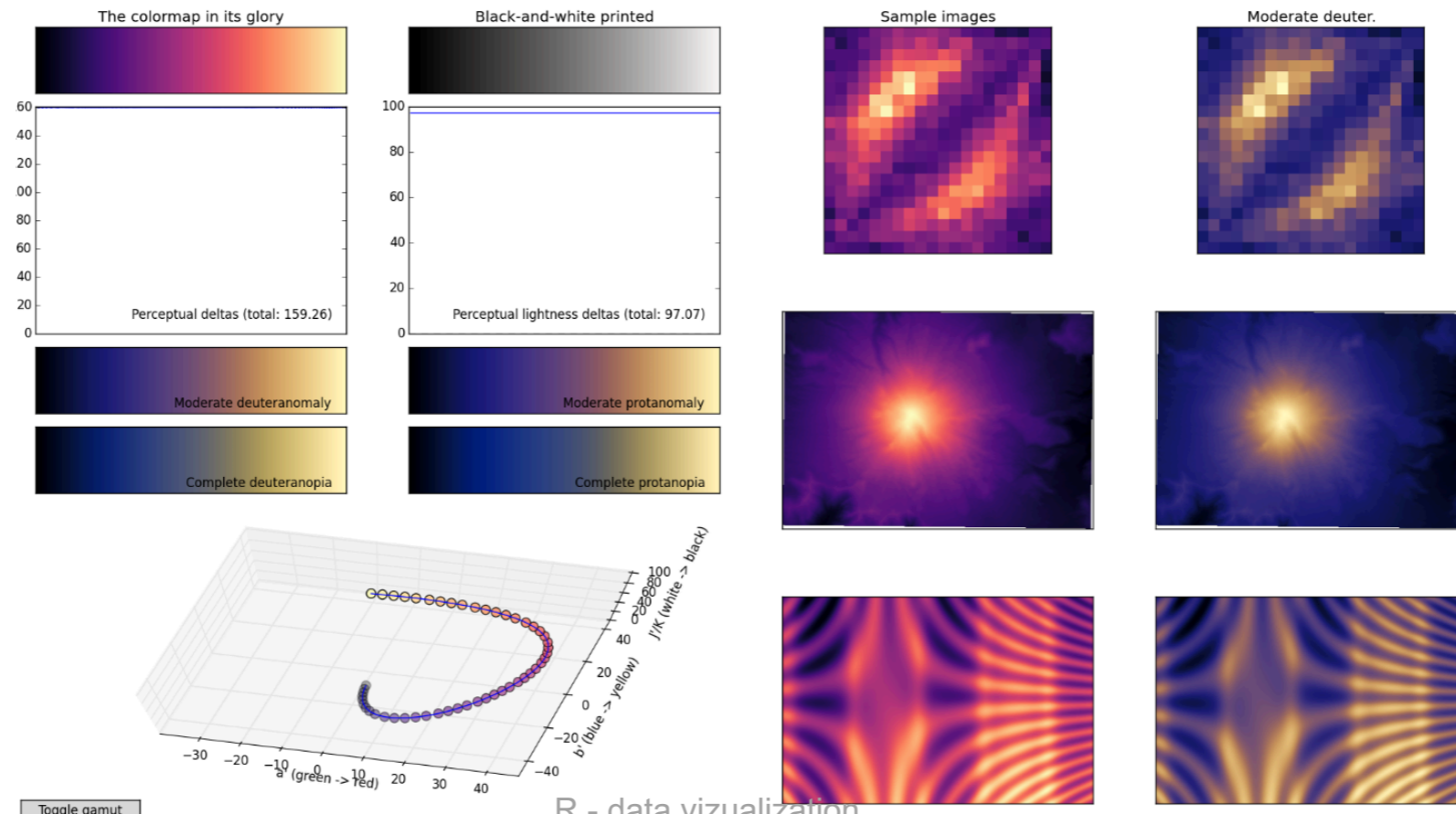
The screenshot displays the `palette_explorer()` interface, which is organized into several sections:

- Brewer Palettes:** Four sliders allow selection of the number of colors (3, 7, 9, or 20) for different Brewer palette types.
- Sequential Palettes:** A slider for contrast range (0 to 1) and a checkbox for "Automatic" selection.
- Categorical Palettes:** A checkbox for "Stretch" to adjust categorical color distributions.
- Diverging Palettes:** A slider for contrast range (0 to 1) and a checked checkbox for "Automatic" selection.
- Viridis Palettes:** A slider for the number of colors (3 to 20).
- Options:** A section at the bottom with checkboxes for "Print color values", "Code generator" (with radio buttons for "Direct code" and "tmap layer function call"), and "Color blindness simulator" (with radio buttons for "Normal" and "Deuteranopia").

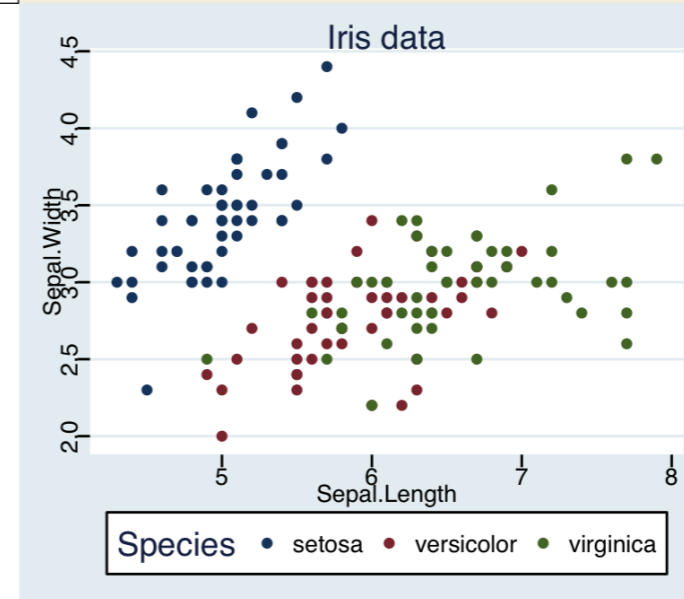
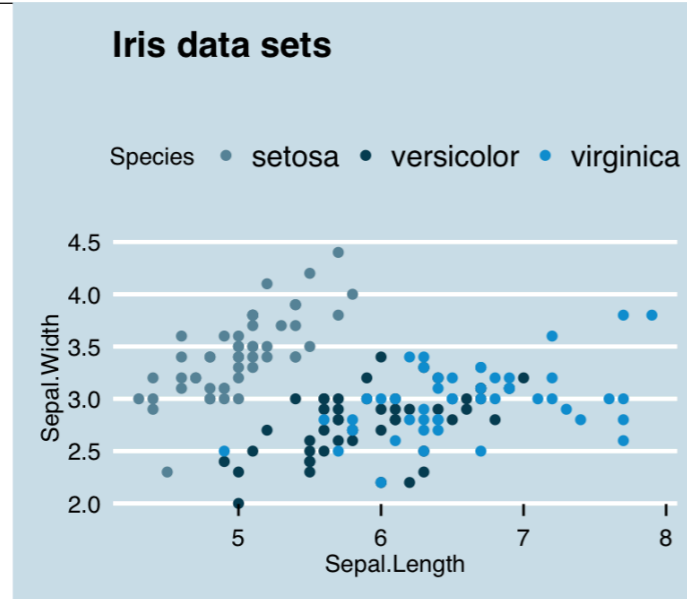
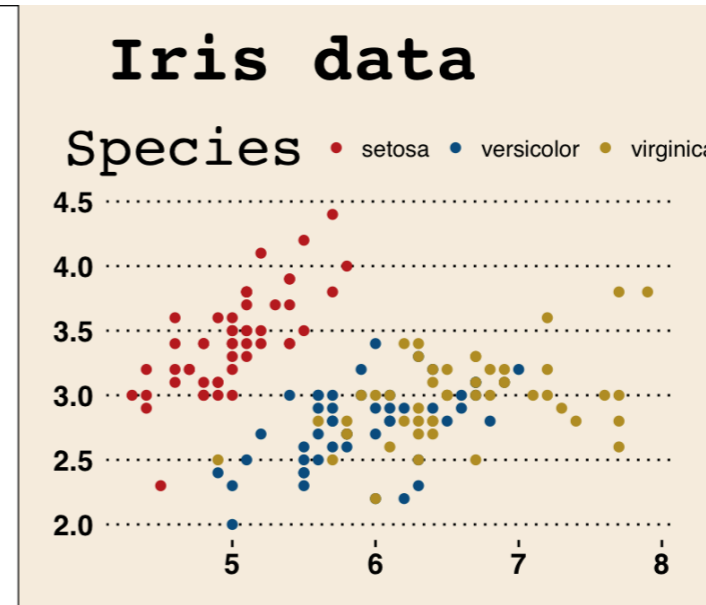
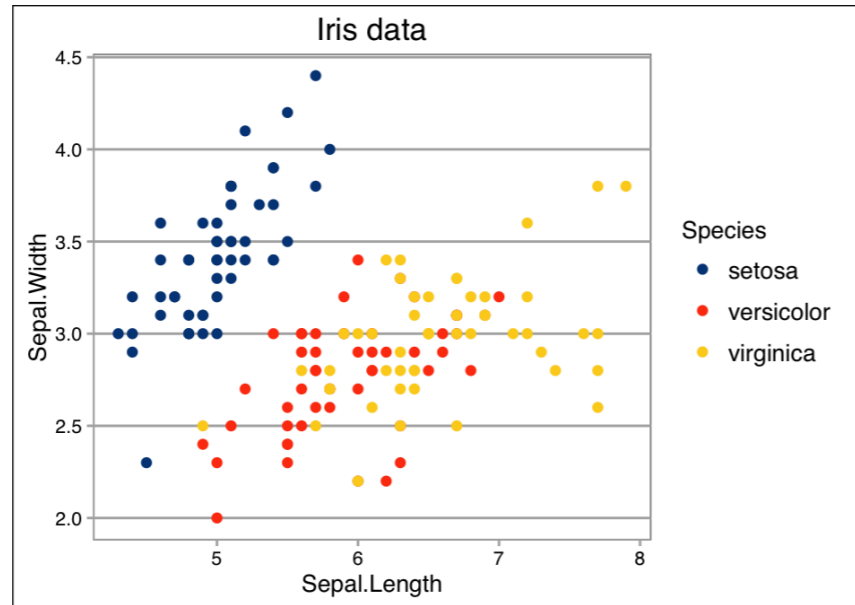
On the right side, a grid of color swatches is shown for various palettes, including Blues, BuGn, BuPu, GnBu, Greens, Greys, Oranges, OrRd, PuBu, PuBuGn, PuRd, Purples, RdPu, Reds, YlGn, YlGnBu, YlOrBr, and YlOrRd. Below the grid, R code snippets are provided for each palette, such as `get_brewer_pal("Blues", n = 7, contrast = c(0.25, 0.9))` and `viridisLite::viridis(20)`.

# Palettes

- Use color blind friendly palettes
  - viridis (e.g. `scale_colour_viridis_c()`)



# Themes



# Extensions

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## ggplot2 extensions - gallery

- [Add Your Extension!](#)
- [exts.ggplot2.tidyverse.org](https://exts.ggplot2.tidyverse.org)
- [Navbar Link](#)



142 registered extensions available to explore

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- Name
- Author
- Github stars

Github stars ▼

Sort

search name, author, descrip Text Filter

▼

- 
- Alex Zanidean (1)
- AllanCameron (1)
- almeidaxan (1)
- aphalo (4)
- arcresu (1)

**Let's do a graph**



# Your mission

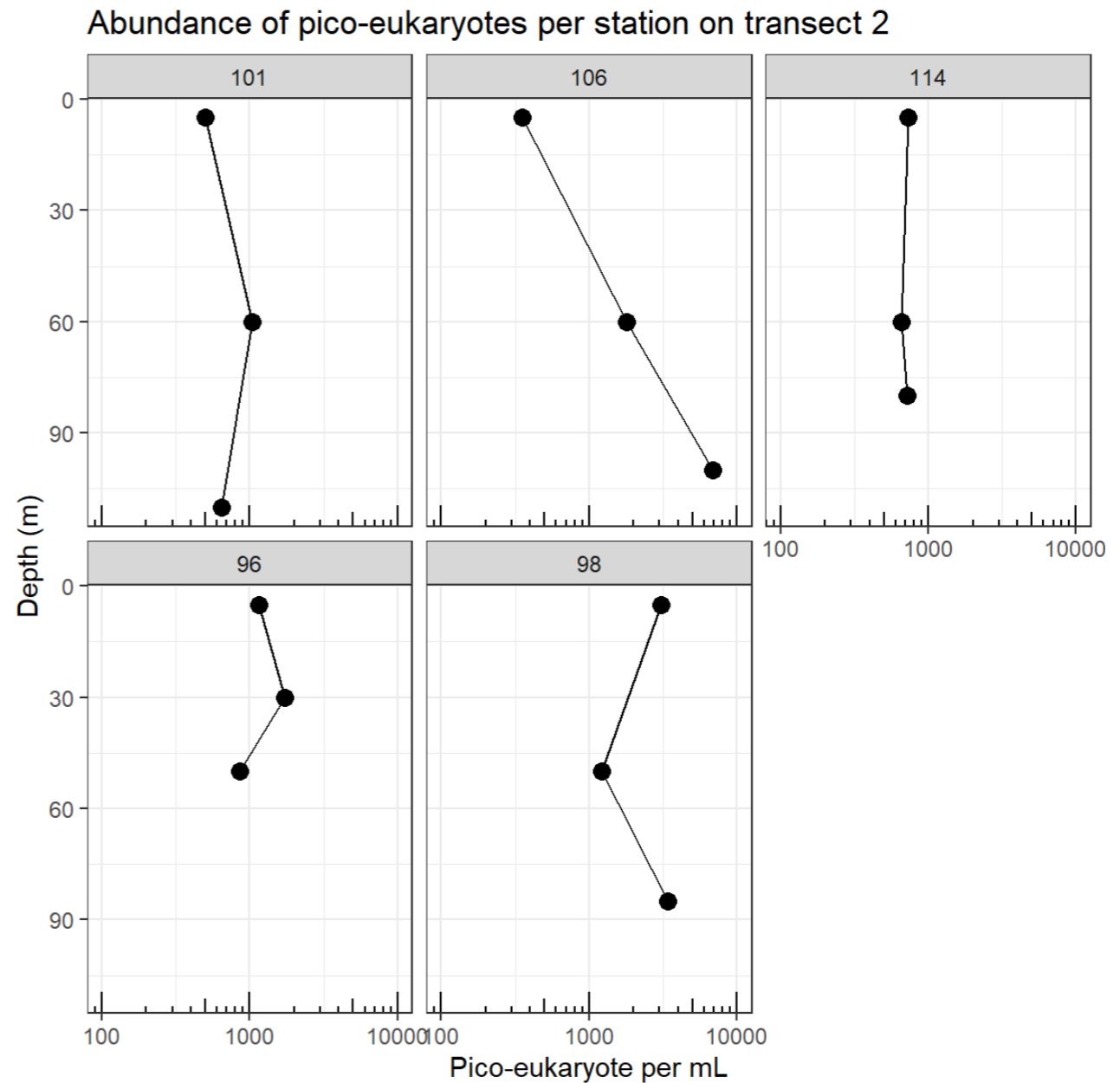
---

# Reproduce graph on right

- Only transect 2
- One panel per station
- Increasing depth
- Log scale for x
- White background

## Instructions

- Work by group of 2 (1 expert, 1 less expert)
- We will correct later in the week



# Recap

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- Conceptualize your graph before coding
- Decide what element is fixed and what varies
- It takes time to get what you want...
- Exploratory vs. final

**Next time: Markdown and Quarto**