

# R COURSE

Data visualization

Daniel Vaulot

2025-01-16



# 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

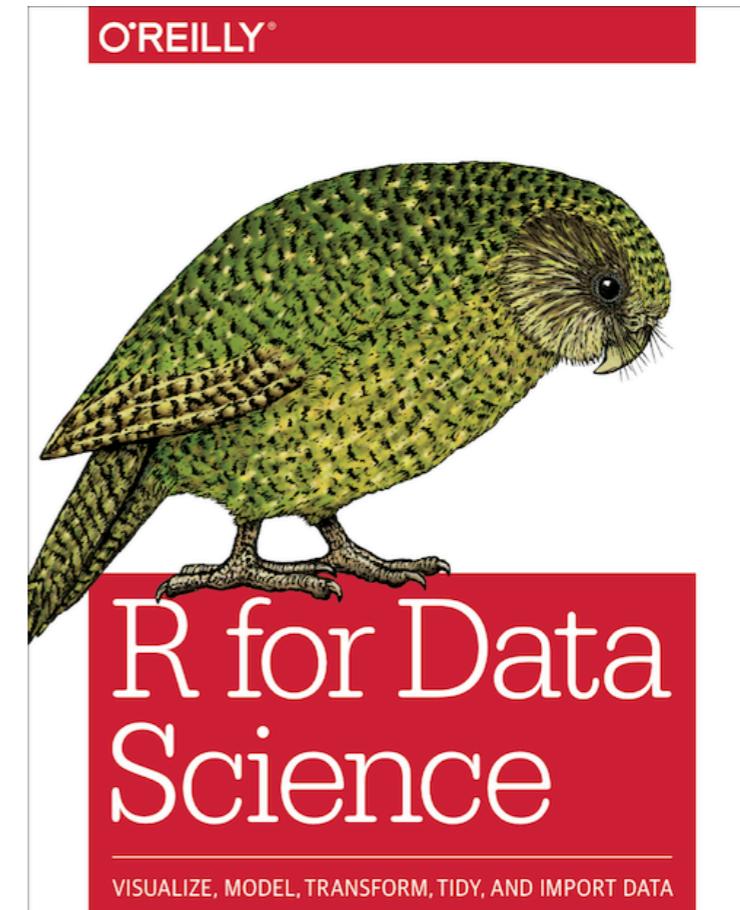
- ggplot2
- patchwork

## Reading

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

## Resources

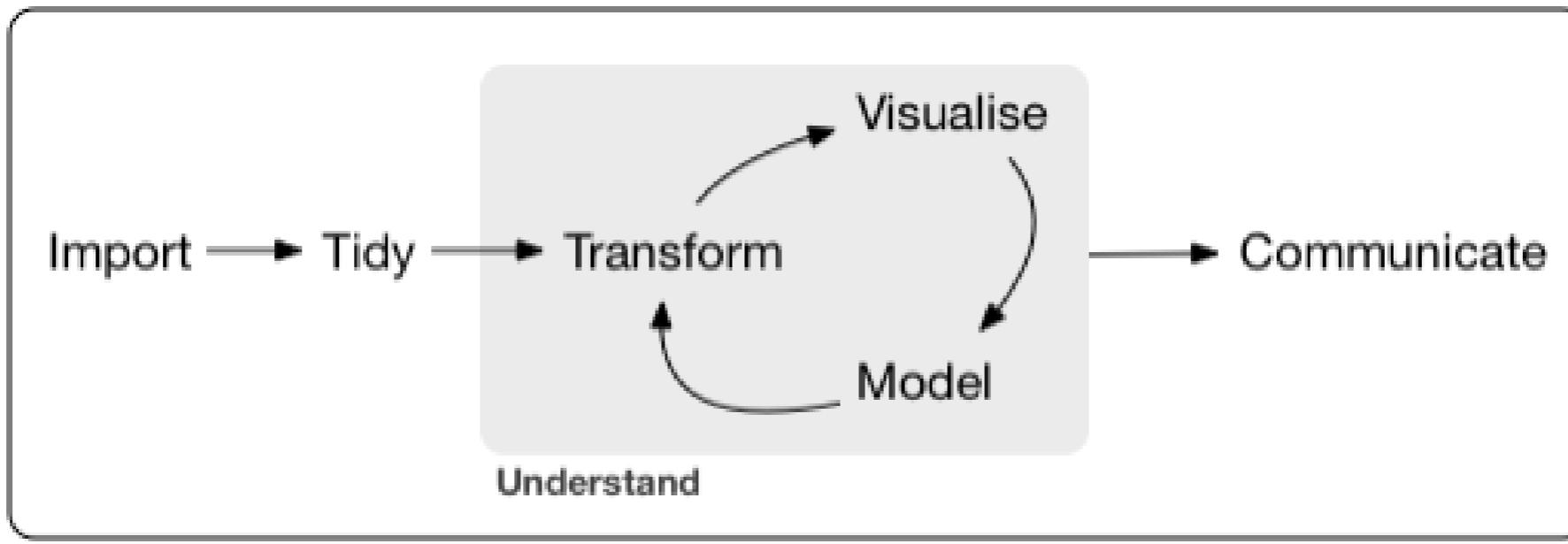
- [Fundamental of data visualization](#)
- [Data visualization: practical introduction](#)



Hadley Wickham &  
Garrett Grolemund

# 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

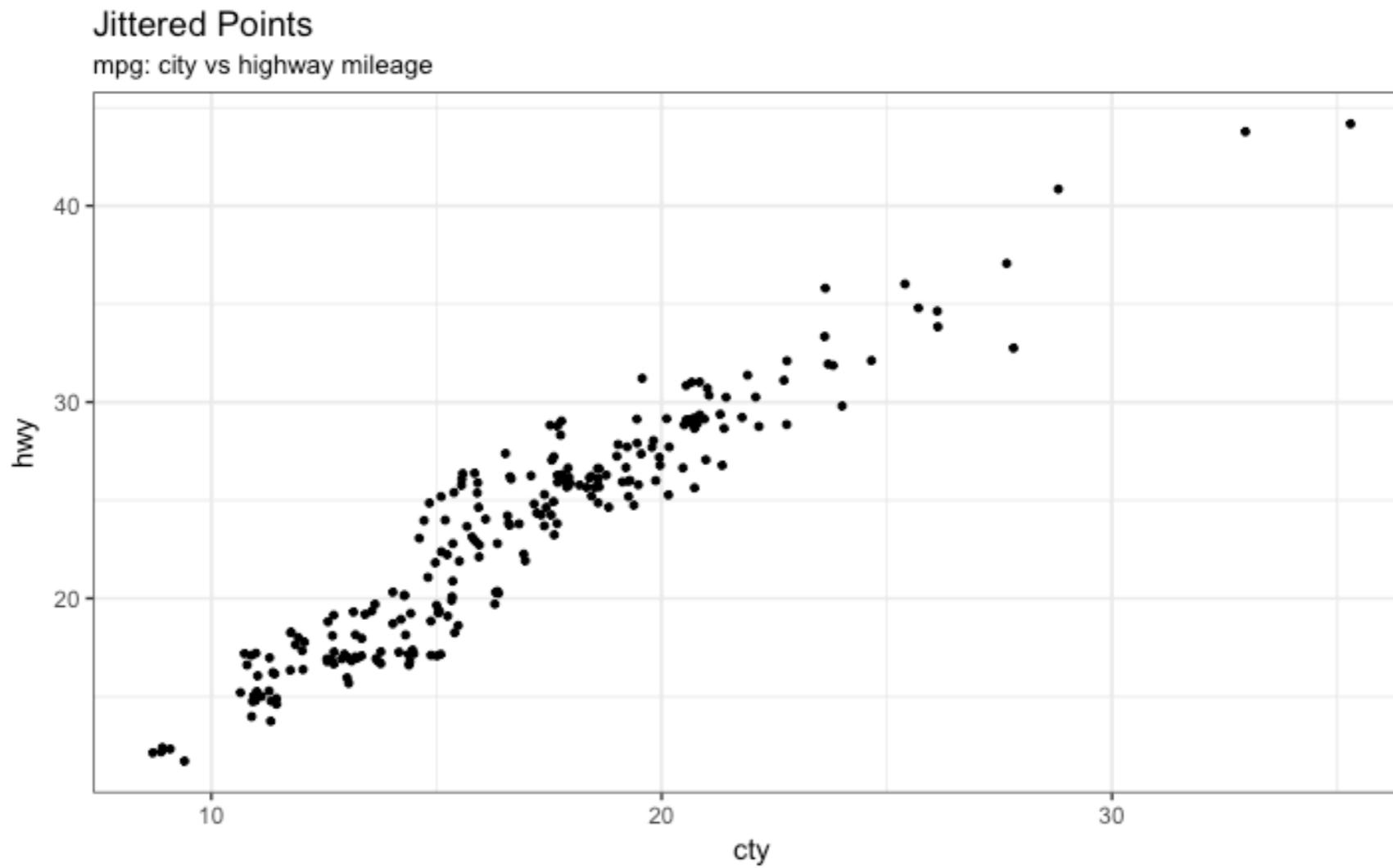
Source: [Michael Friendly](#)

# Graph types

# Jitter

---

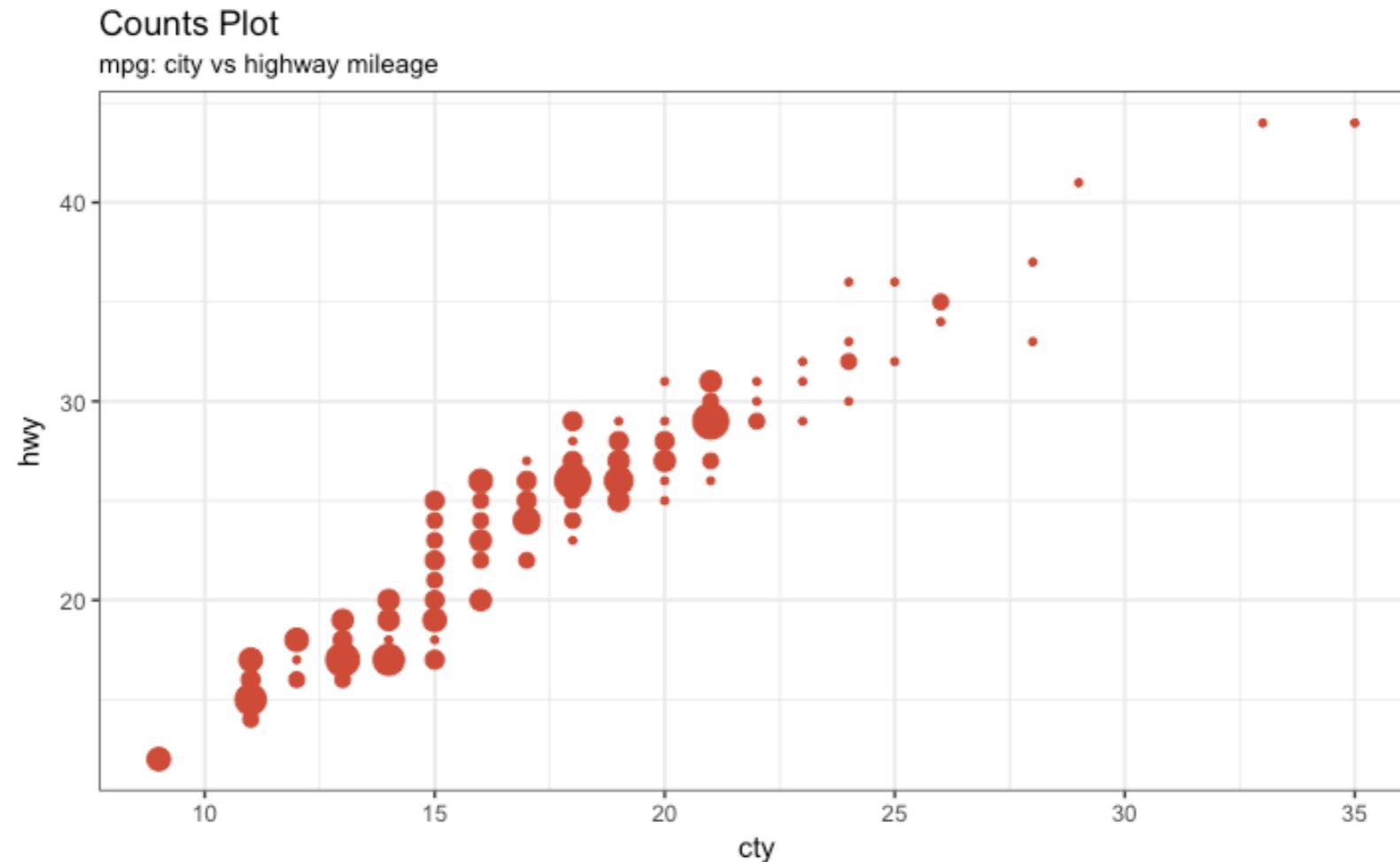
- Two variables numerical



# Bubble

---

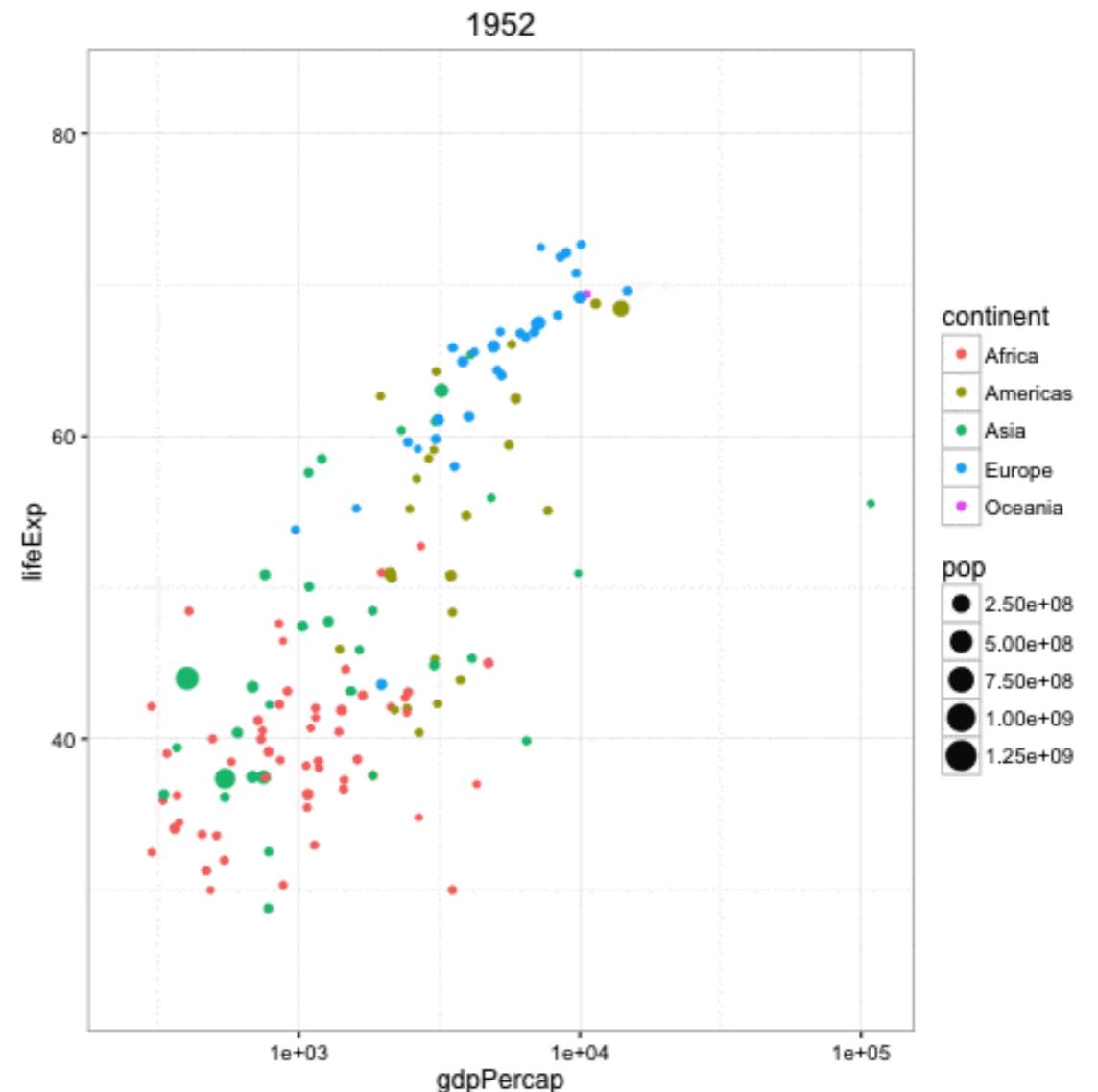
- Two variables numerical
- **Add another variable numerical**



# Animate

---

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

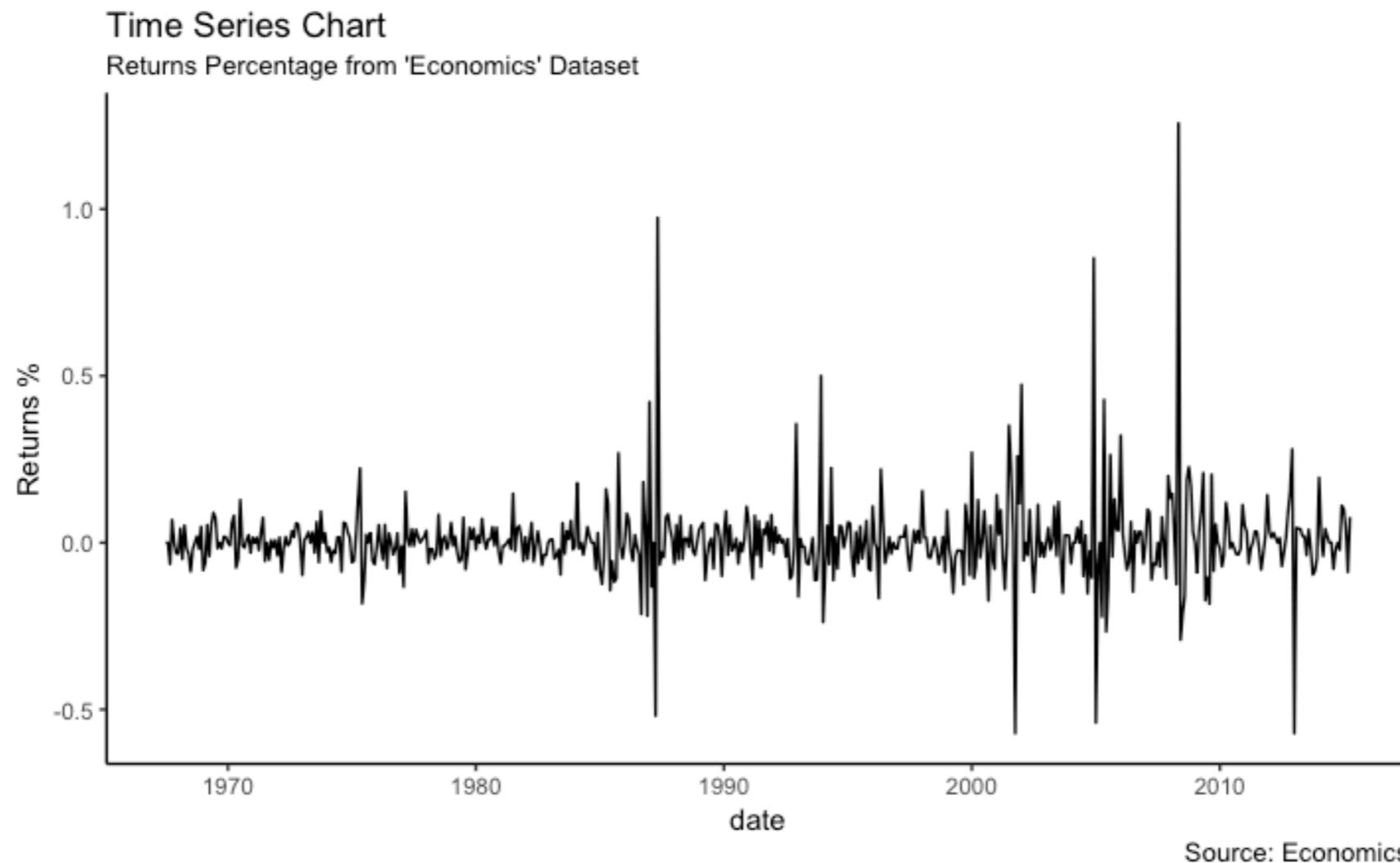




# Times series

---

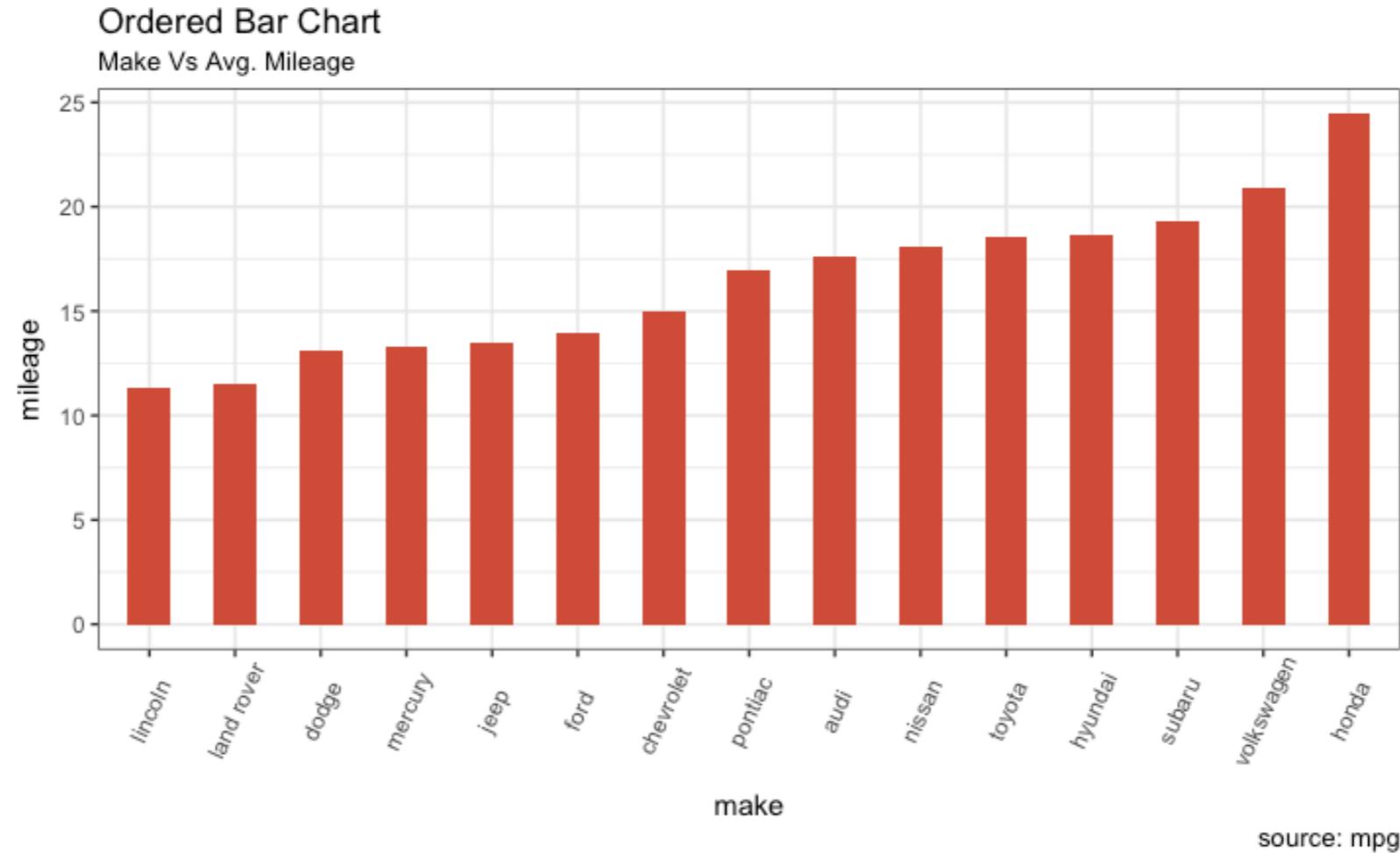
- Line graph



# Bargraphs

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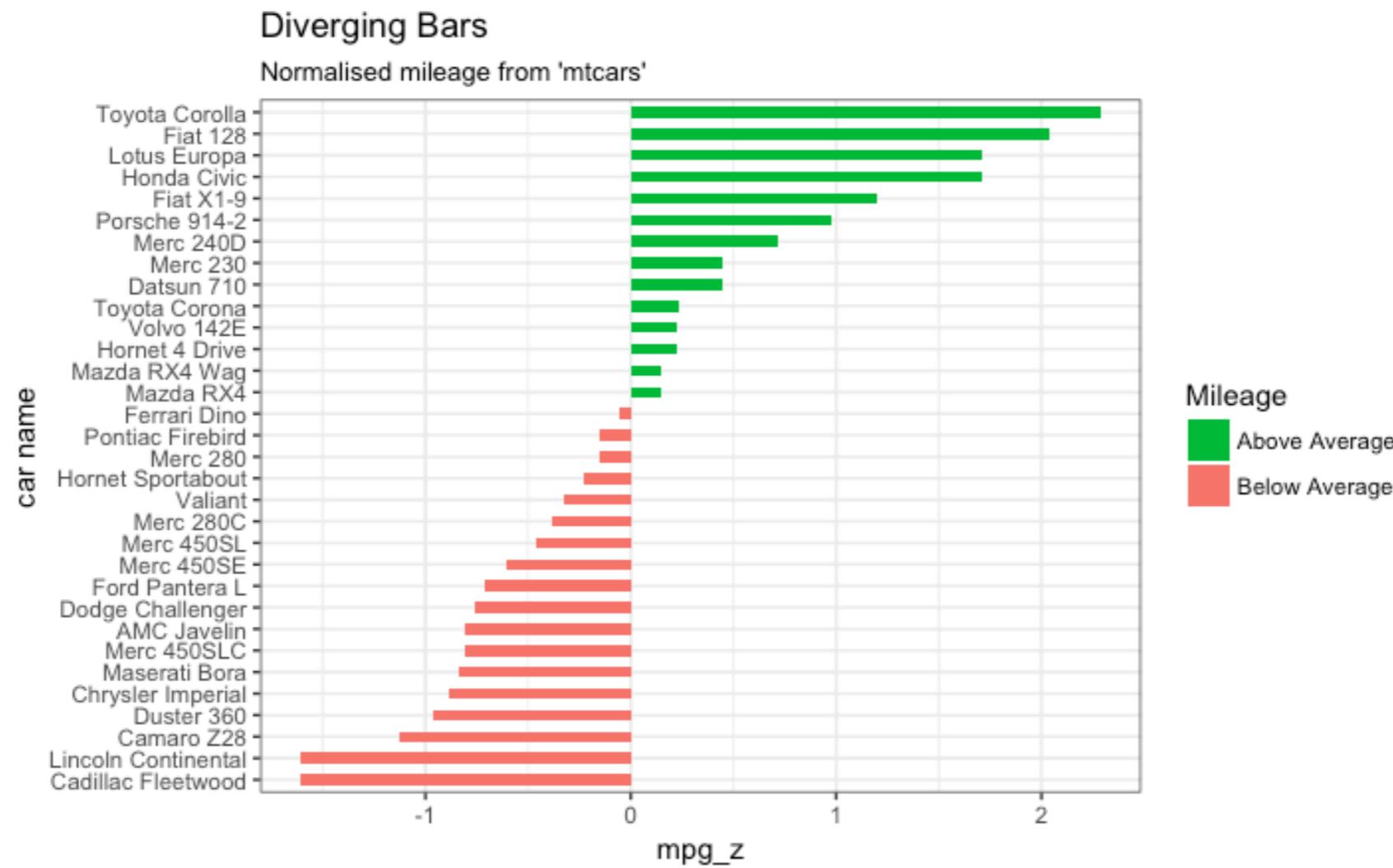
- One variable categorical
- One variable numerical



# Bargraphs

---

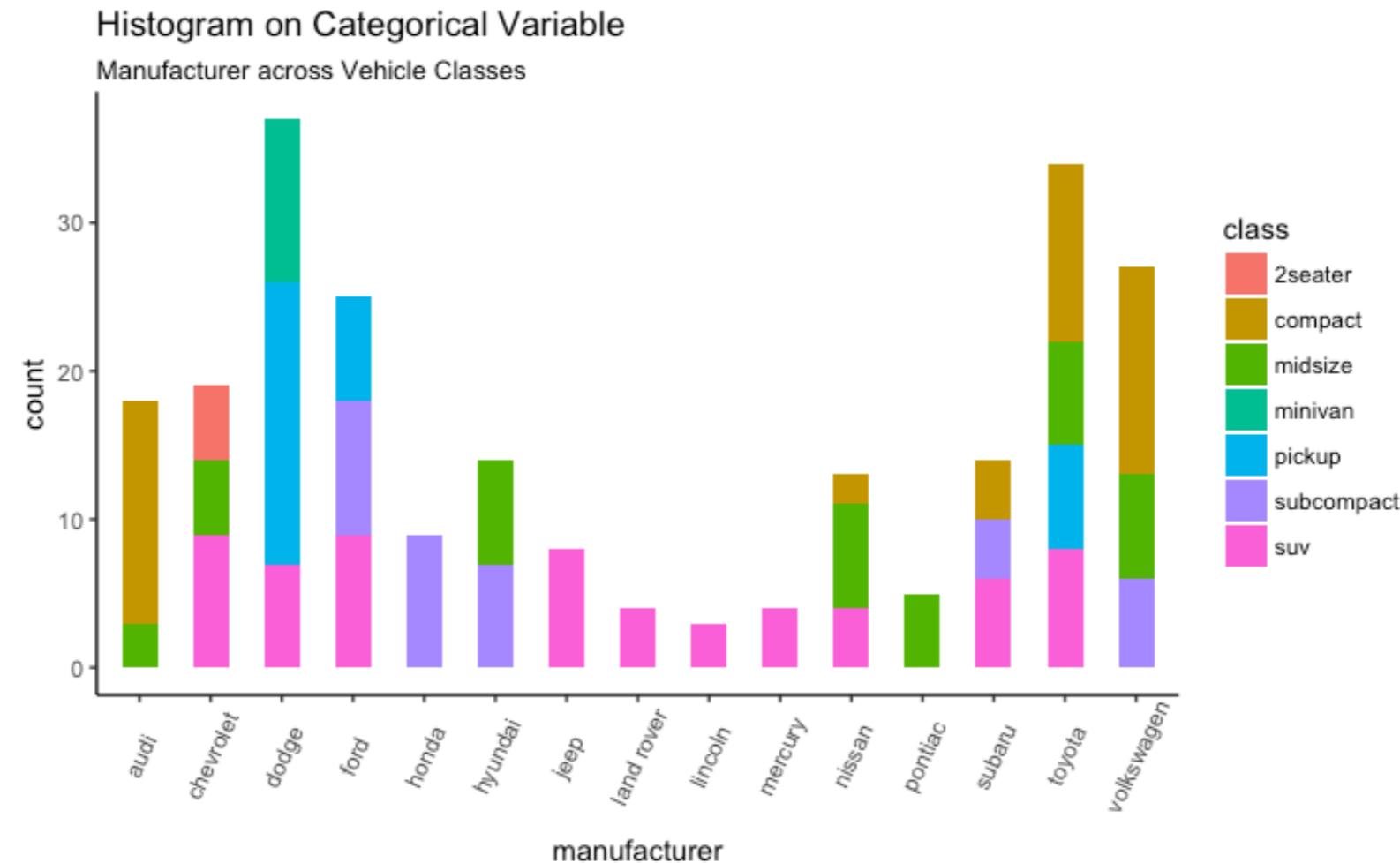
- Rotate



# Bargraphs

---

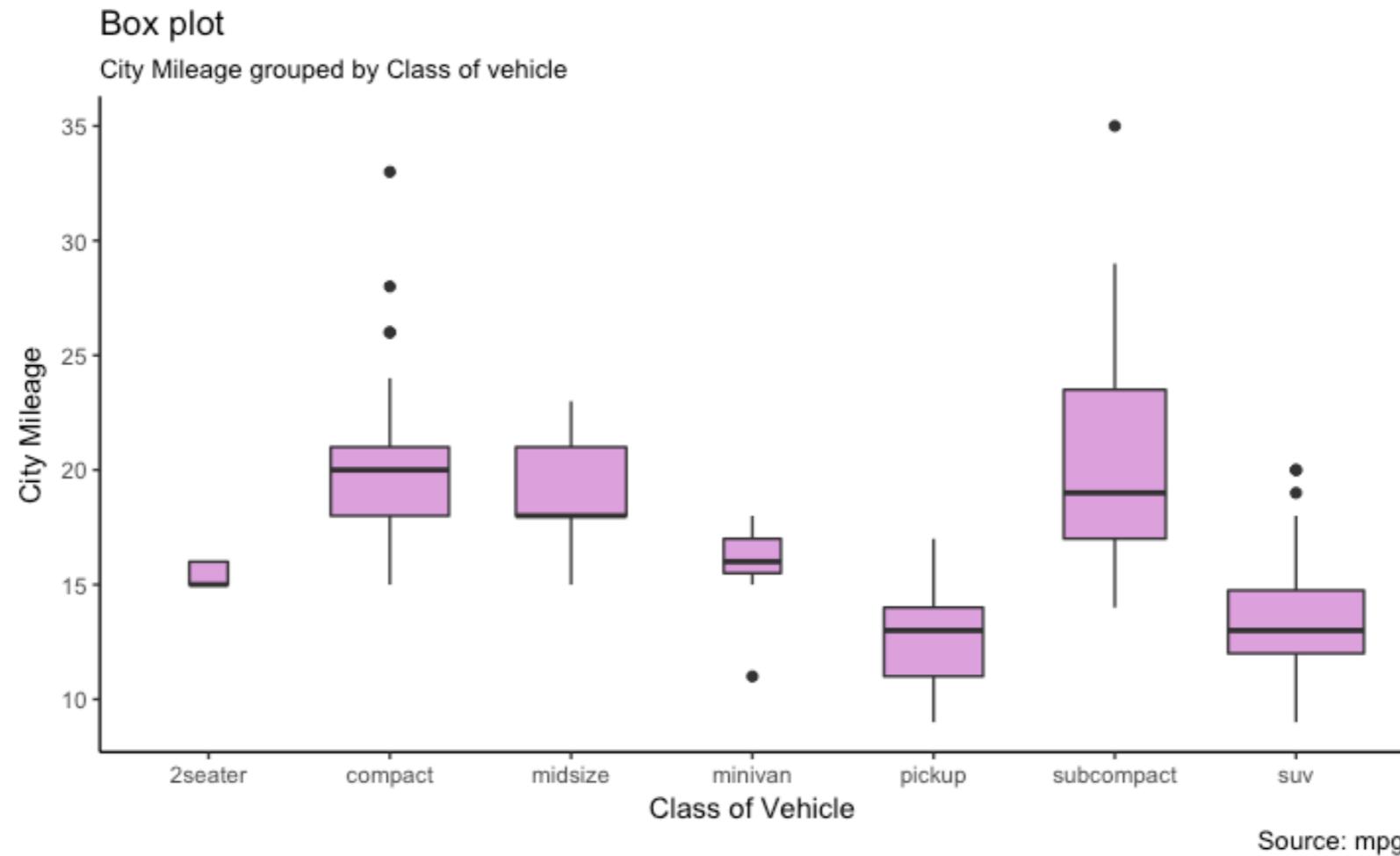
- Two variable categorical
- One variable numerical



# Boxplots

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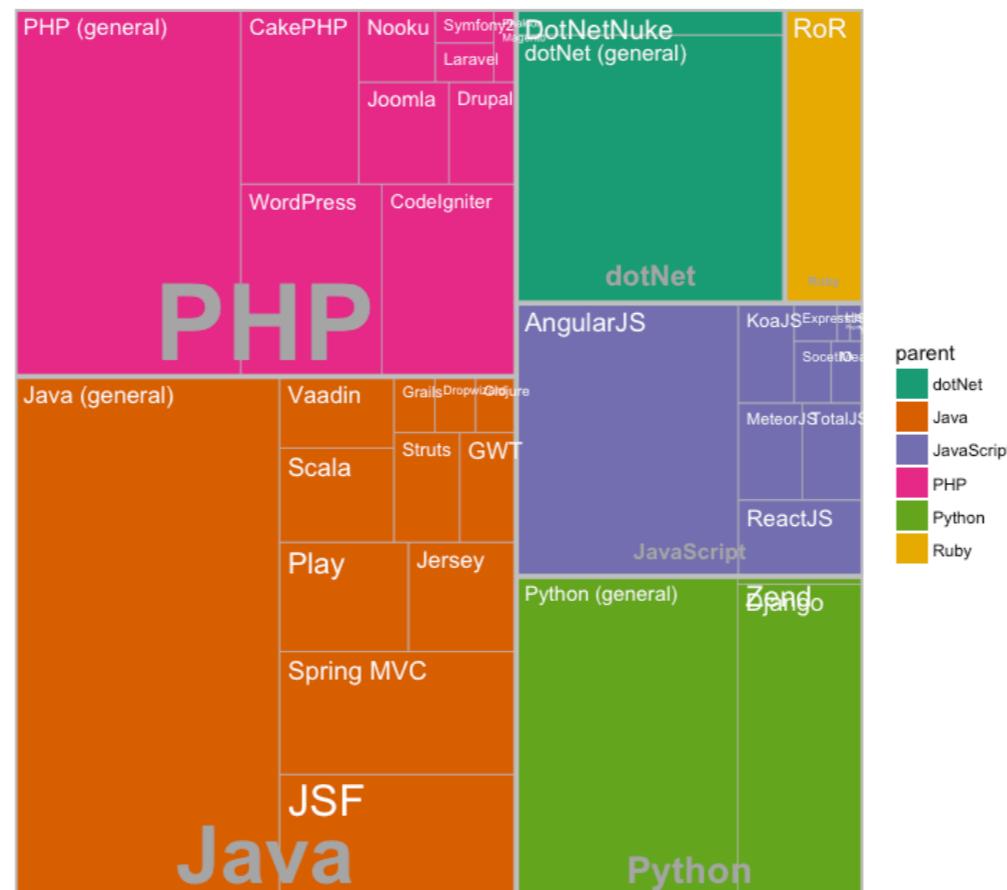
- One variable categorical
- One variable numerical but with many values



# Treemaps

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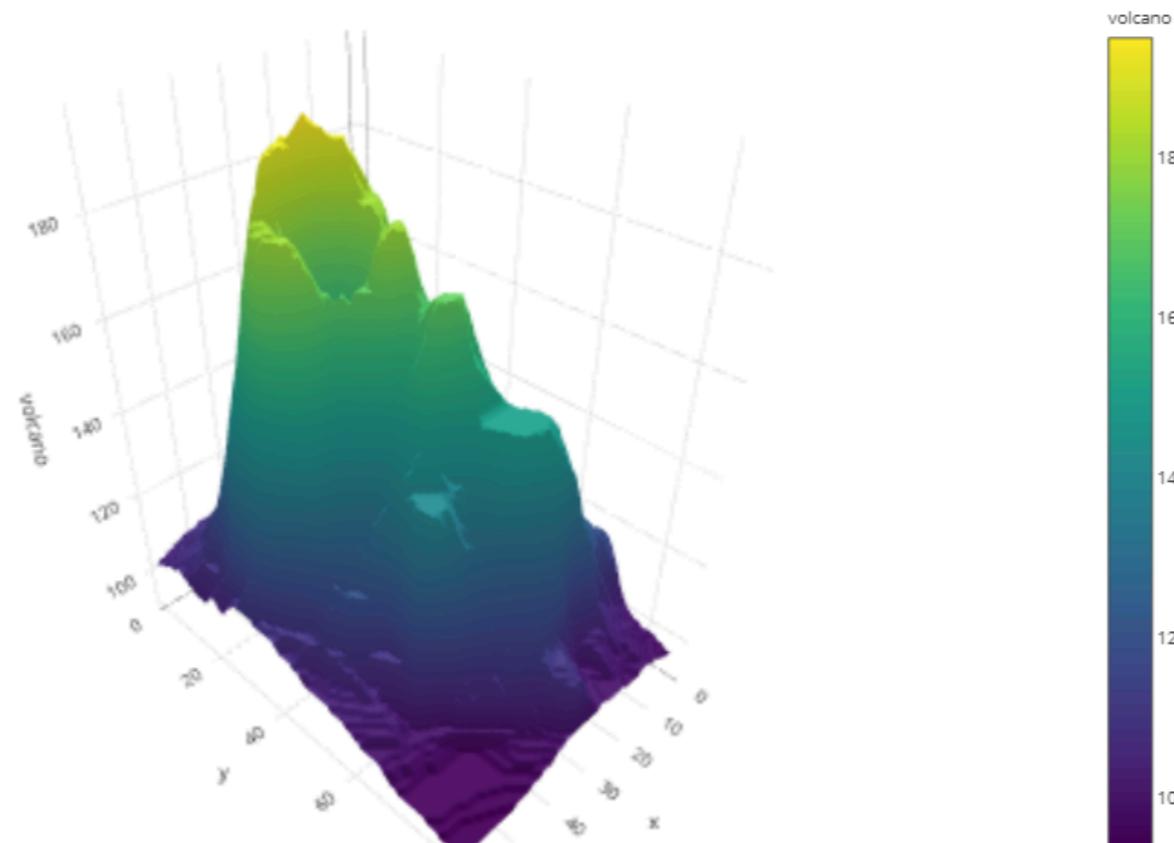
- One variable categorical
- One variable numerical
- Much better than pie charts



# 3D

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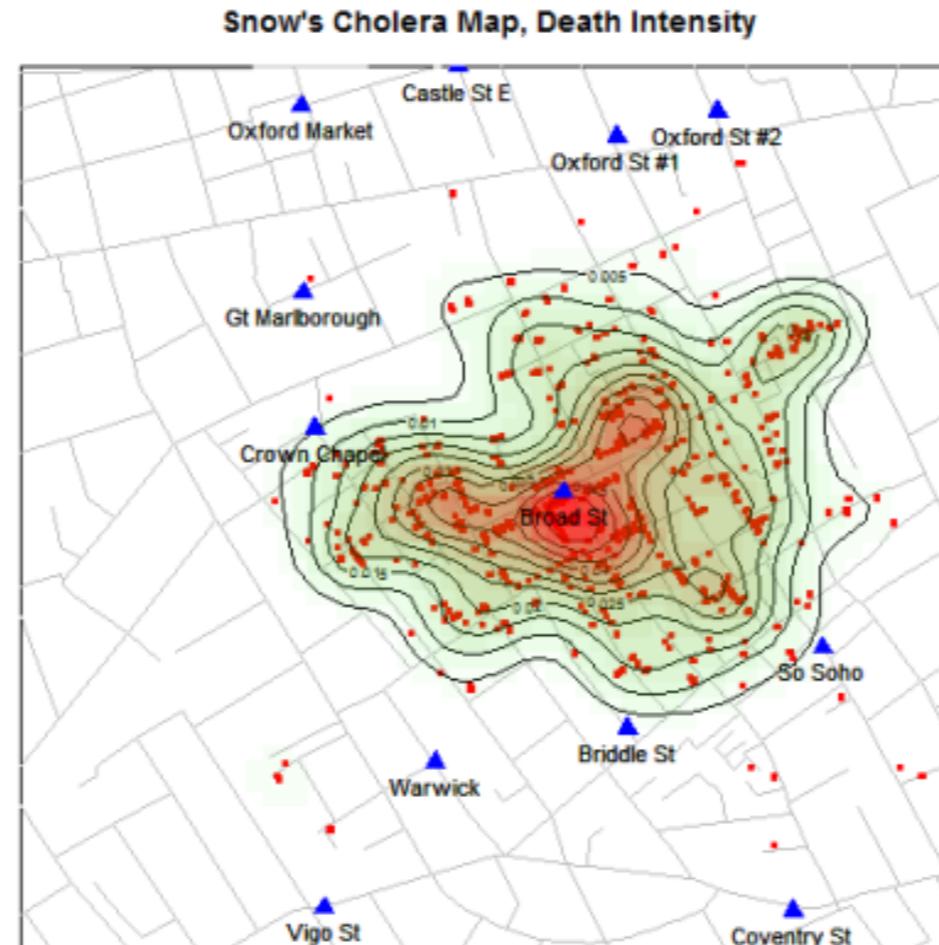
- Three variable numerical
- Avoid unless it is a simple shape



# Contours

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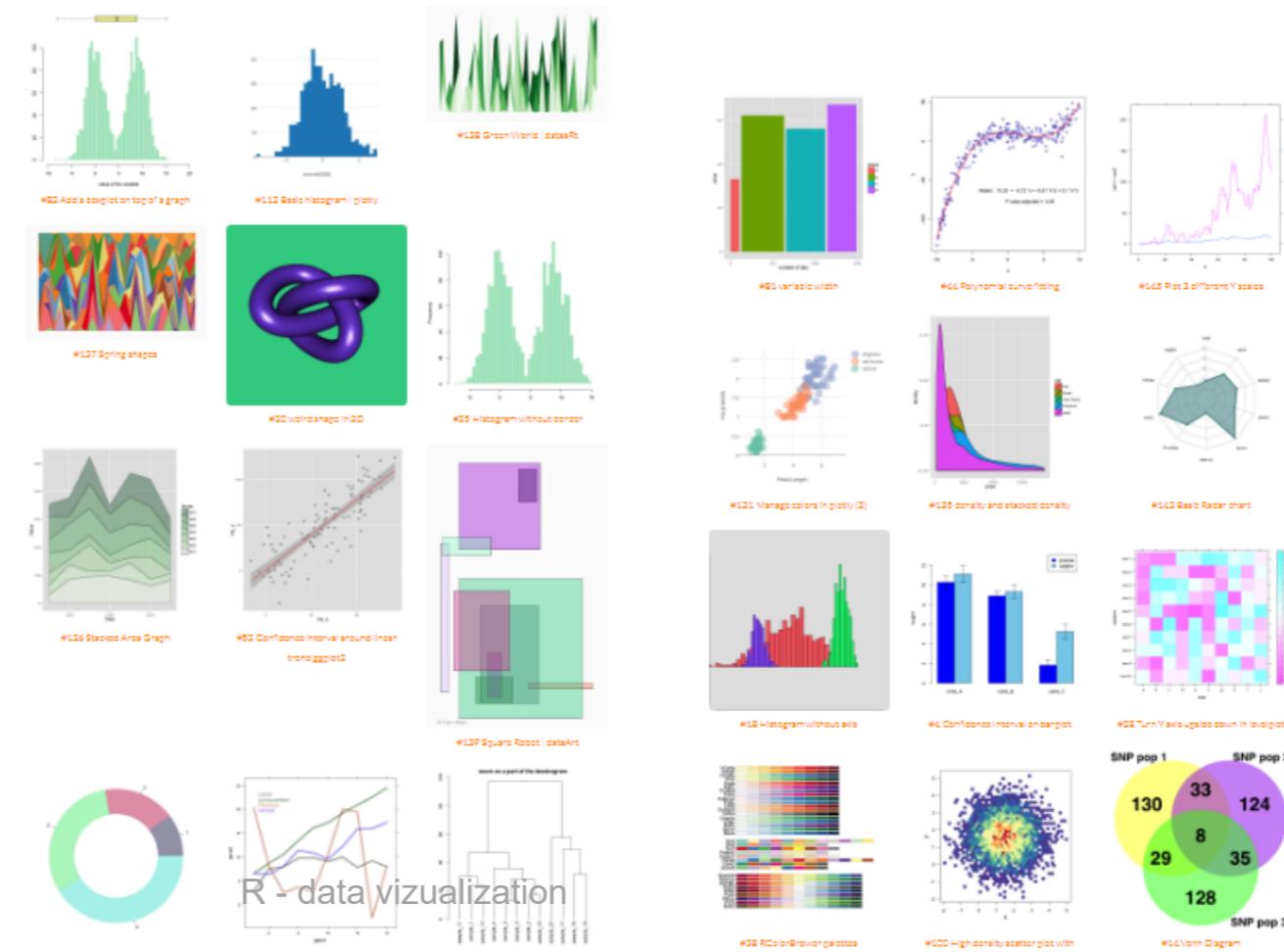
- Three variable numerical
- Better than 3D



# Many...

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- 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/>



# 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") %>%
  tidyverse::fill(station)
```

sample number	transect	station	date	time	depth	level	latitude	longitude	picoeuk	nanoeuk	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

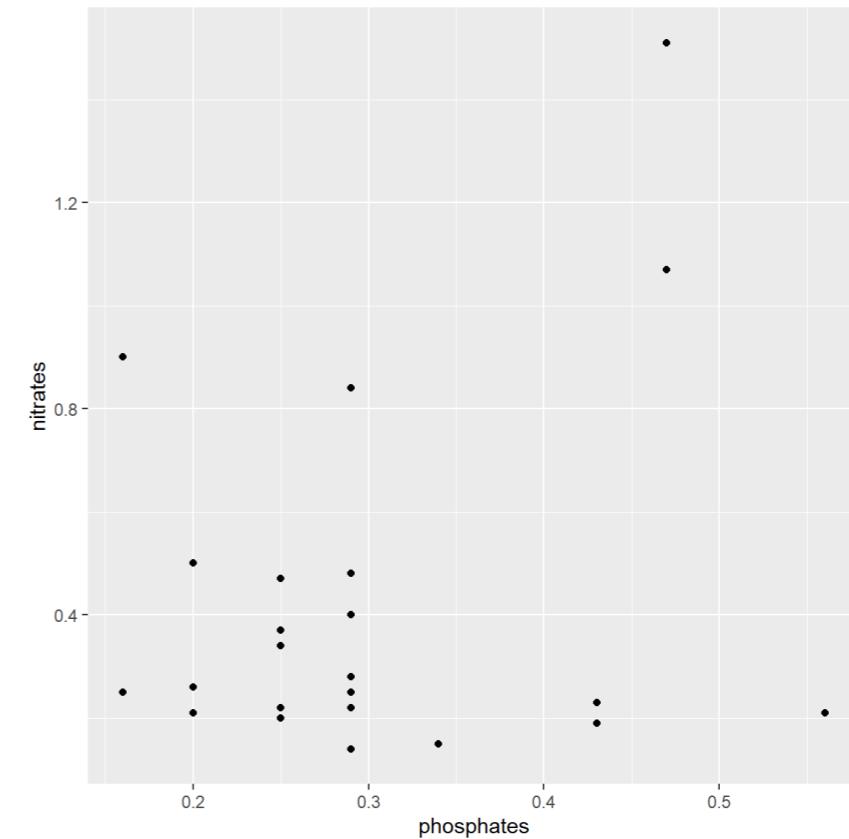
# A simple plot

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

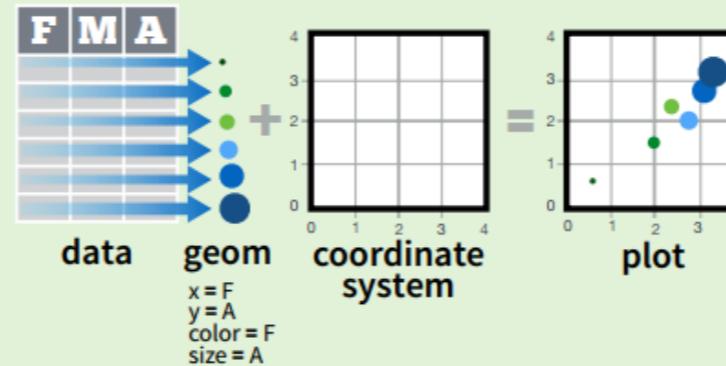
```
# All functions are from ggplot2
# package unless specified

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:

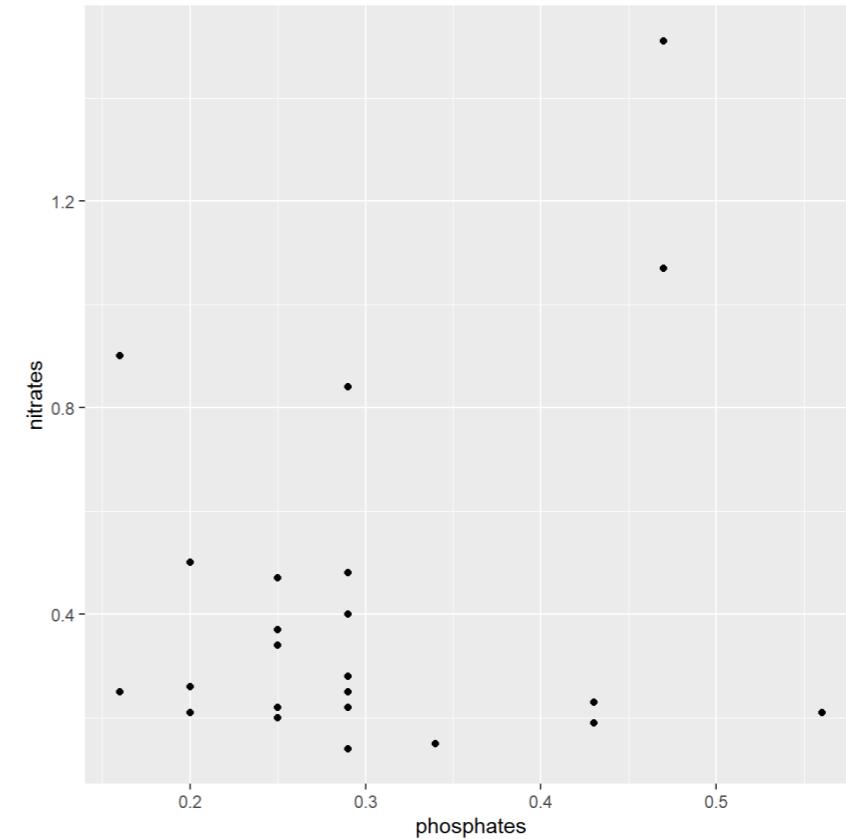
- **data**: a data frame: quantitative, categorical; local or data base query
- **aesthetic mapping** of variables into visual properties: size, color, x, y
- **geometric objects (“geom”)**: points, lines, areas, arrows, ...
- **coordinate system (“coord”)**: Cartesian, log, polar, map

# Alternatively

---

- Move mapping into ggplot function

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

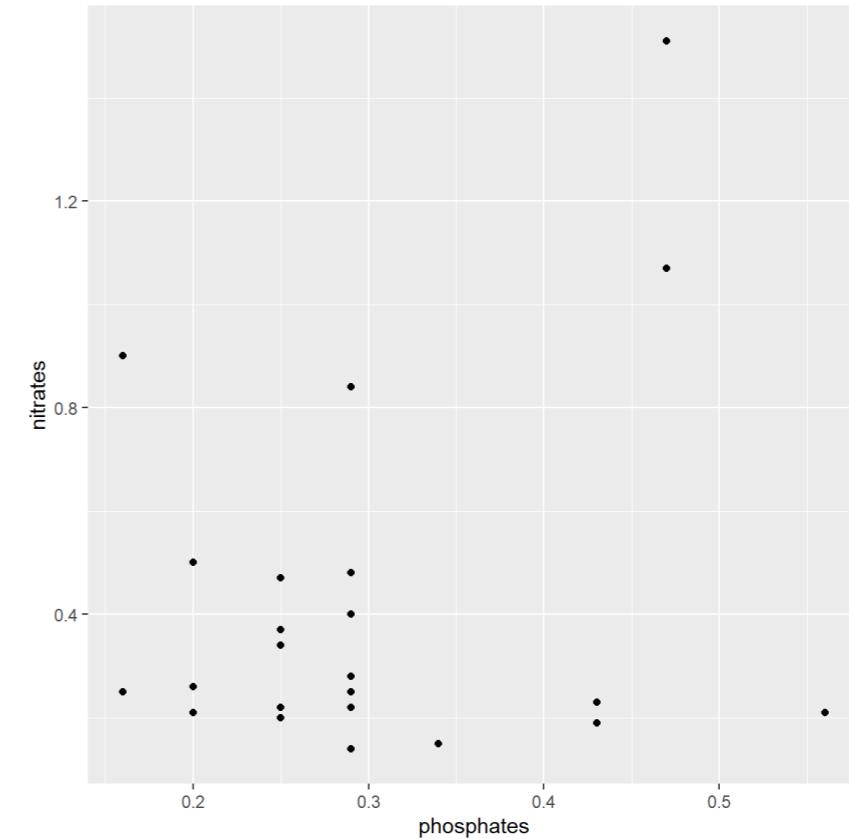


# Alternatively

---

- Remove function arguments

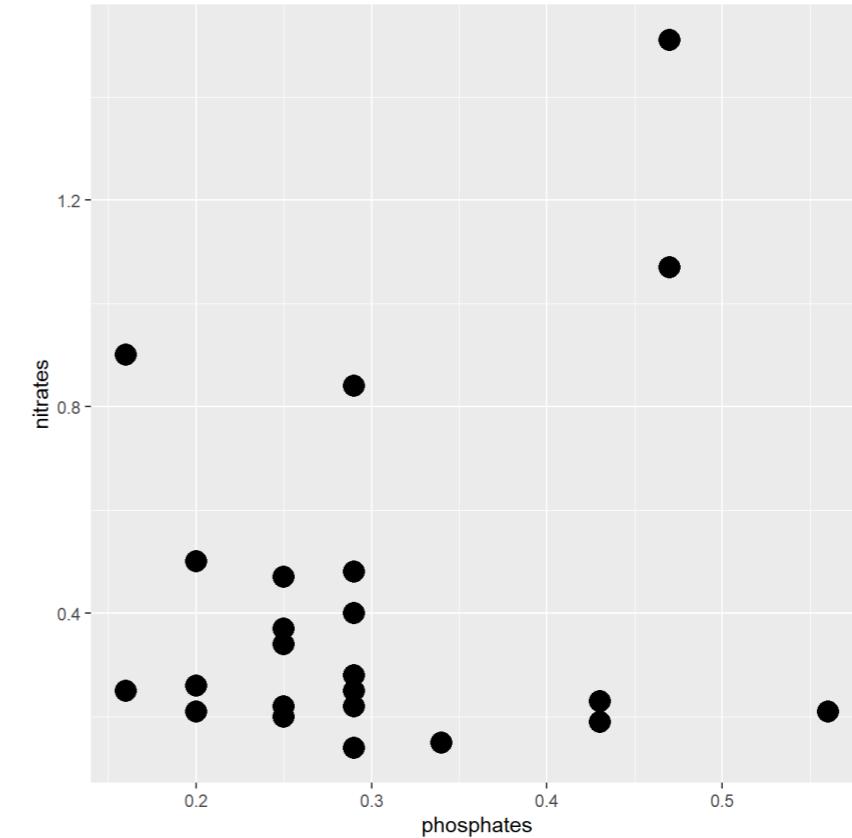
```
ggplot(samples,  
       aes(x=phosphates,  
            y=nitrates)) +  
  geom_point()
```



# Makes dots bigger

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

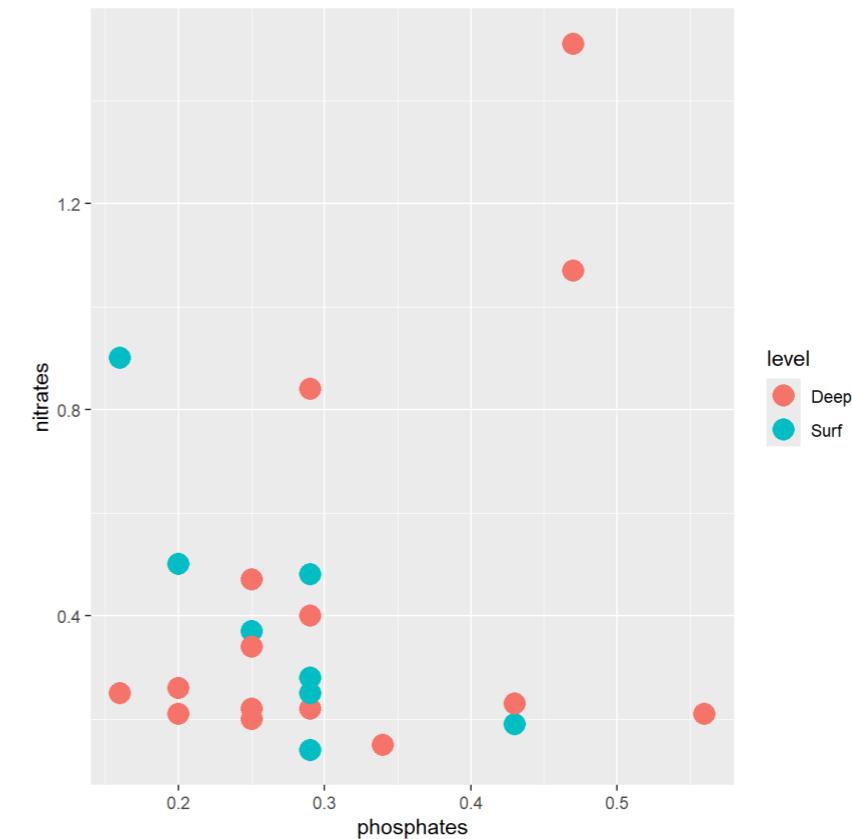
```
ggplot(samples,  
       aes(x=phosphates,  
            y=nitrates)) +  
  geom_point(size=5)
```



# 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...

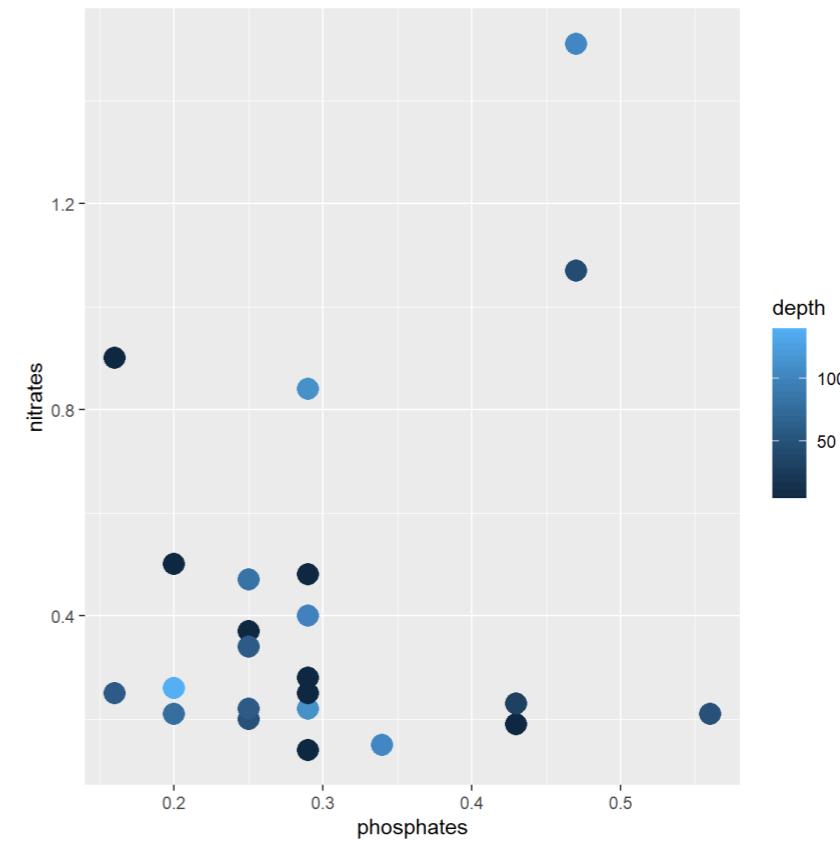
```
ggplot(samples,  
       aes(x=phosphates,  
            y=nitrates)) +  
  geom_point(color=level, size=5)
```

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**

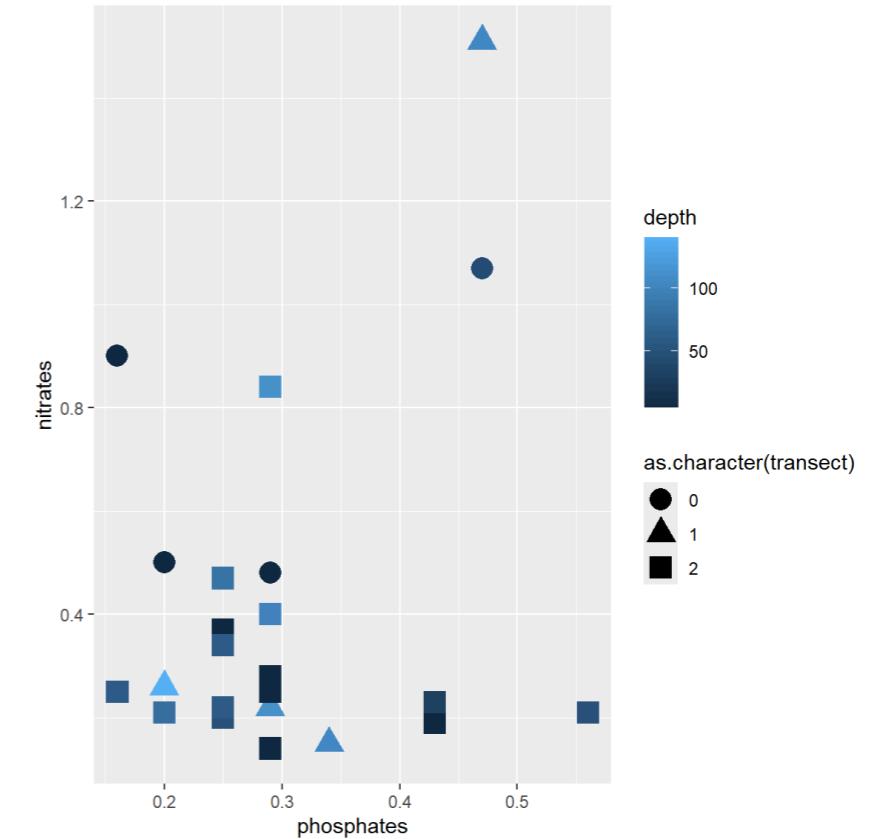
```
ggplot(samples,  
       aes(x=phosphates,  
            y=nitrates,  
            color=depth,  
            shape=transect)) +  
  geom_point(size=5)
```

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)

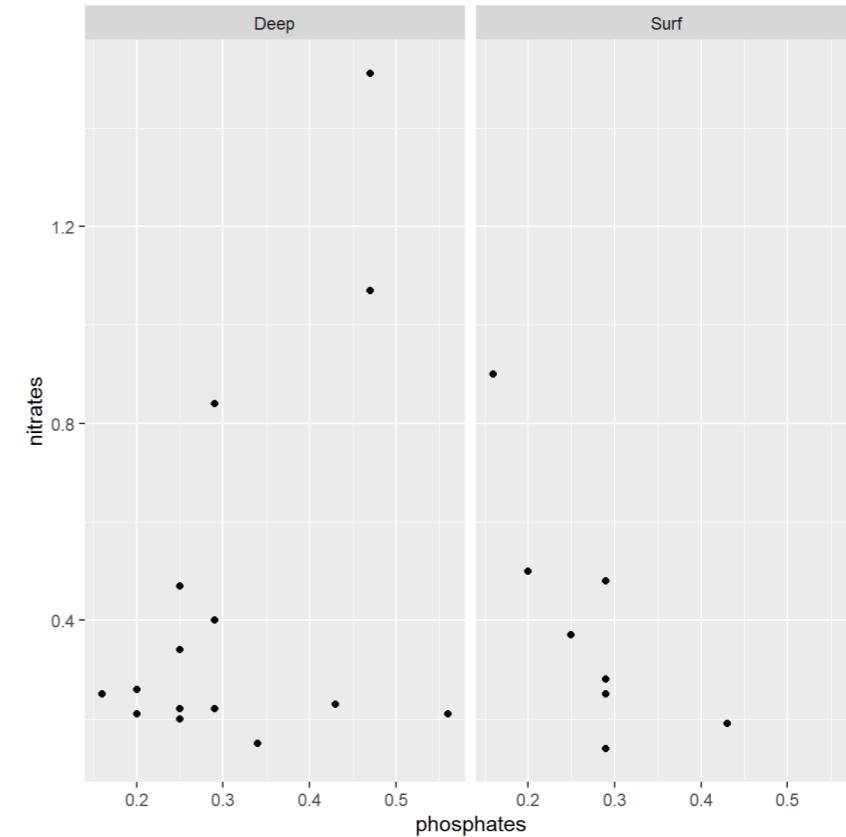
- Add: **shape=as.character(transect)**

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



# Panels depending on one variable

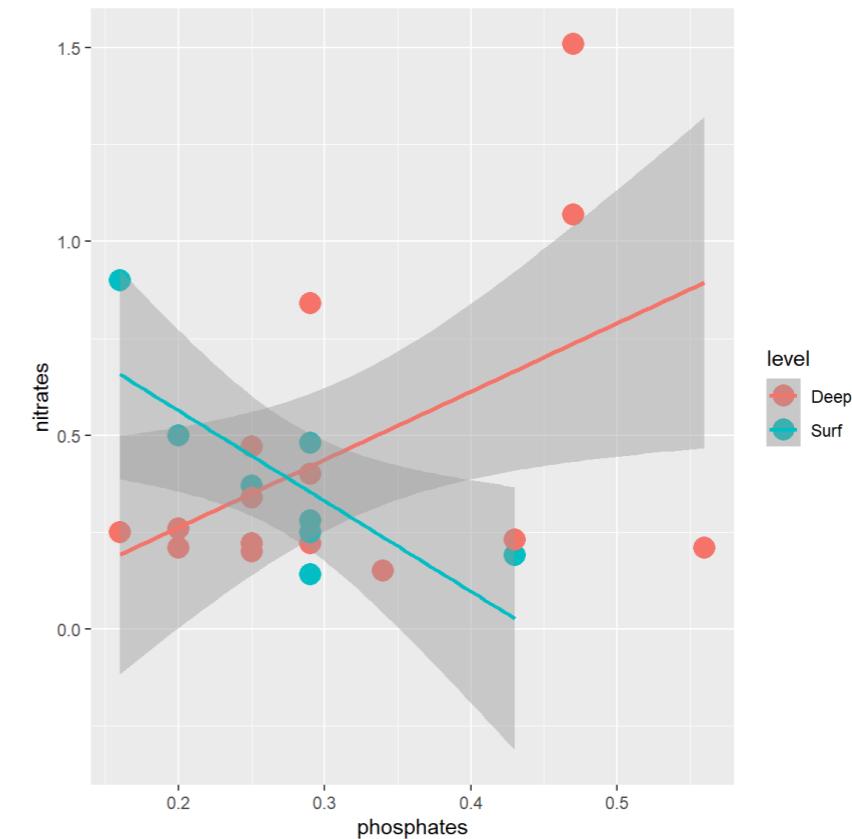
```
ggplot(samples,  
       aes(x=phosphates,  
            y=nitrates)) +  
  geom_point() +  
  facet_wrap(~ level)
```



# Adding a regression line

- Add: **geom\_smooth()**
- You can choose the type of smoothing “lm” is for linear model

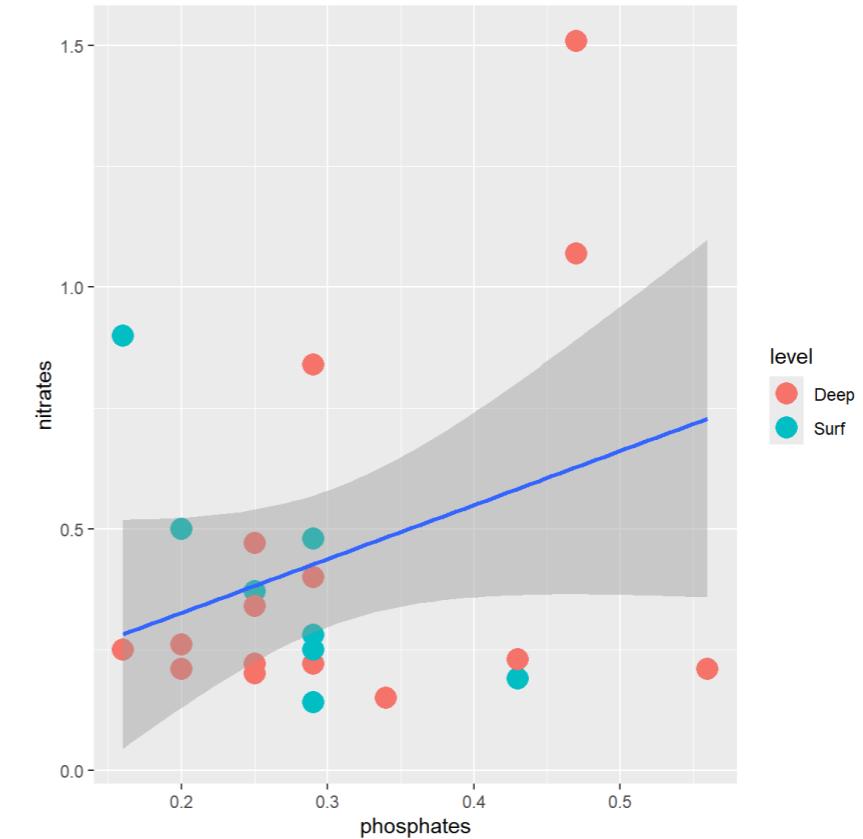
```
ggplot(samples,  
       aes(x=phosphates,  
            y=nitrates,  
            color=level)) +  
  geom_point(size=5) +  
  geom_smooth(mapping = aes(x=phosphates,  
                            y=nitrates),  
              method="lm")
```



# Adding a regression line

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

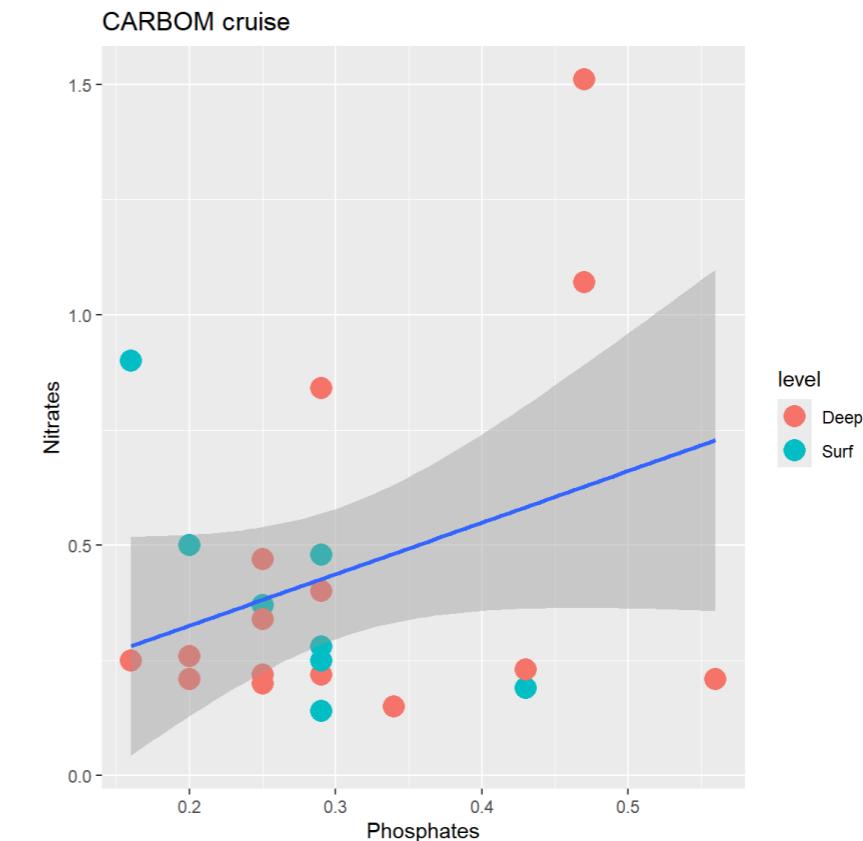
```
ggplot(samples,  
       aes(x=phosphates,  
            y=nitrates)) +  
  geom_point(aes(color=level),  
             size=5) +  
  geom_smooth(mapping = aes(x=phosphates,  
                            y=nitrates),  
              method="lm")
```



# Finalizing the graph

- Adding labels and legends

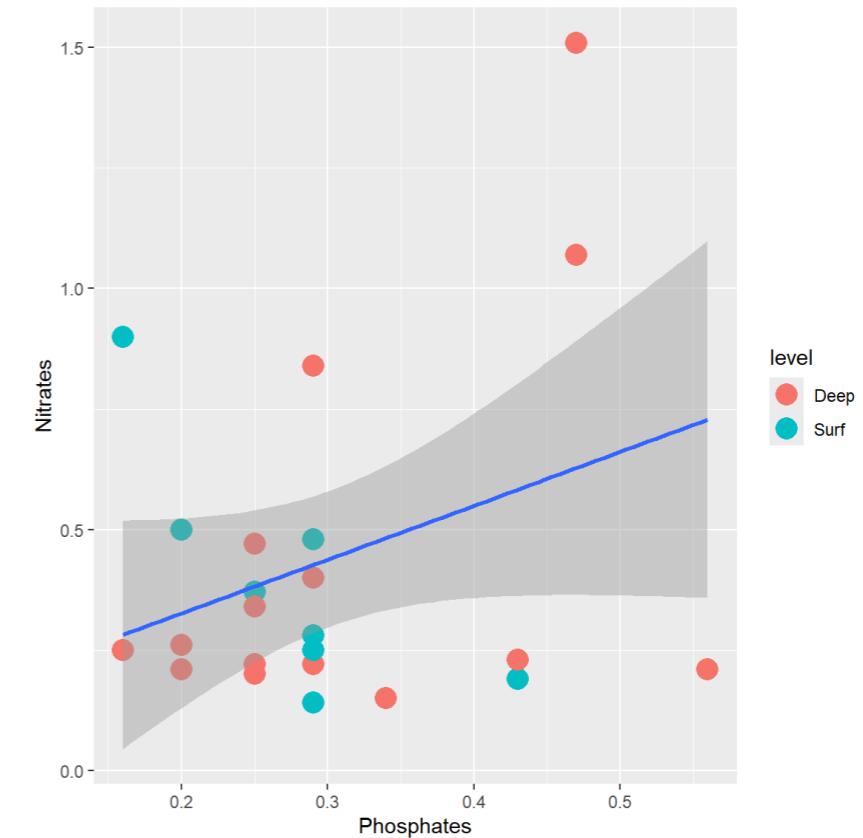
```
ggplot(samples) +  
  geom_point(mapping = aes(x=phosphates,  
                            y=nitrates,  
                            color=level),  
              size=5) +  
  geom_smooth(mapping = aes(x=phosphates,  
                            y=nitrates),  
              method="lm") +  
  xlab("Phosphates")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



# Multigraphs (patchwork package)

# First graph

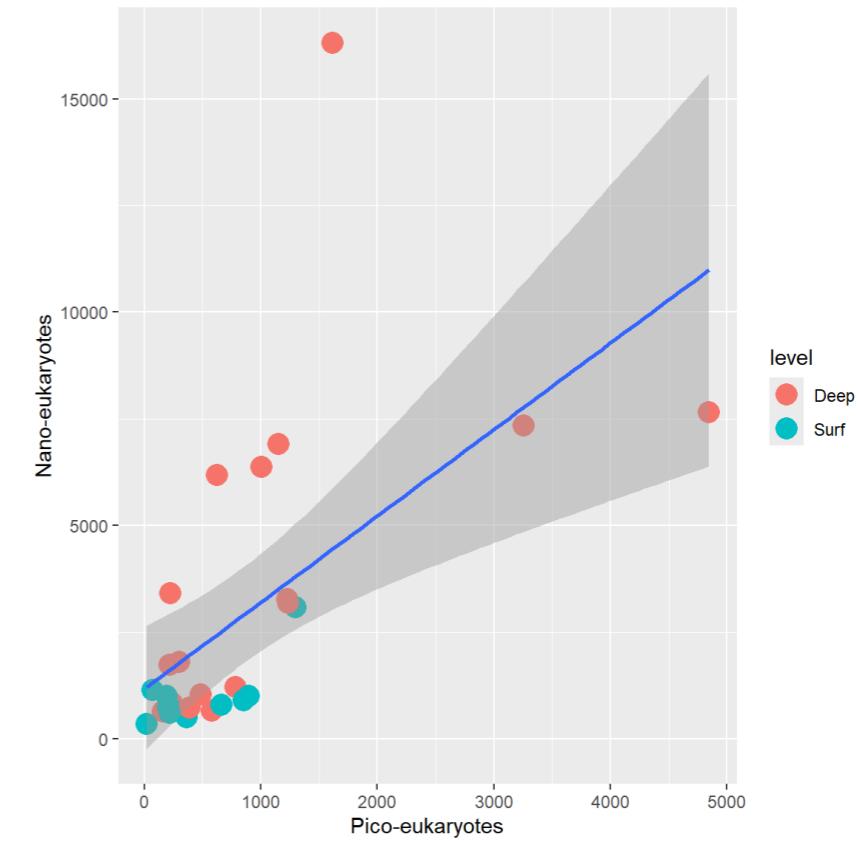
```
g1 <- ggplot(samples) +  
  geom_point(mapping = aes(x=phosphates,  
                            y=nitrates, color=  
                            level), size=5)  
  geom_smooth(mapping = aes(x=phosphates,  
                            y=nitrates),  
              method="lm") +  
  xlab("Phosphates")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



## Second graph

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

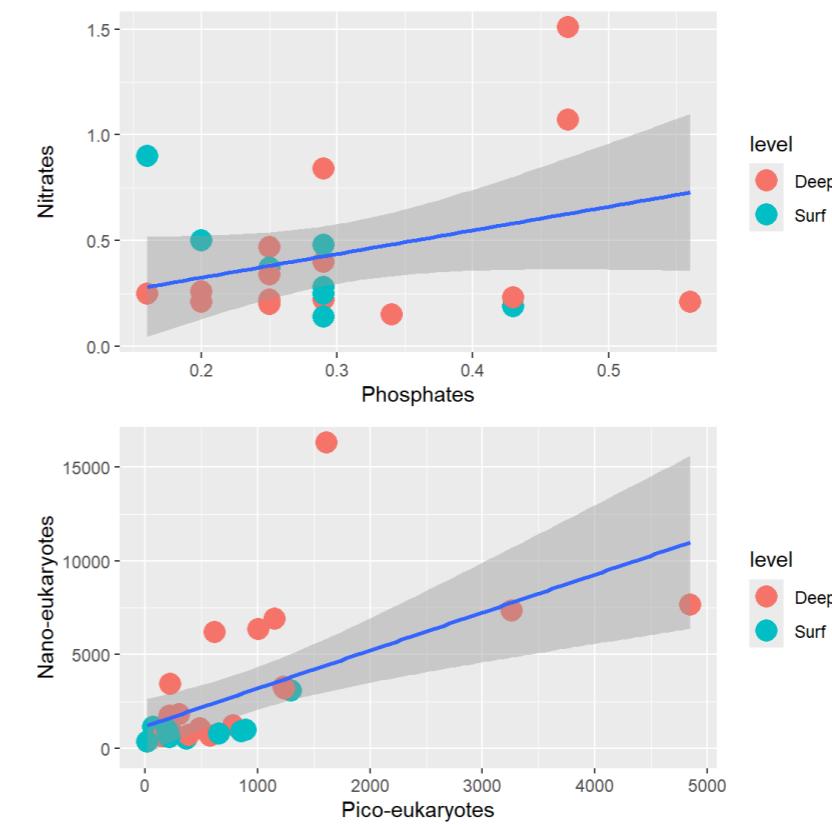
g2



# Package patchwork

- Other packages:
  - `gridExtra`
  - `cowplot`

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

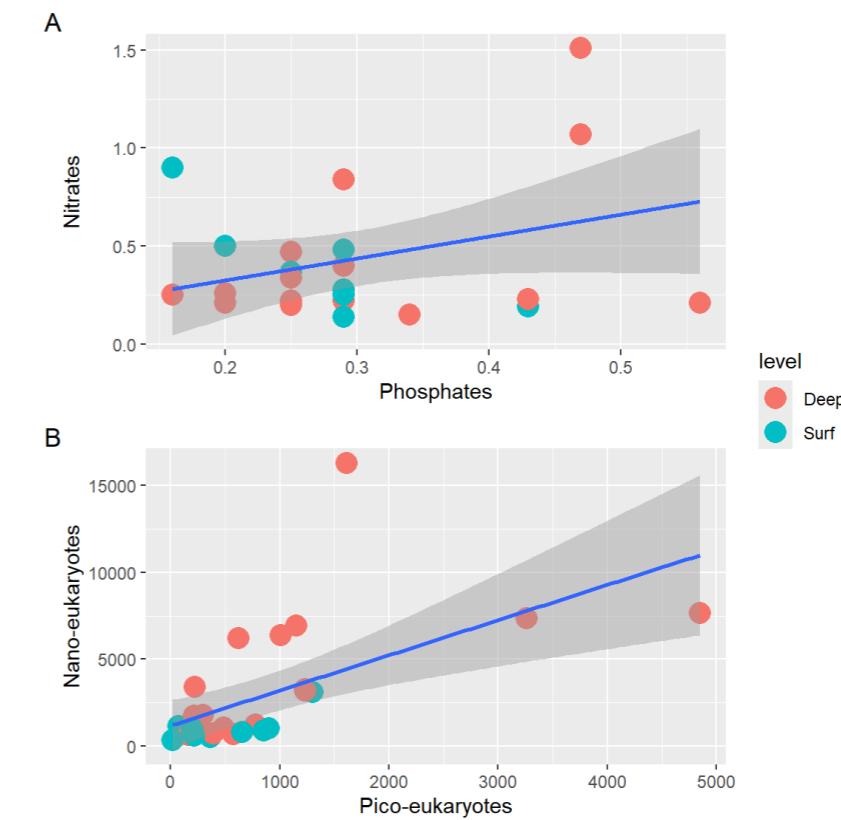


# Package patchwork

- Adding annotation
- Collecting legends

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

R - data vizualization



# Package esquisse

esquisse 2.0.1.9000

Reference

Articles

Changelog

Search for



## Get started with esquisse

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

```
library(esquisse)
```



### On this page

[Launch the addin](#)

[Import data into {esquisse}](#)

[Create a plot](#)

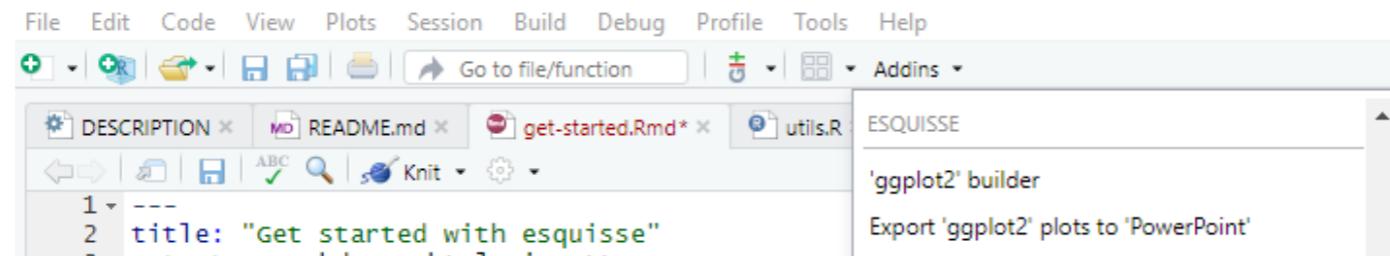
[Controls](#)

[Export](#)

[Addin options](#)

### Launch the addin

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



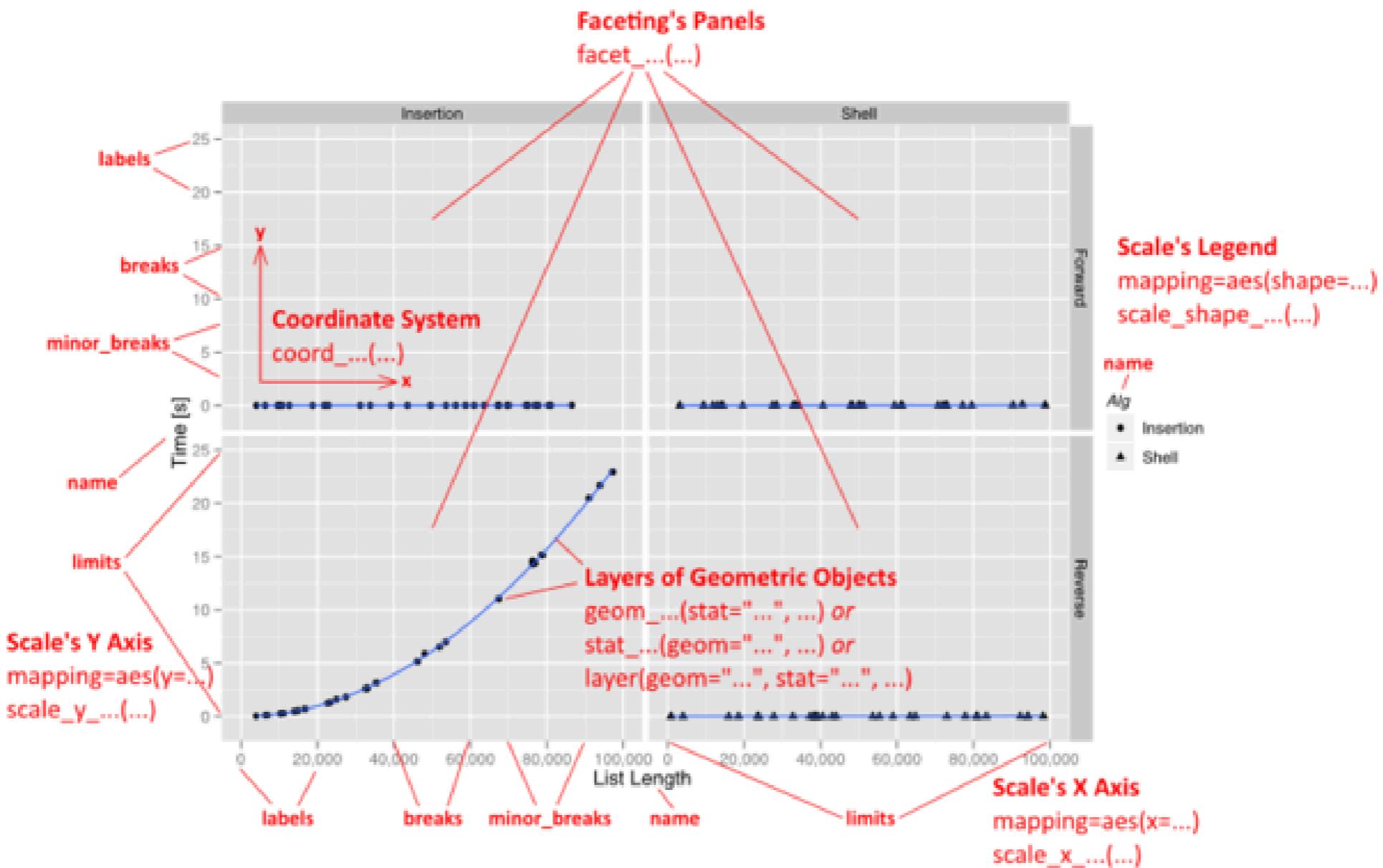
Or in the R console :

```
esquisser()
```

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

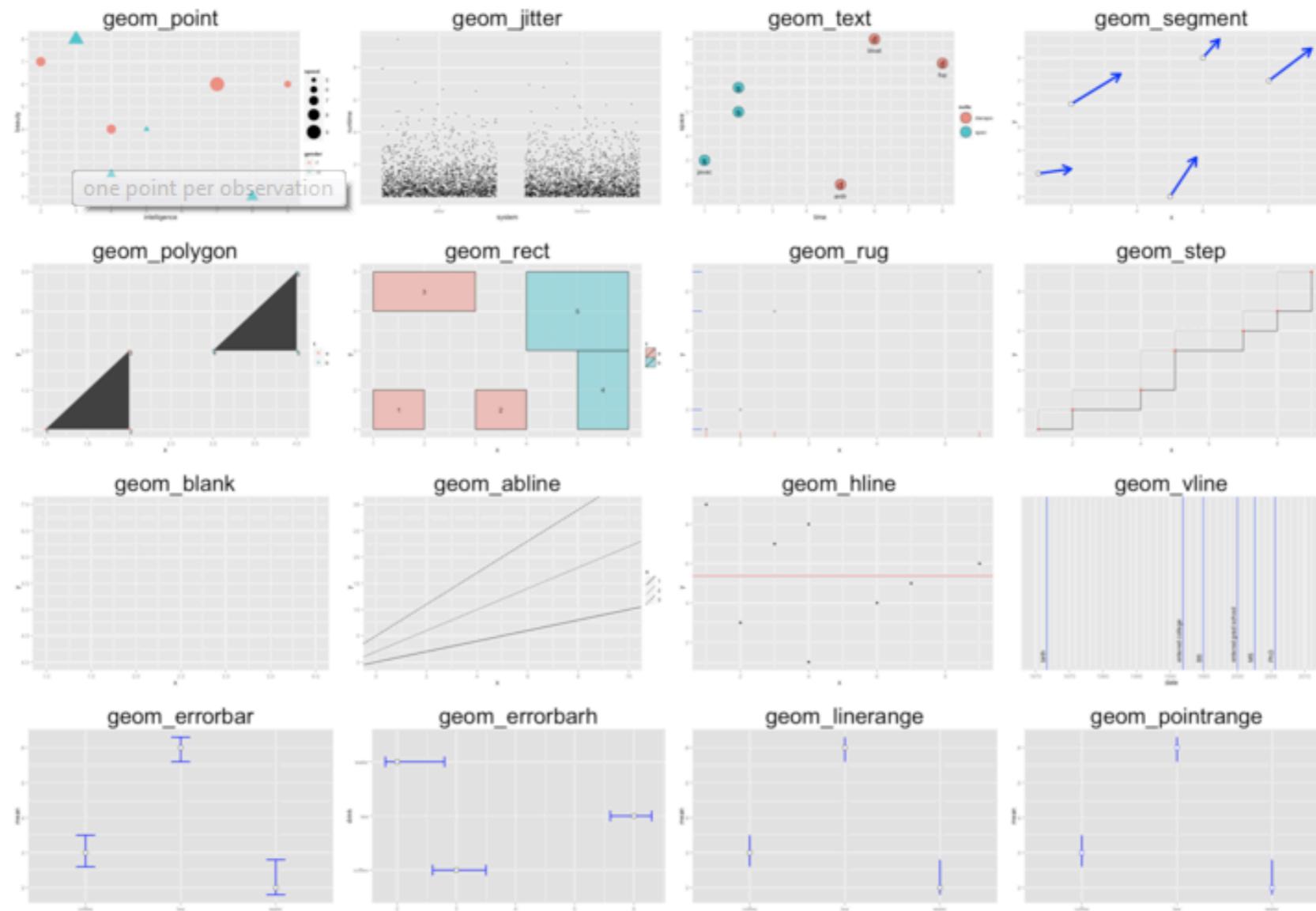
# **ggplot2 syntax**

# Anatomy of a plot



# Geometries

---



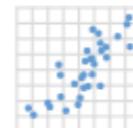
# Continuous x and y

## continuous x , continuous y

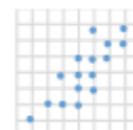
```
e <- ggplot(mpg, aes(cty, hwy))
```



**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



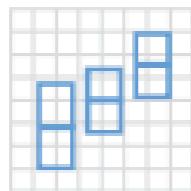
**e + geom\_text(aes(label = cty), nudge\_x = 1,  
nudge\_y = 1, check\_overlap = TRUE)**, x, y, label,  
alpha, angle, color, family, fontface, hjust,  
lineheight, size, vjust

# Plotting error

---

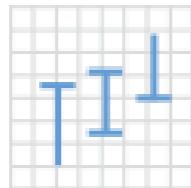
## visualizing error

```
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))
```

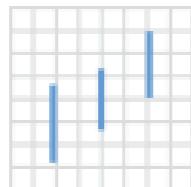


**j + geom\_crossbar(fatten = 2)**

x, y, ymax, ymin, alpha, color, fill, group, linetype,  
size

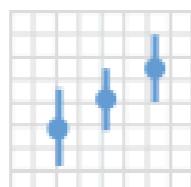


**j + geom\_errorbar(), x, ymax, ymin, alpha, color,  
group, linetype, size, width (also  
geom\_errorbarh())**



**j + geom\_linerange()**

x, ymin, ymax, alpha, color, group, linetype, size



**j + geom\_pointrange()**

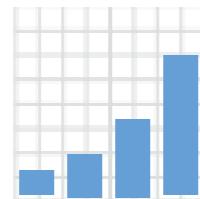
x, y, ymin, ymax, alpha, color, fill, group, linetype,  
shape, size

## Discrete x - Continuous y

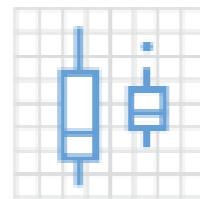
---

### discrete x , continuous y

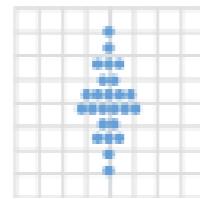
```
f <- ggplot(mpg, aes(class, hwy))
```



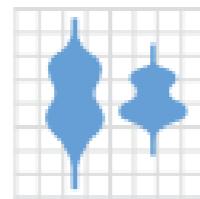
**f + geom\_col()**, x, y, alpha, color, fill, group,  
linetype, size



**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



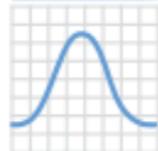
**f + geom\_violin(scale = "area")**, x, y, alpha, color,  
fill, group, linetype, size, weight

# Continuous x

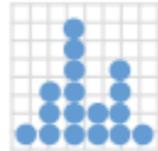
---

## ONE VARIABLE continuous

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



**c + geom\_density(kernel = "gaussian")**  
x, y, alpha, color, fill, group, linetype, size, weight



**c + geom\_dotplot()**  
x, y, alpha, color, fill



**c + geom\_histogram(binwidth = 5)**  
x, y, alpha, color, fill, group, linetype, size, weight

# 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

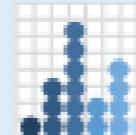
Use with x or y aesthetics (x shown here)

**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

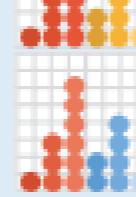
## COLOR AND FILL SCALES (CONTINUOUS)



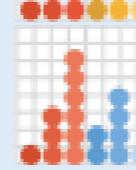
**o <- c + geom\_dotplot(aes(fill = ..x..))**



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



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



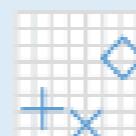
**o + scale\_fill\_gradient2(low="red", high="blue", mid = "white", midpoint = 25)**



**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

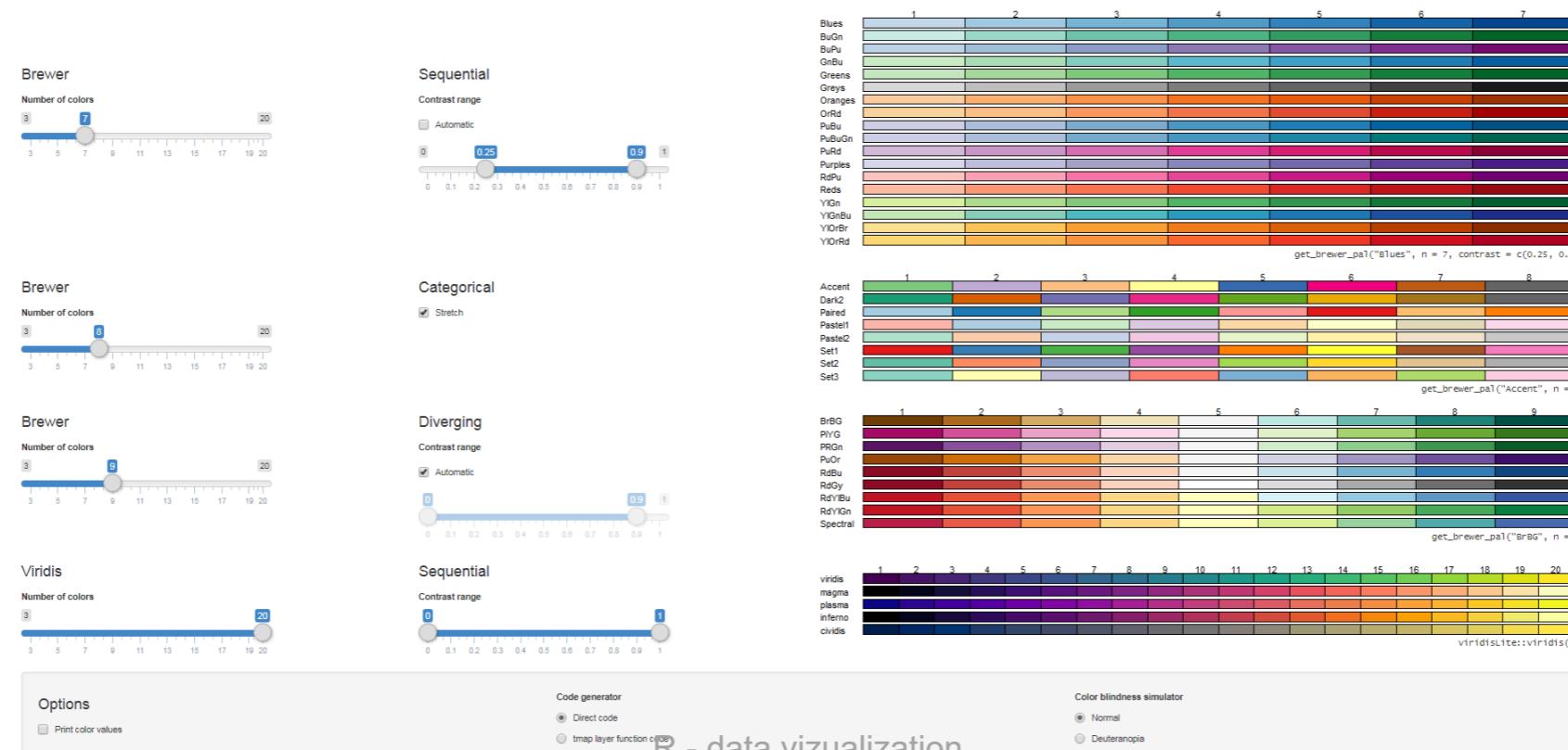


**p + scale\_radius(range = c(1,6))**

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

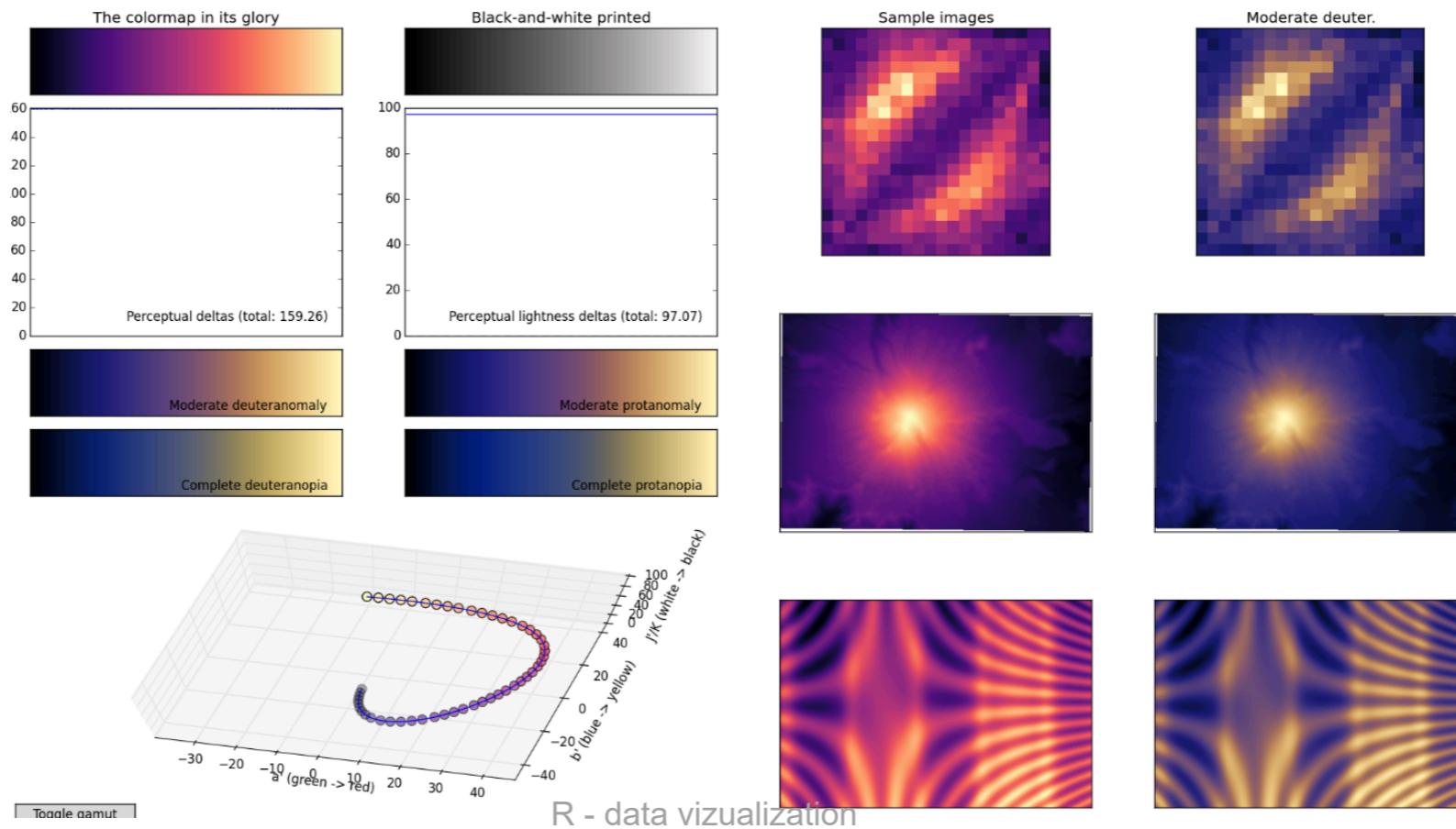
# Palettes

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

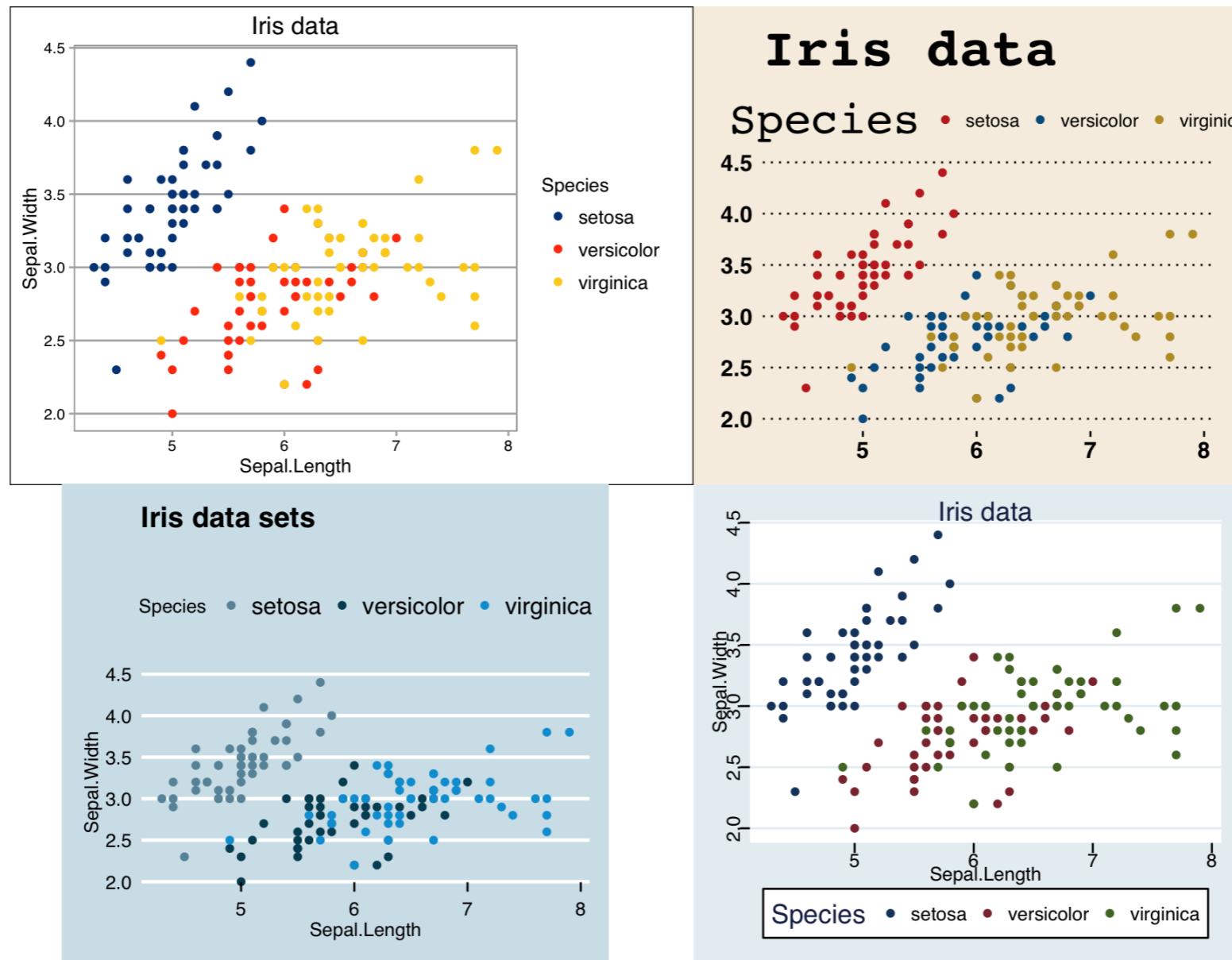


# Palettes

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



# Themes



# Extensions

---

## ggplot2 extensions - gallery

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



142 registered extensions available to explore

### ▼ Github stars

- Name
- Author
- Github stars

### Github stars ▾

Sort

Text Filter

### ▼

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

# Let's do a graph

# Your mission

---

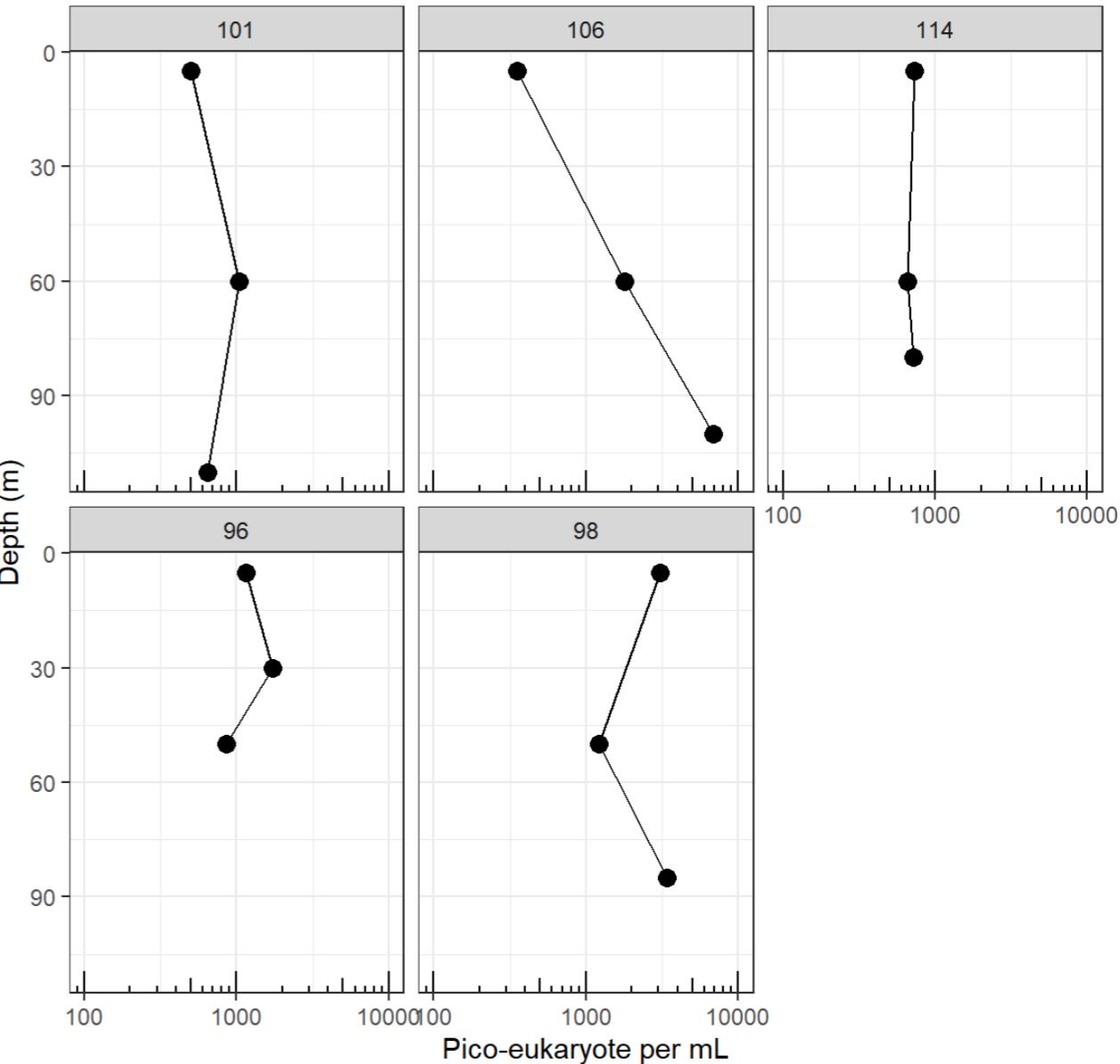
# Reproduce graph on right

- 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

Abundance of pico-eukaryotes per station on transect 2



# Recap

---

- 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**