

# Visualización de datos en R

Elena Quintero

13/01/2025

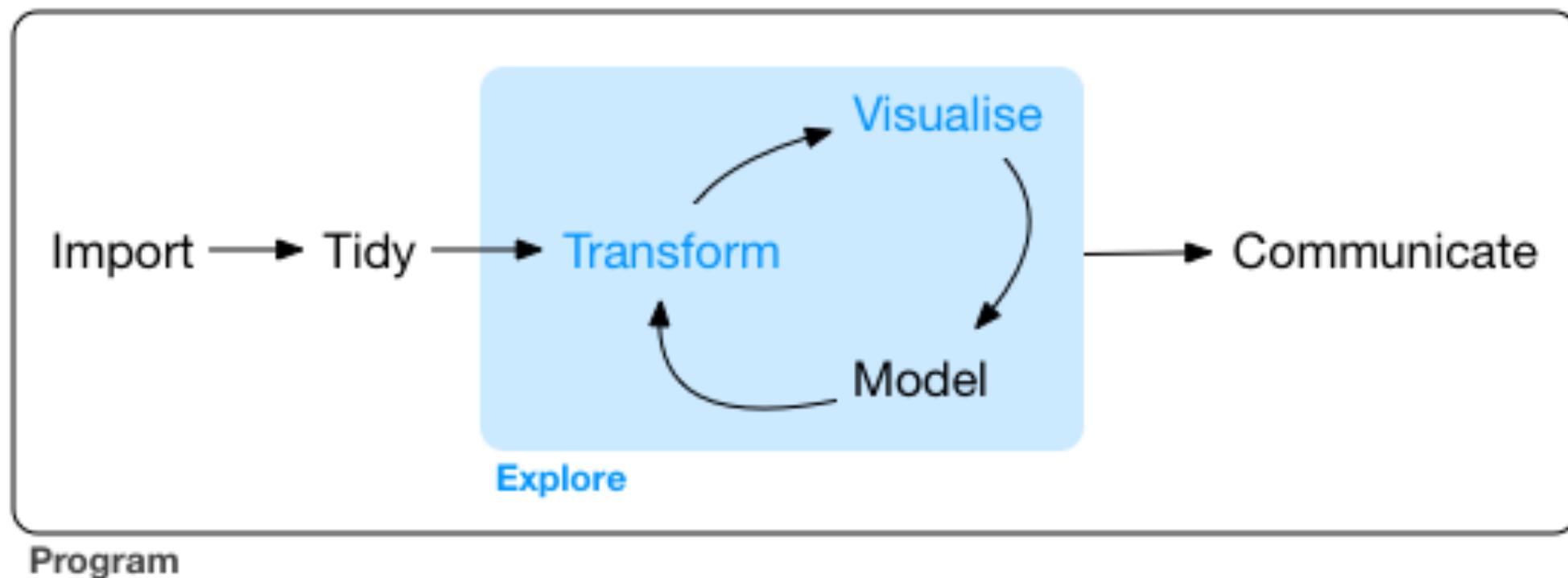
# Carpeta con material

<https://rstats-courses.github.io/CursoR-AEET-2025/materiales.html>

# Exploración de datos

La exploración de datos nos permite verificar su calidad, generar y probar hipótesis de forma rápida, identificando pistas prometedoras para analizar más a fondo luego.

La visualización de los datos es un buen comienzo, pero por sí sola no suele ser suficiente, ya que a menudo requiere transformar los datos previamente.



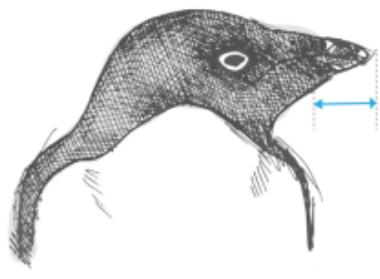
# Beneficios de usar ggplot

- Reproducible
- Consistencia gramática
- Muy flexible y permite controlar gran cantidad de detalles
- Fácil para uso básico
- Comunidad de usuarios activos

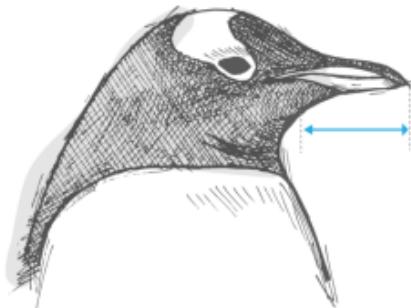
## Palmer Penguins Bill Length

Palmer Archipelago is a group of islands off the northwestern coast of the Antarctic Peninsula.  
The histograms show that females has shorter bills than males in every species

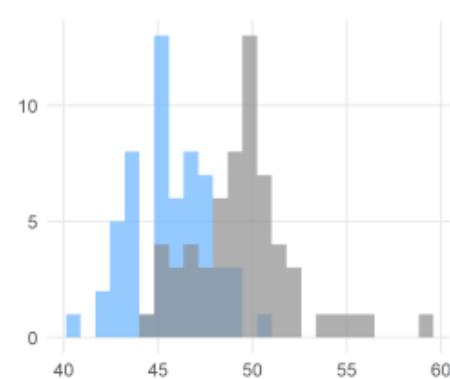
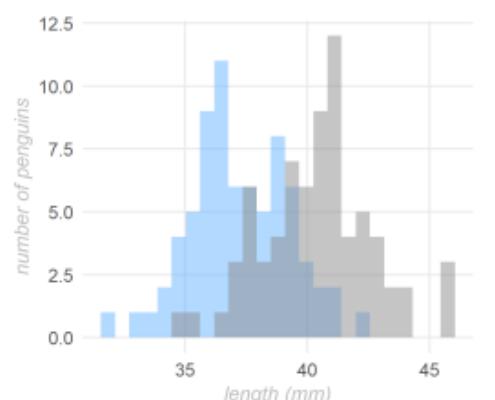
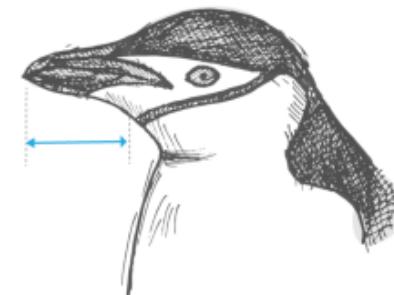
ADELIE



GENTOO



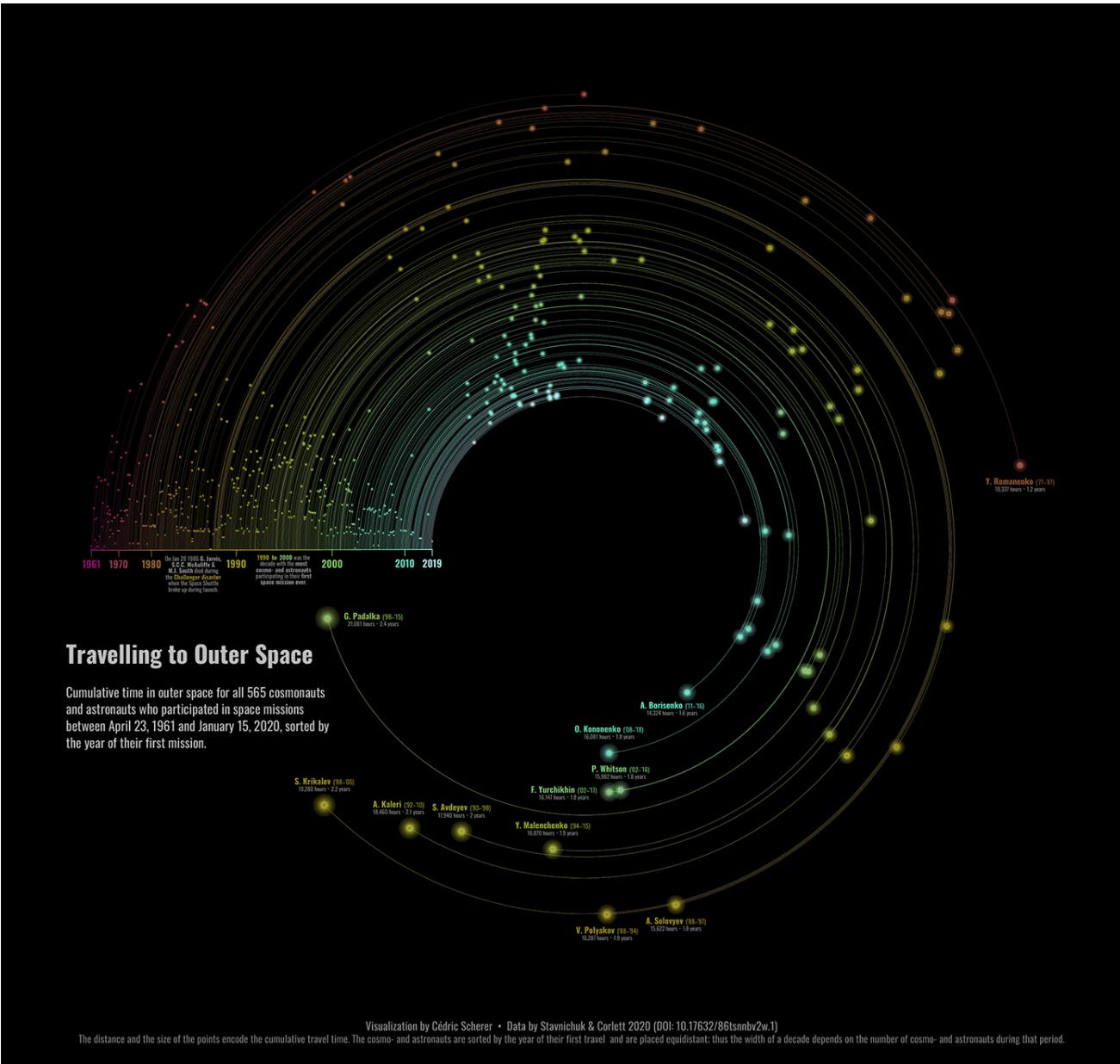
CHINSTRAP



female male

Visualization: Laura Navarro Soler | Data: Gorman, Williams & Fraser (2014)

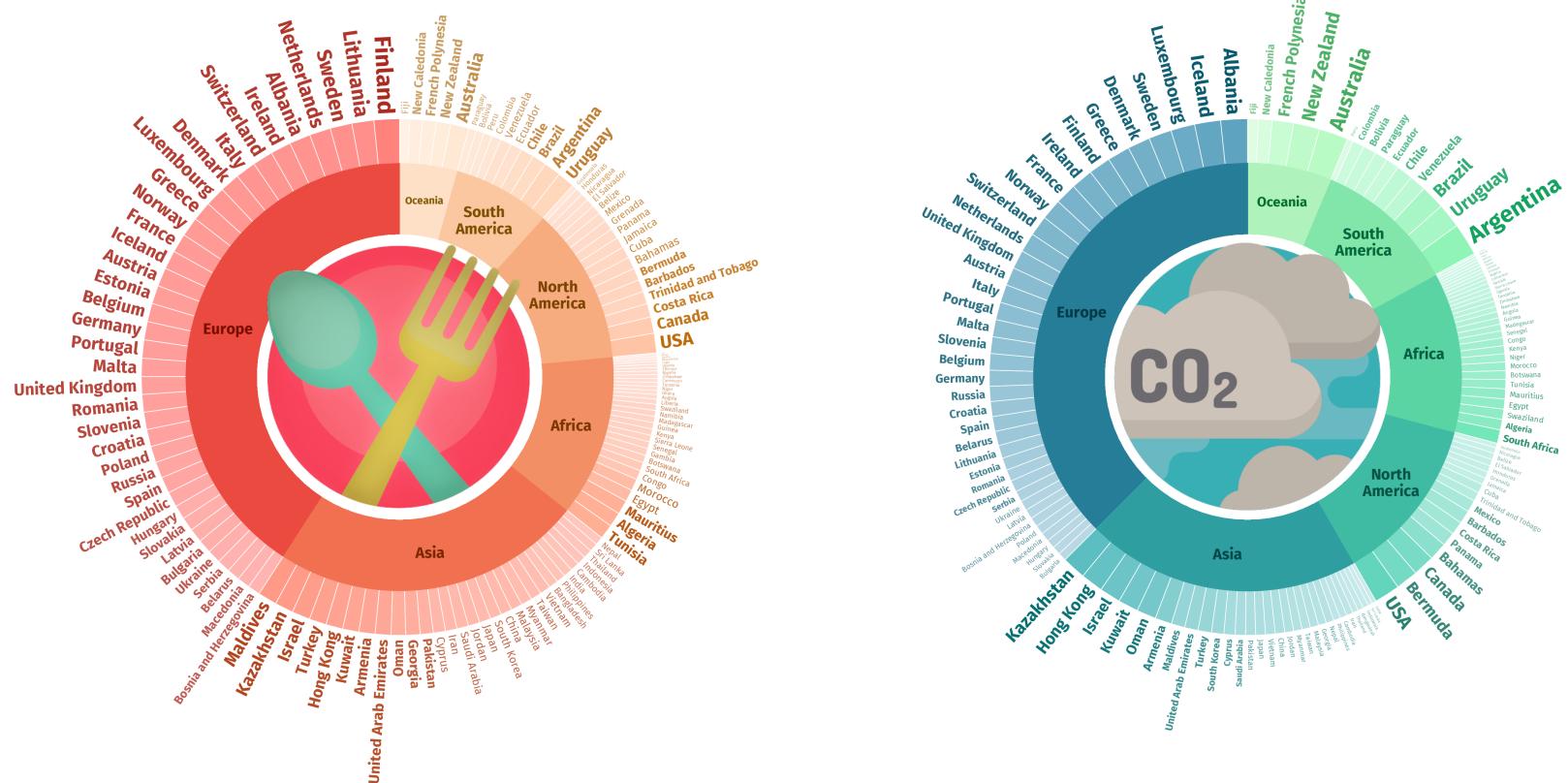
Laura Navarro | [code here](#)



Cédric Scherer | [code here](#)

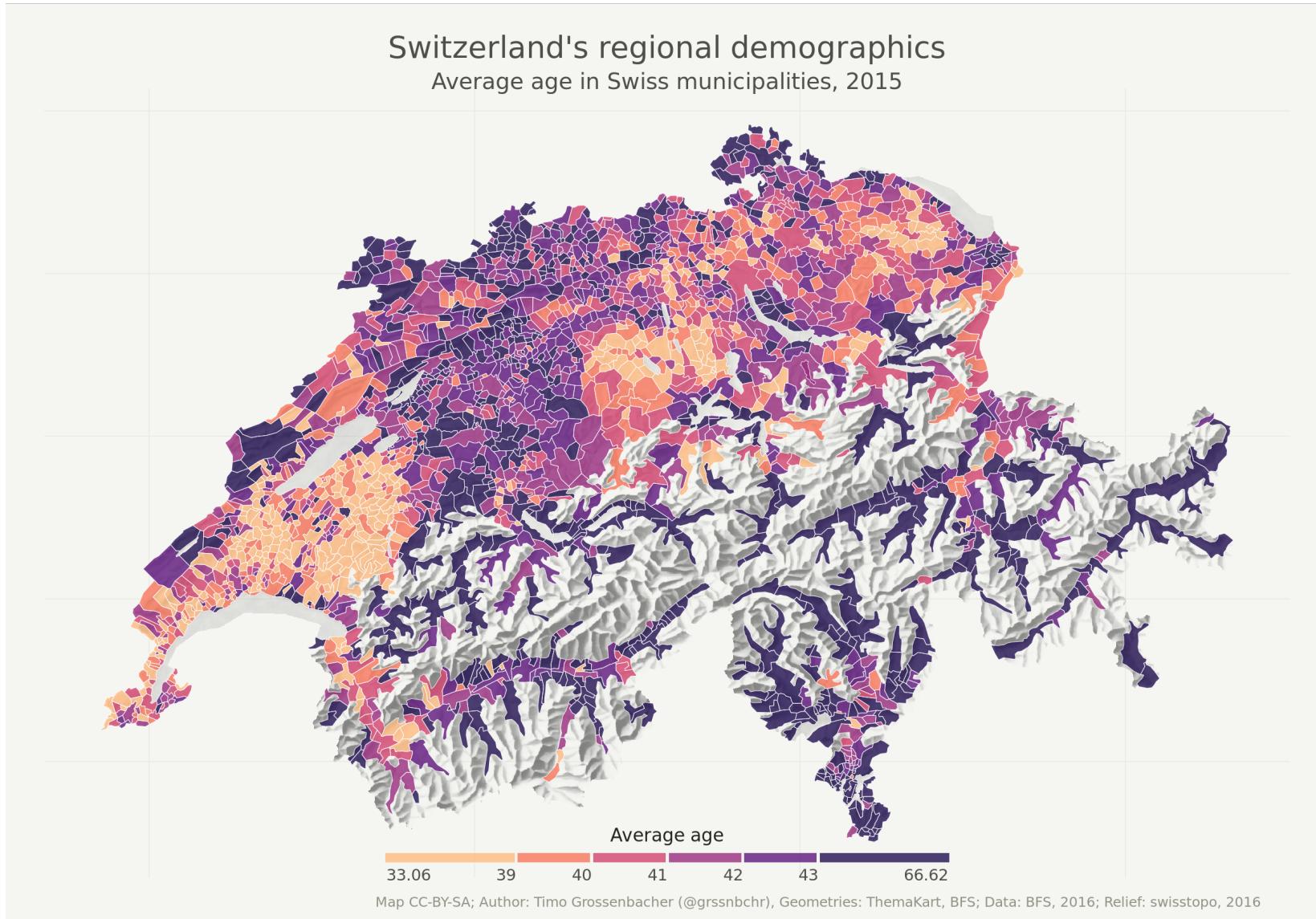
# Food Carbon Footprint Index 2018

Global comparison of different diets in terms of **Average Consumption** (kg/person/year) of both animal and non-animal products as well as **Carbon Emissions** (kg CO<sub>2</sub>/person/year) per continent and country. Font size and color intensity indicate each country's estimate with **countries printed in bold** belonging to the upper 50% of consumers and CO<sub>2</sub> emitters, respectively.



Visualization by Cédric Scherer • Data by Food and Agriculture Organization of the United Nations (FAO) via nuz • Icons by FreePik

# Cédric Scherer



Timo Grossenbacher | [code here](#)

# What do most people die from?

Cardiovascular Diseases



Cancers



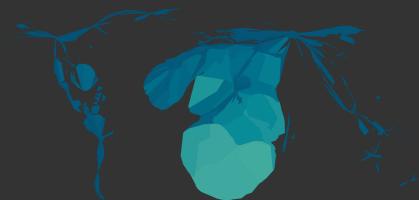
Diabetes



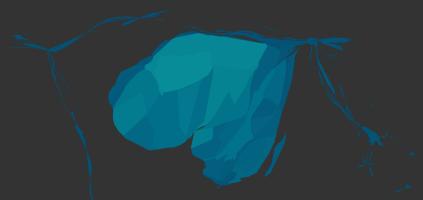
The leading causes of death across the world still vary significantly.

These cartograms show causes of deaths in 2016 that exceeded 20 percent of total deaths in at least 1 country.

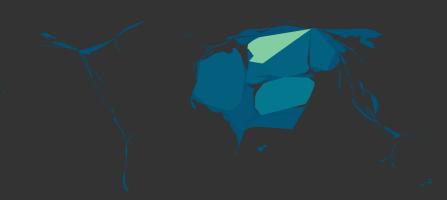
HIV Infections & Aids



Malaria Infections



War & Conflicts

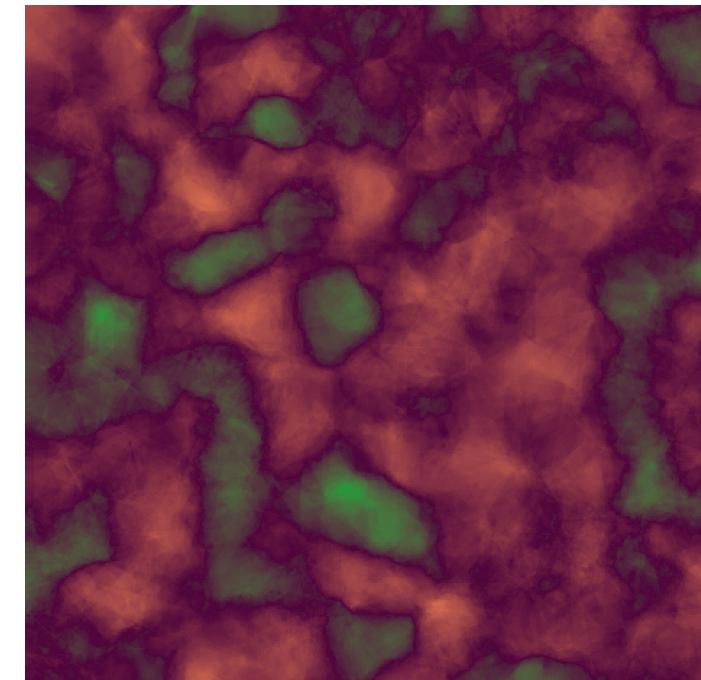
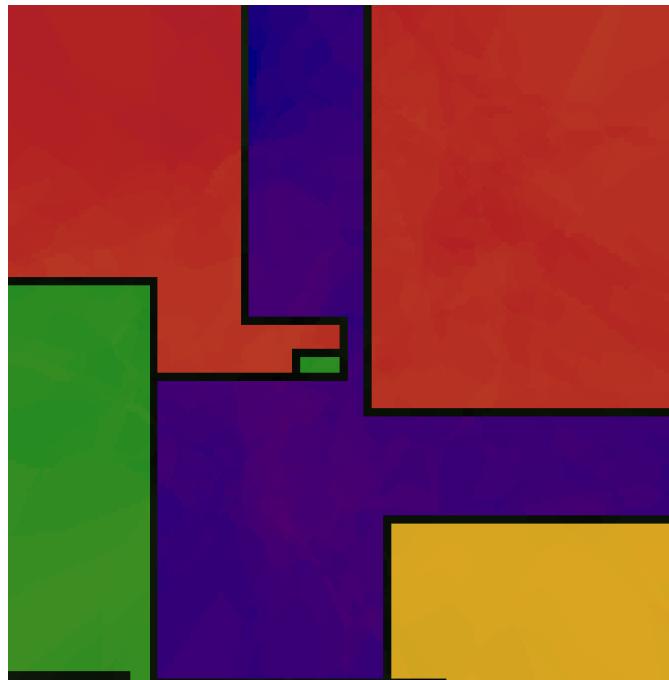
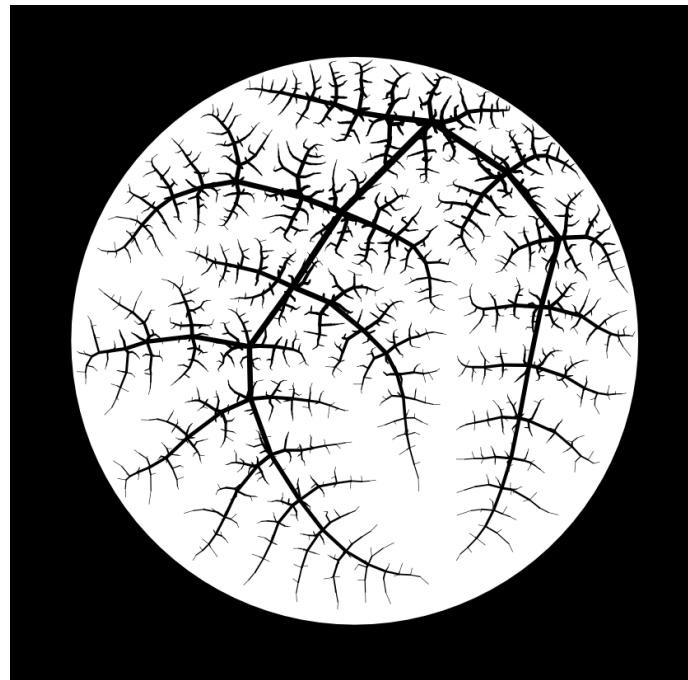


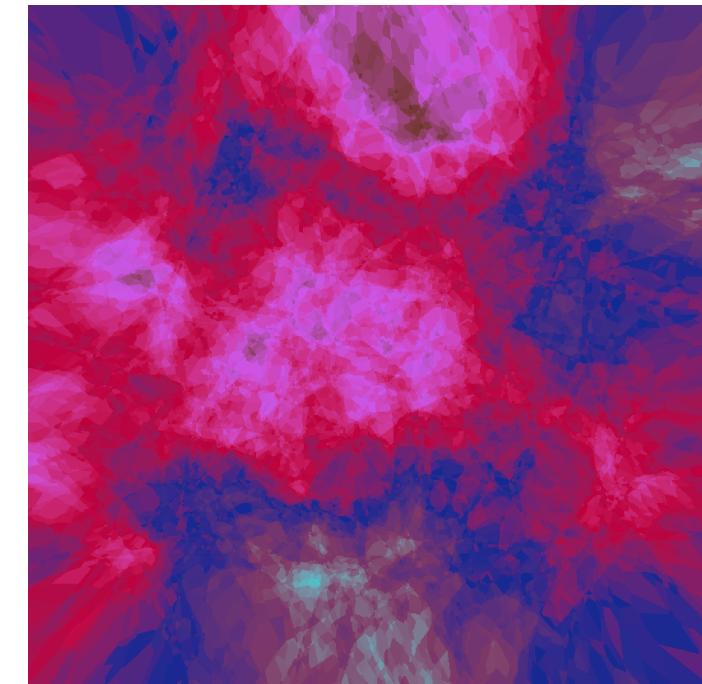
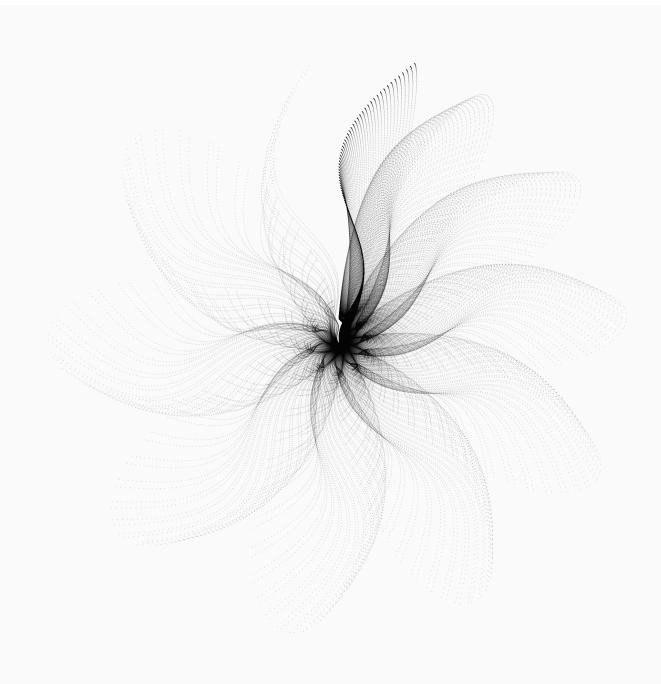
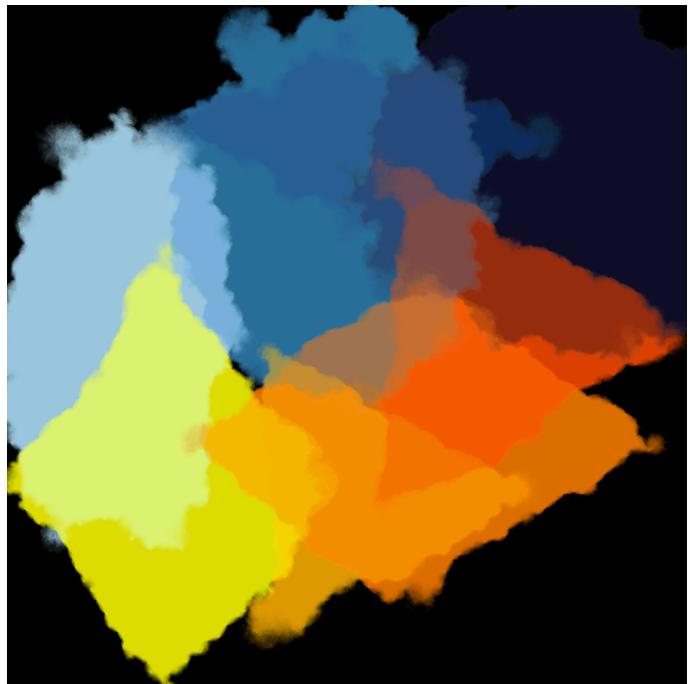
The data refers to the specific cause of death, which is distinguished from risk factors for death, such as air pollution, diet and other lifestyle factors.

Visualization by Cédric Scherer • Data by OurWorldInData.org

Cédric Scherer | [code here](#)

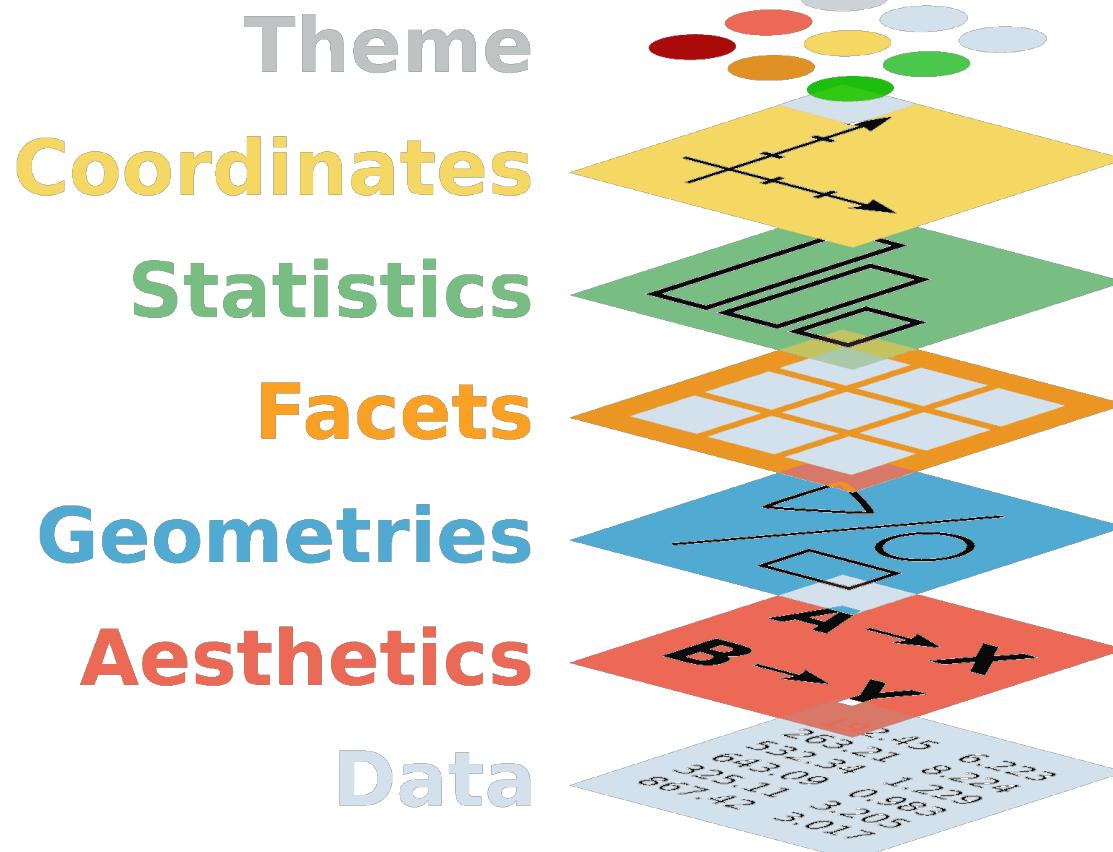
## aRtsy: Generative Art with R and ggplot2





# Gramática de ggplot

Basado en la gramática *layered grammar of graphics* - Bertin 1983, Wilkinson et al. 2005, Wickham 2010.



# Capas

- **Data** - dataset a graficar; necesita formato tidy data
- **Aesthetics** - describe cómo se asignan las variables del dataset (y sus escalas) a las propiedades visuales (x, y, size, colour, fill, group, sise, shape... )
- **Geometries** - determina la forma de representar los datos (geoms: points, lines, bars, boxplot... )
- Scales - maneja las escalas de los aesthetics (x & y format, colors continuous or discrete, sizes, shapes... )
- Facets - crea subplots (facet\_wrap or facet\_grid... )
- Themes - apariencia general del gráfico, no ligada a los datos (title, x.axis.text, legend... )
- Statistics - resume los datos con estadísticos. Muchas veces ya va implícito en el “geom” (stats: count, density, bins, means, density... )
- Coordinate system - determina el sistema de coordenadas a usar en los ejes (cartesian, polar, map projections... )

\*Capas necesarias

# Dataset

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DOI: 10.1002/ecy.4463

DATA PAPER



## Co-mast: Harmonized seed production data for woody plants across US long-term research sites

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Rebecca S. Snell<sup>16</sup> | Jess K. Zimmerman<sup>17</sup> | Johannes M. H. Knops<sup>18</sup> |  
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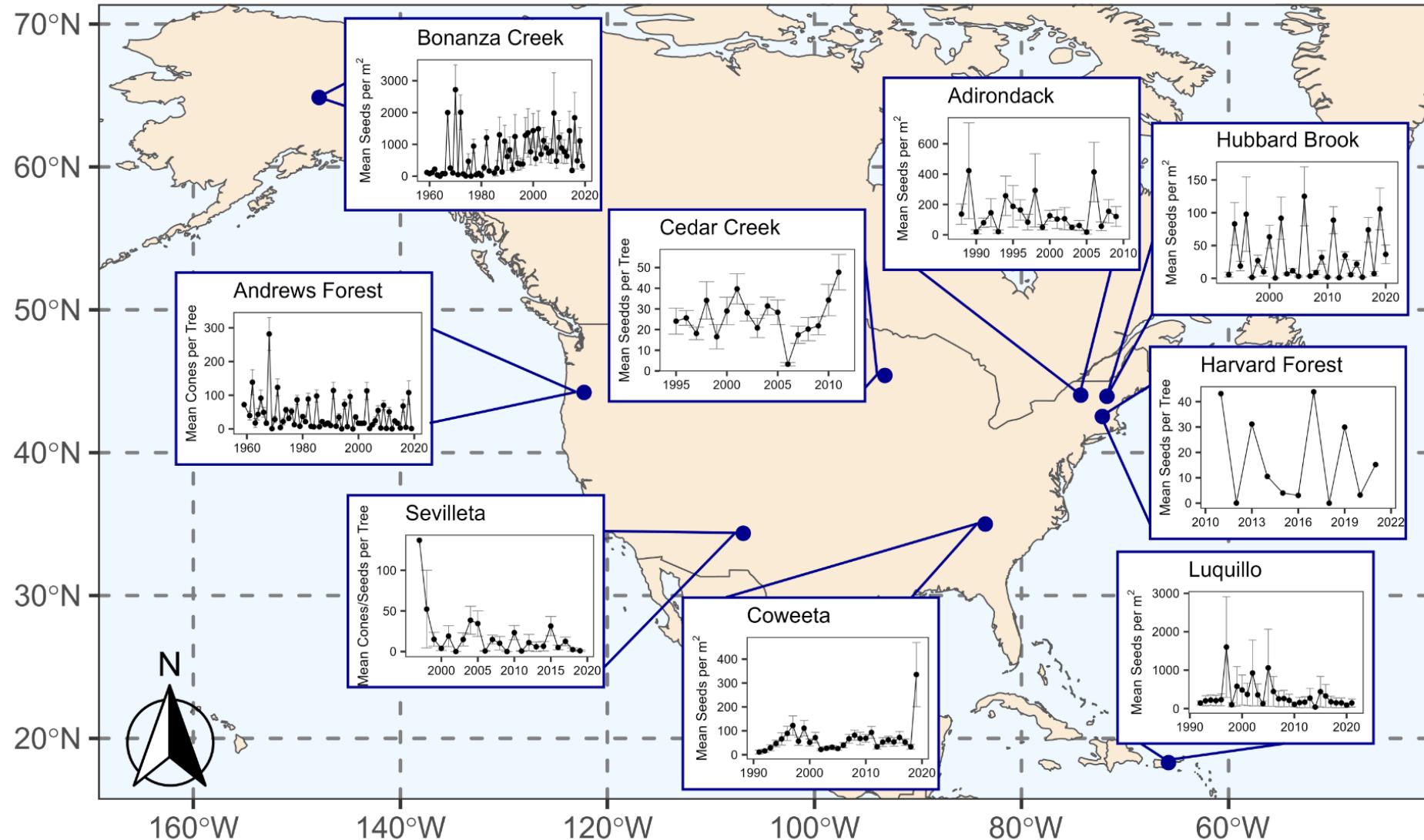
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### Abstract

Plants display a range of temporal patterns of inter-annual reproduction, from relatively constant seed production to “mast seeding,” the synchronized and highly variable interannual seed production of plants within a population. Previous efforts have compiled global records of seed production in long-lived

<https://doi.org/10.1002/ecy.4463>

# Dataset



# Cargar paquetes

```
library(here)
library(tidyverse)
library(tidylog)
```

# Leer datos

```
dt <- read_csv(here("data/clean_data.csv"))
```

# Graficar

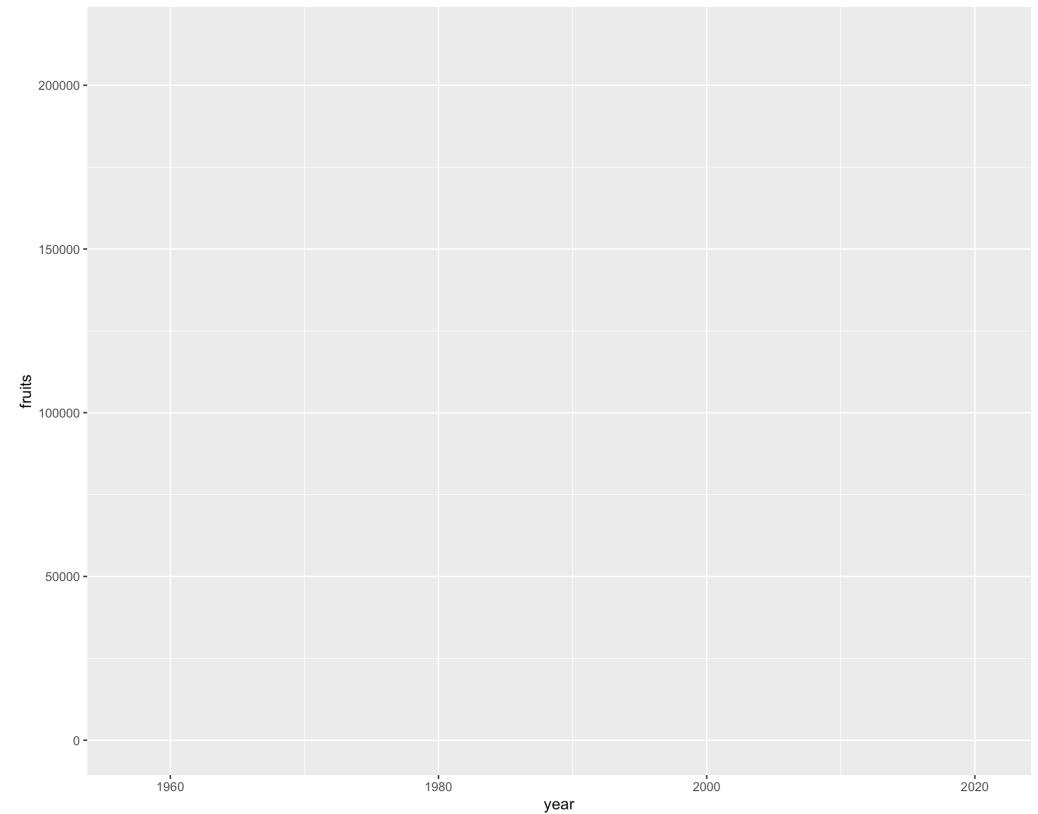
## 1. Indicar el dataset

```
ggplot(data = dt)
```

# Graficar

## 2. Definir los aesthetics (variables a visualizar)

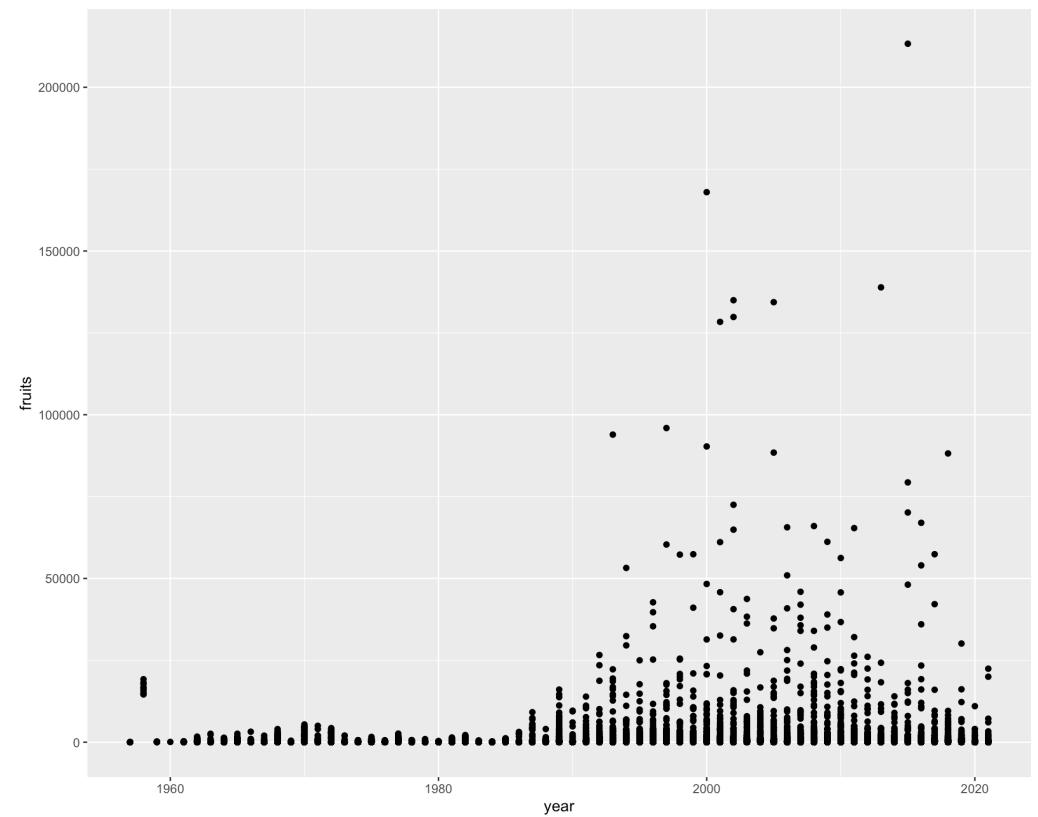
```
ggplot(data = dt,  
       aes(x = year, y = fruits))
```



# Graficar

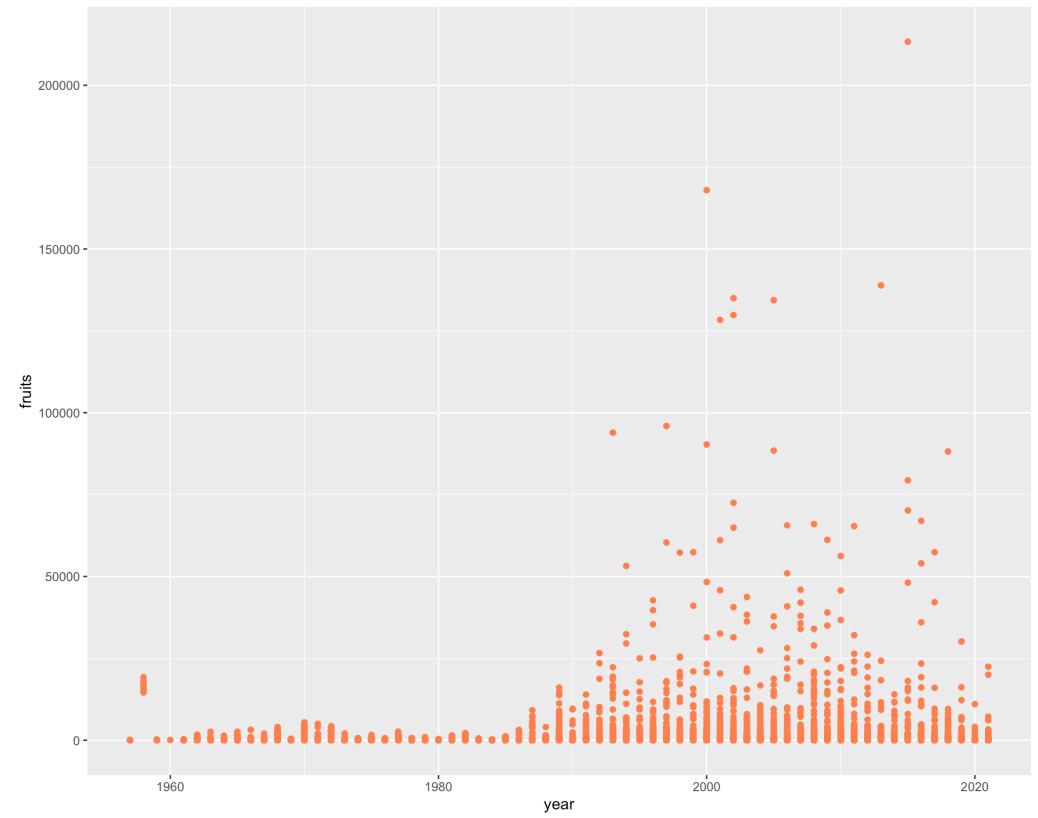
## 3. Determinar la forma de representar los datos con GEOMs

```
ggplot(data = dt,  
       aes(x = year, y = fruits)) +  
  geom_point()
```



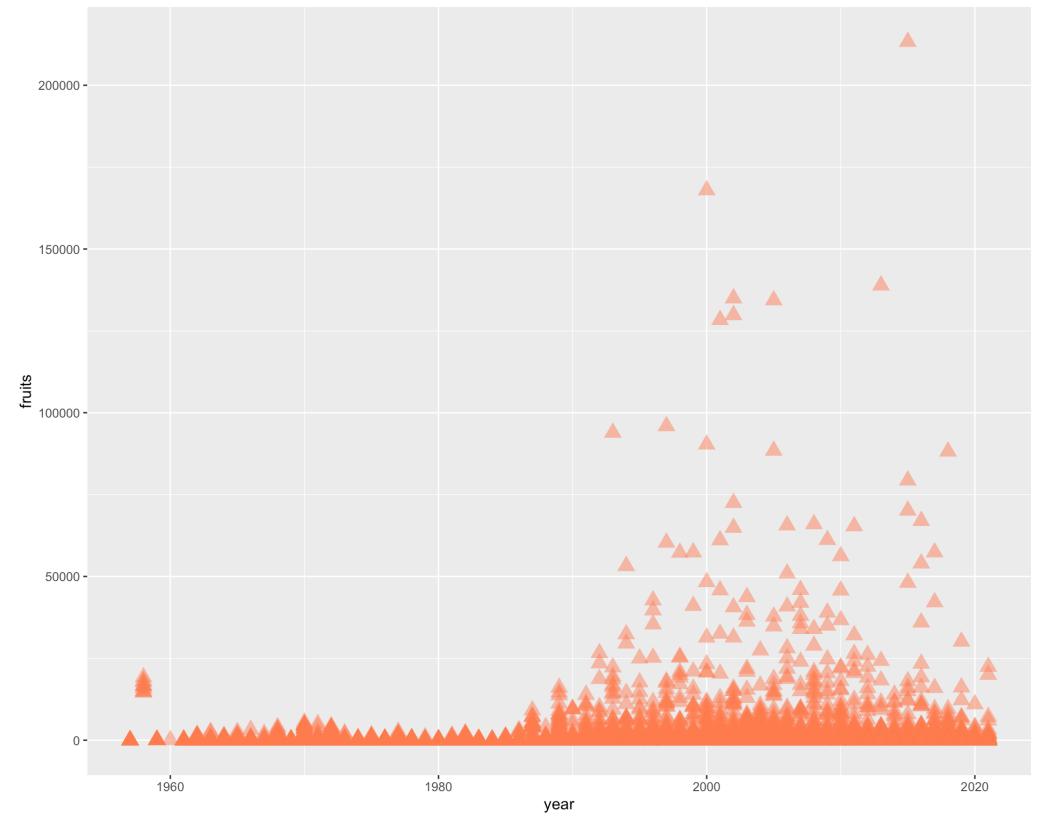
# Ajustar detalles de visualización

```
ggplot(dt,  
       aes(x = year, y = fruits)) +  
  geom_point(color = "coral")
```



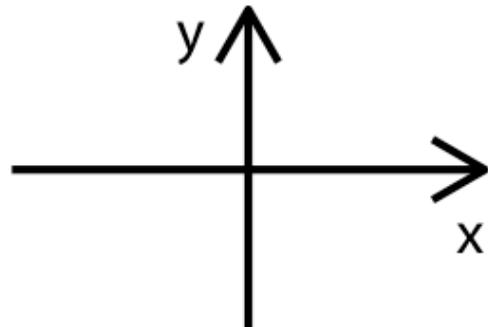
# Ajustar detalles de visualización

```
ggplot(dt,  
       aes(x = year, y = fruits)) +  
  geom_point(color = "coral",  
             shape = "triangle",  
             size = 4,  
             alpha = 0.5)
```

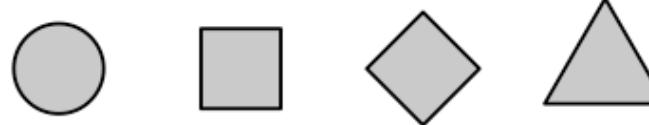


# Valores de los aesthetics (aes):

position



shape



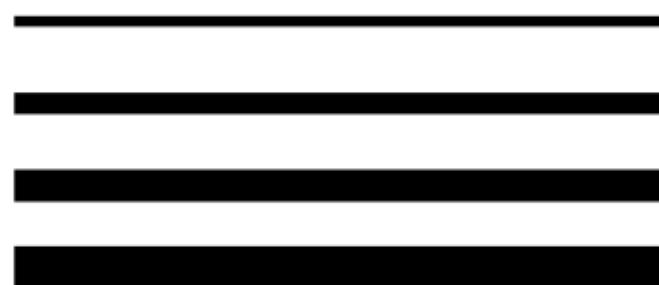
size



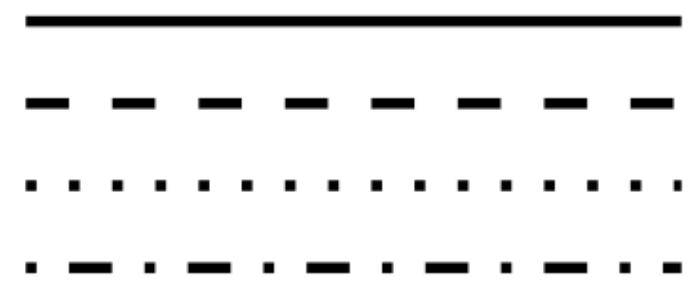
color



line width

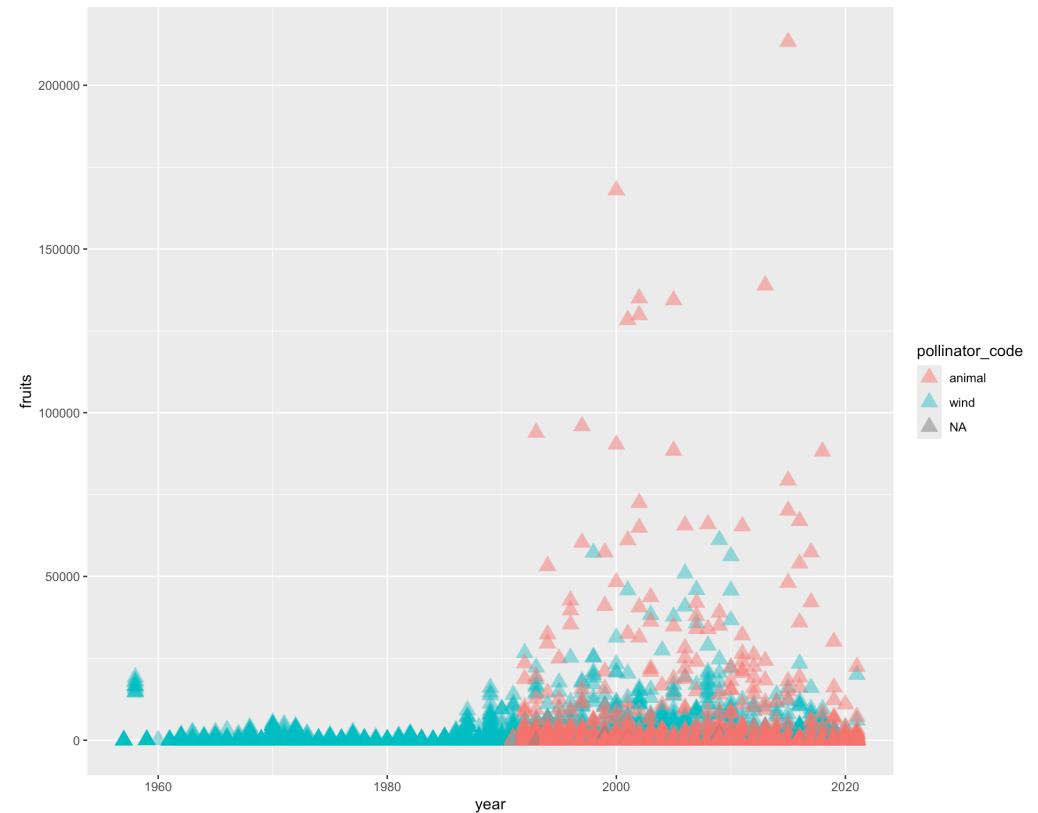


line type



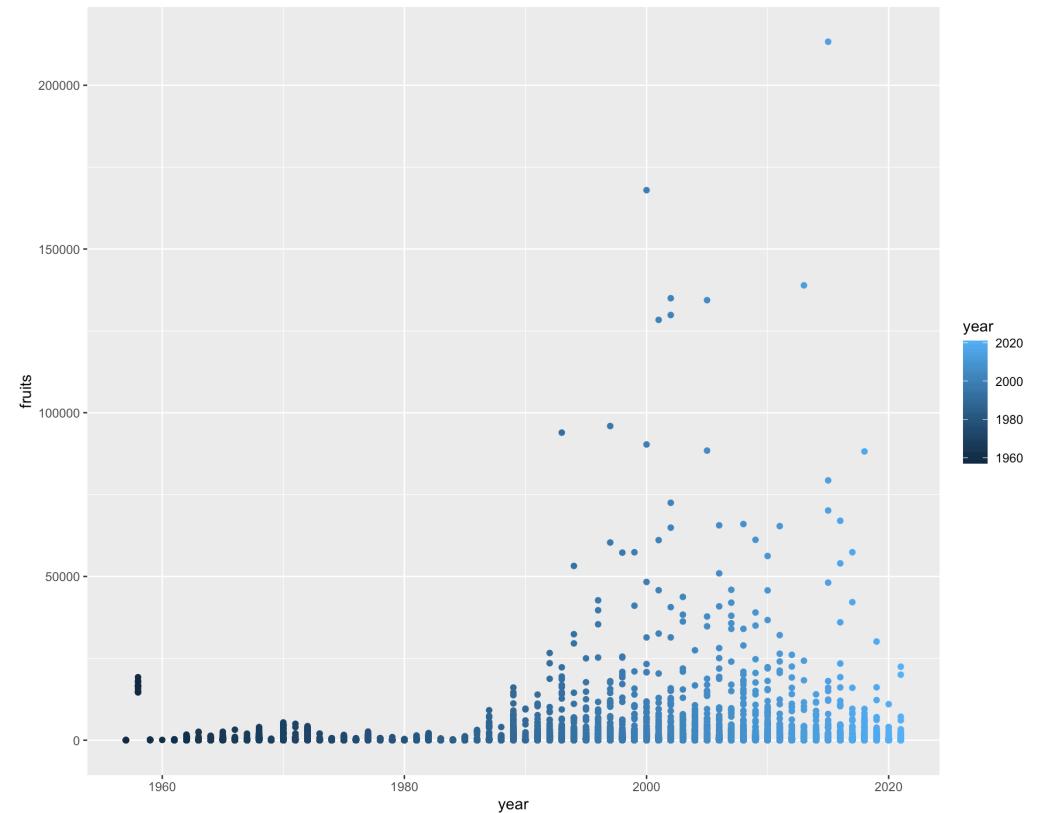
# Ajustar detalles de visualización

```
ggplot(dt,
       aes(x = year, y = fruits,
           color = pollinator_code)) +
  geom_point(shape = "triangle",
             size = 4,
             alpha = 0.5)
```



# Ajustar detalles de visualización

```
ggplot(dt,  
       aes(x = year, y = fruits,  
            color = year)) +  
  geom_point()
```



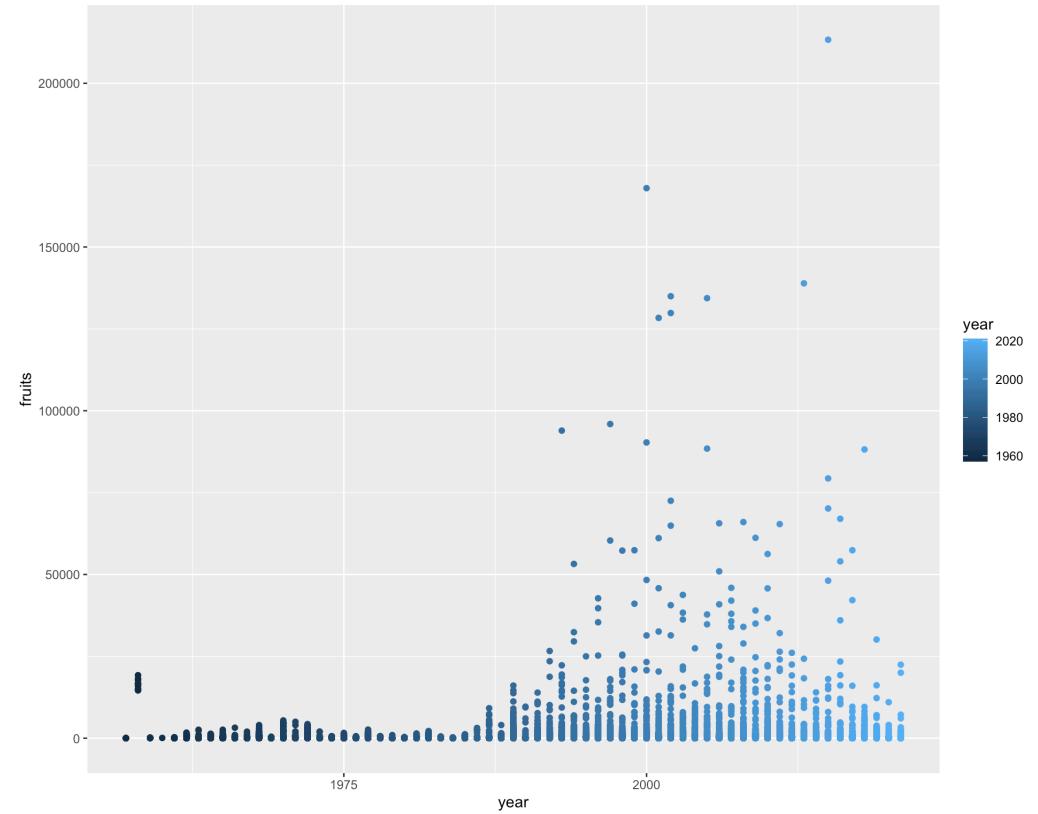
# Escalas

Van referidas a las variables o aesthetics (aes):

- X e Y
- Tamaño de punto (size), de linea (linewidth)
- Forma de punto (shape) y de linea (linetype)
- Color y relleno (colour, fill)

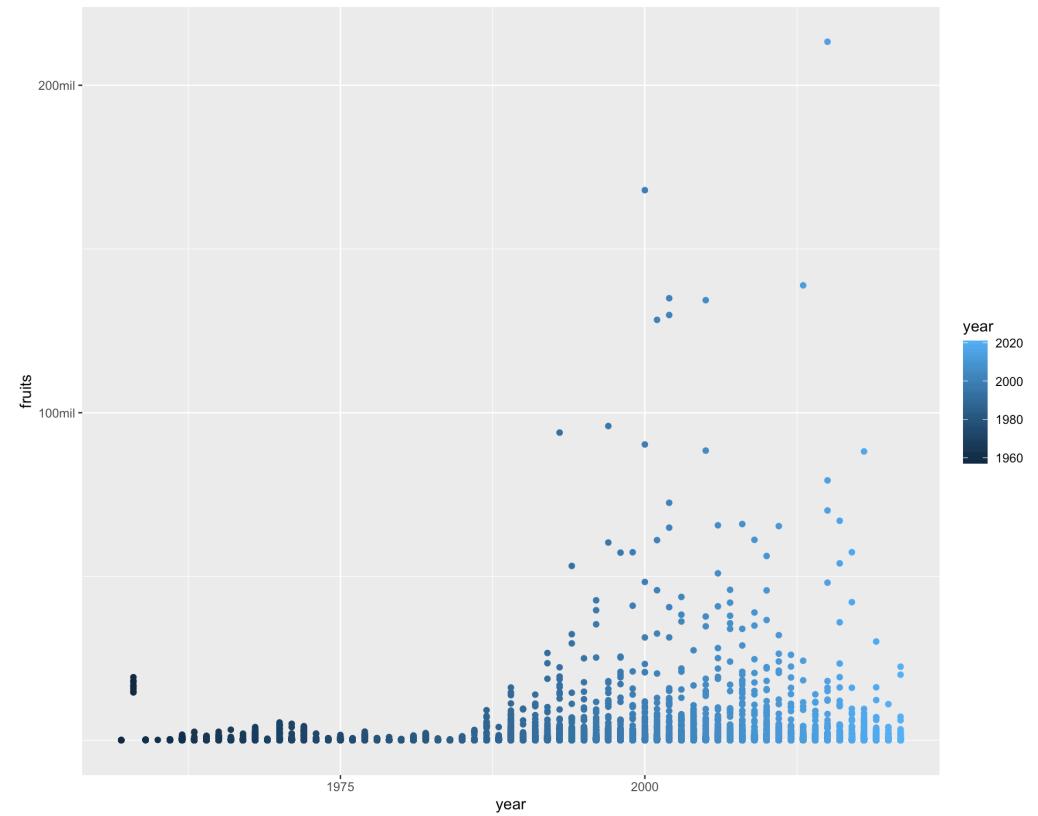
# Escalas - X, Y

```
ggplot(dt,
       aes(x = year, y = fruits,
            color = year)) +
  geom_point() +
  scale_x_continuous(n.breaks = 3)
```



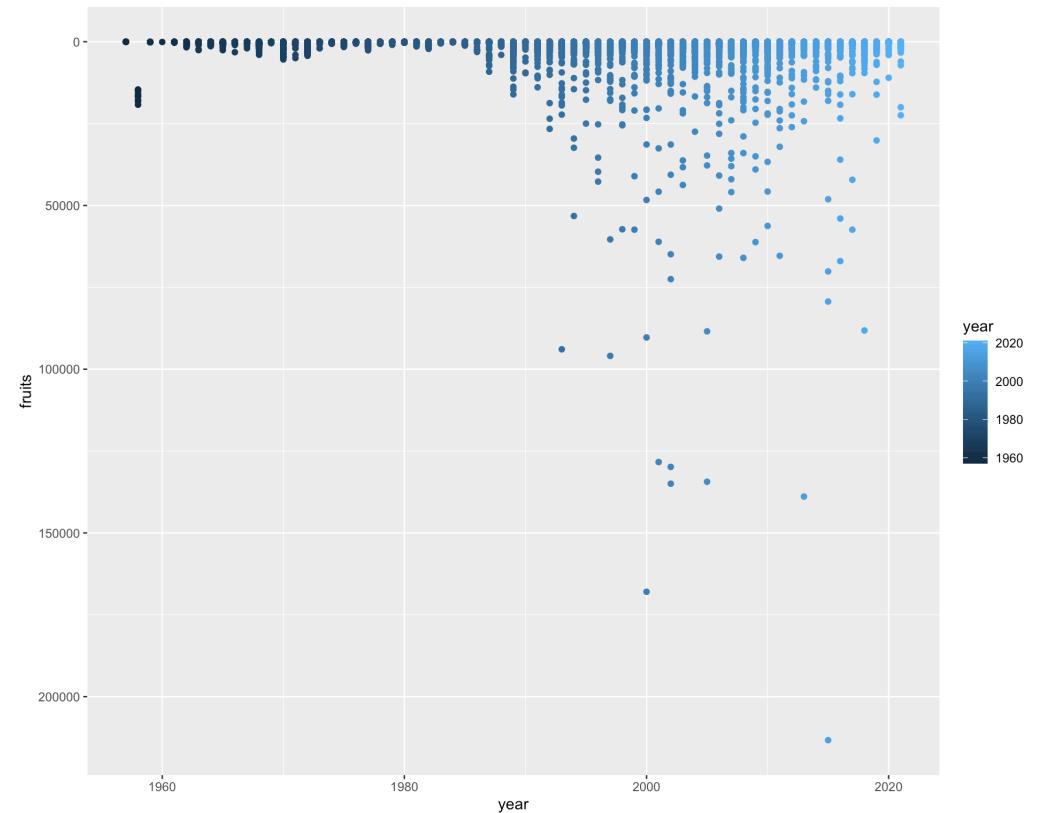
# Escalas - X, Y

```
ggplot(dt,
       aes(x = year, y = fruits,
            color = year)) +
  geom_point() +
  scale_x_continuous(n.breaks = 3) +
  scale_y_continuous(
    breaks = c(100000, 200000),
    labels = c("100mil",
              "200mil"))
```



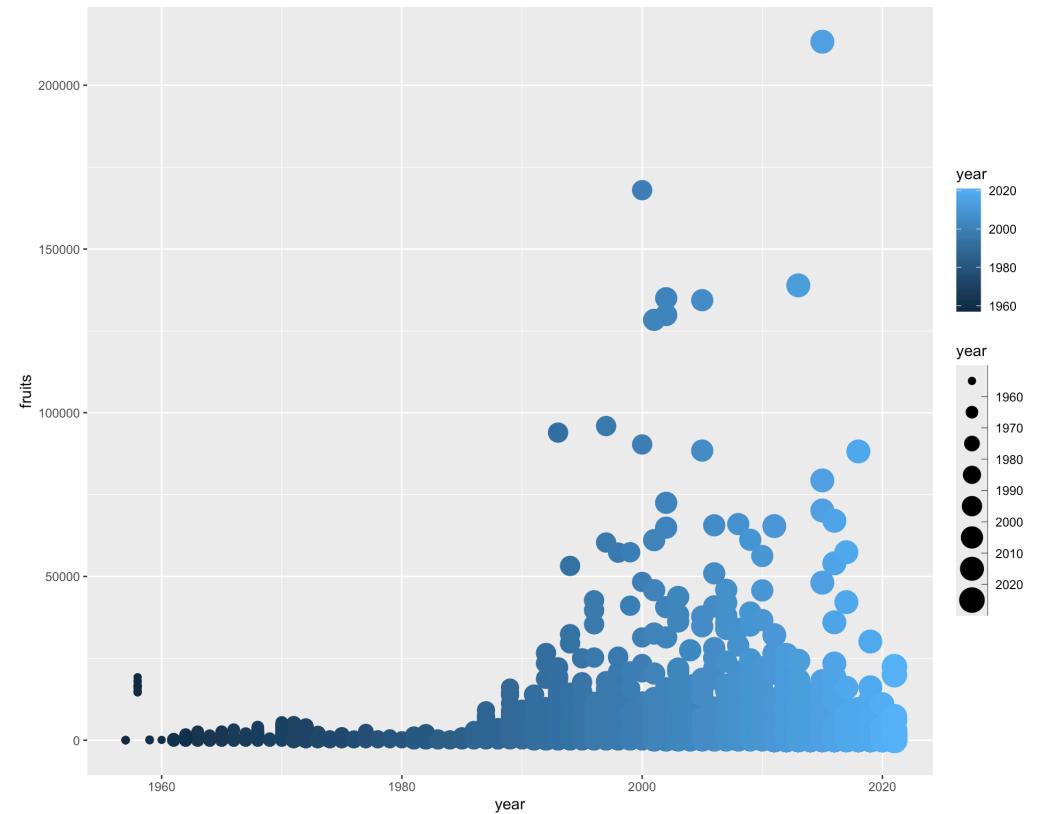
# Escalas - X, Y

```
ggplot(dt,
       aes(x = year, y = fruits,
            color = year)) +
  geom_point() +
  scale_y_reverse()
```



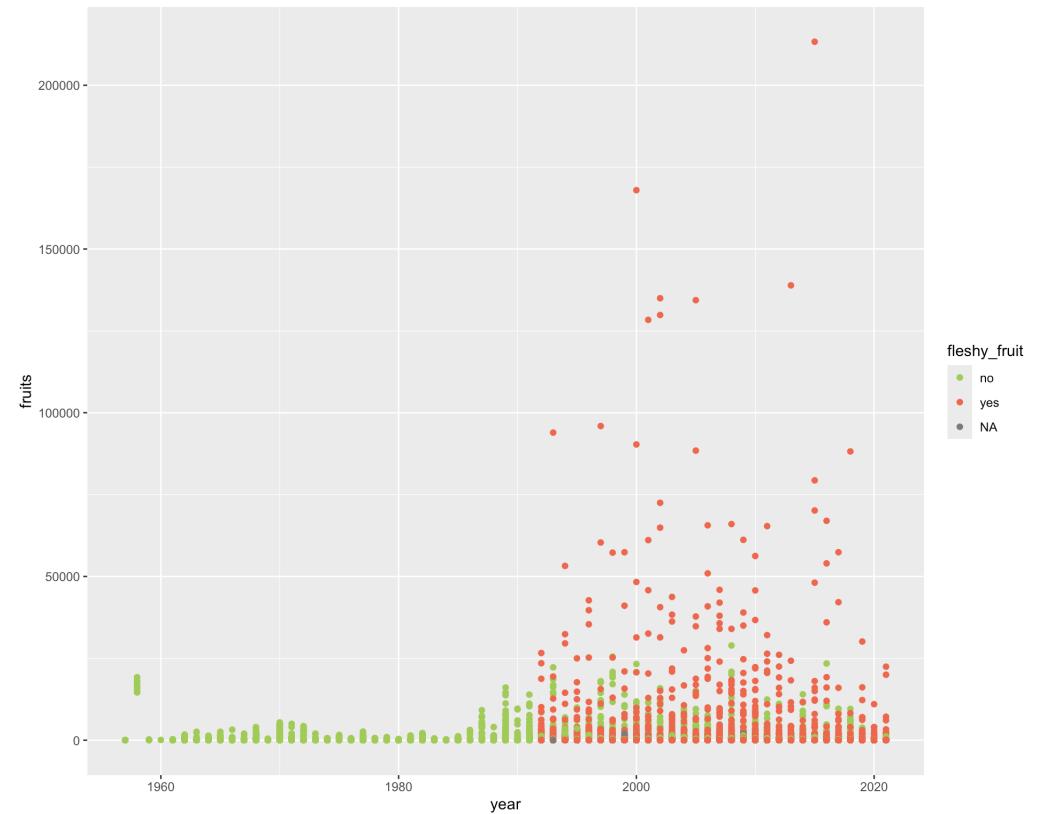
# Escalas - Tamaño

```
ggplot(dt,
       aes(x = year, y = fruits,
           color = year,
           size = year)) +
  geom_point() +
  scale_size_binned(range = c(0.1, 8),
                     n.breaks = 10)
```



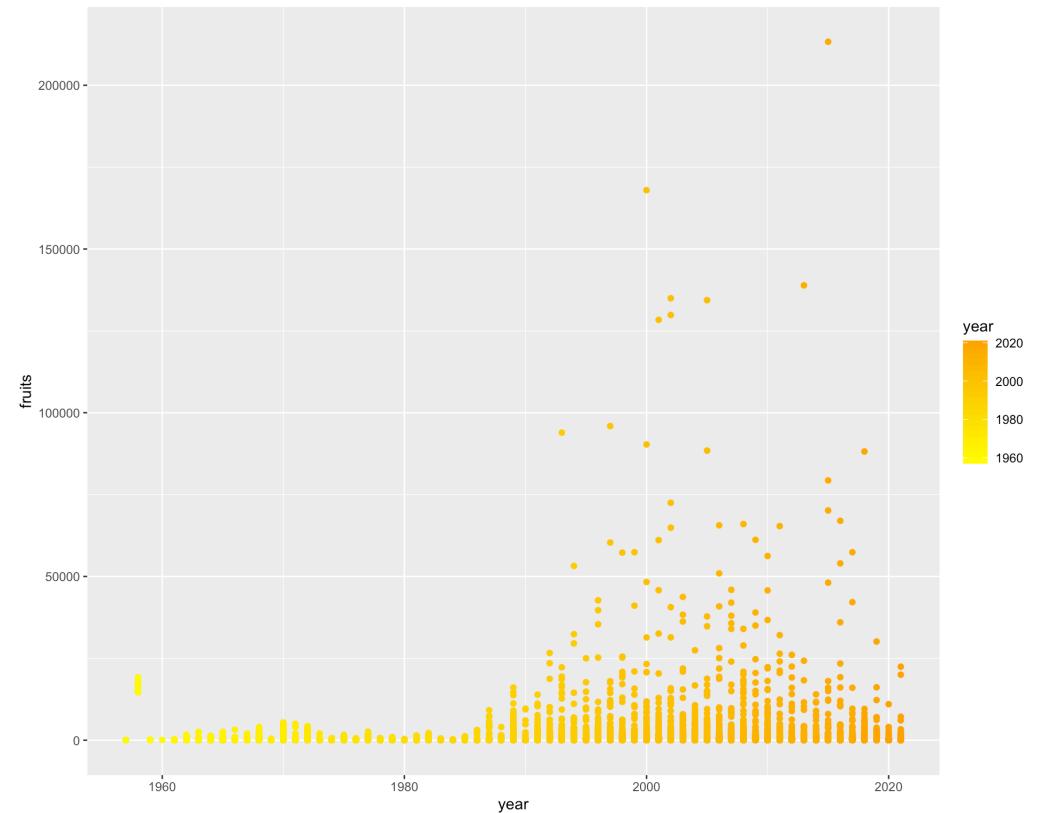
# Escalas - Color (discreto)

```
ggplot(dt,
       aes(x = year, y = fruits,
            color = fleshy_fruit)) +
  geom_point() +
  scale_color_manual(
    values = c("darkolivegreen3",
              "coral2"))
```



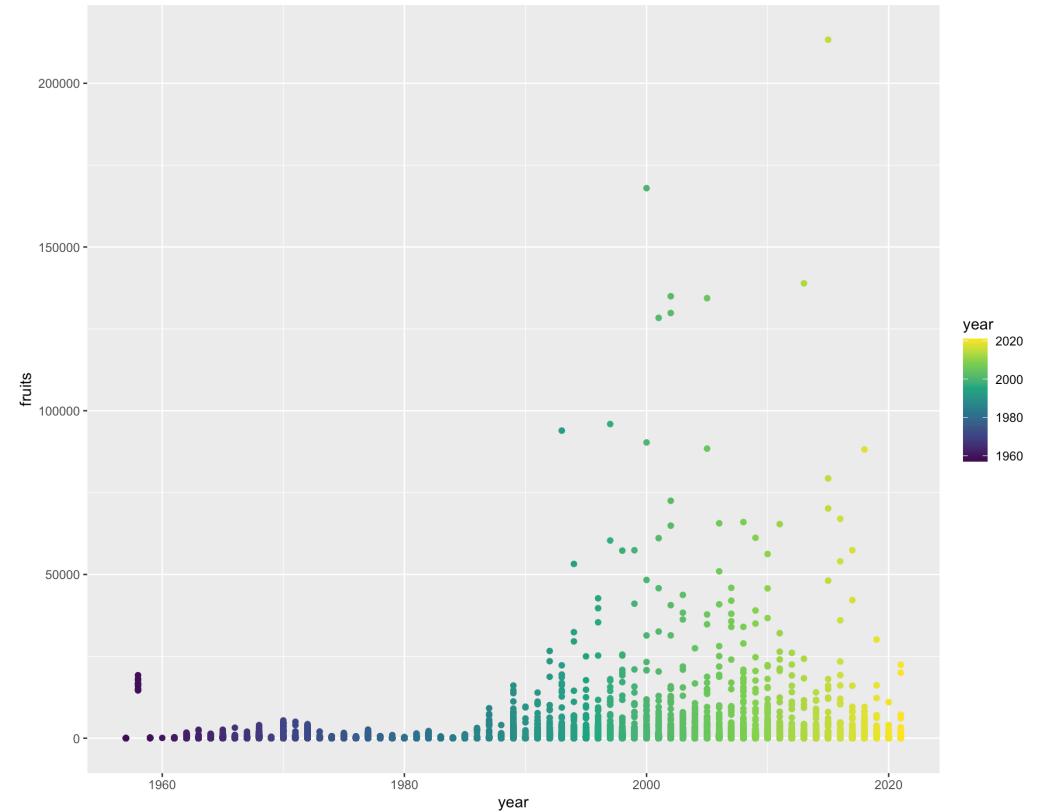
# Escalas - Color (continuo)

```
ggplot(dt,
       aes(x = year, y = fruits,
            color = year)) +
  geom_point() +
  scale_color_gradient(
    low = "yellow", high = "orange")
```



# Escalas - Color

```
ggplot(dt,
       aes(x = year, y = fruits,
            color = year)) +
  geom_point() +
  scale_color_viridis_c()
```



# Escalas - Color

Paletas de color:

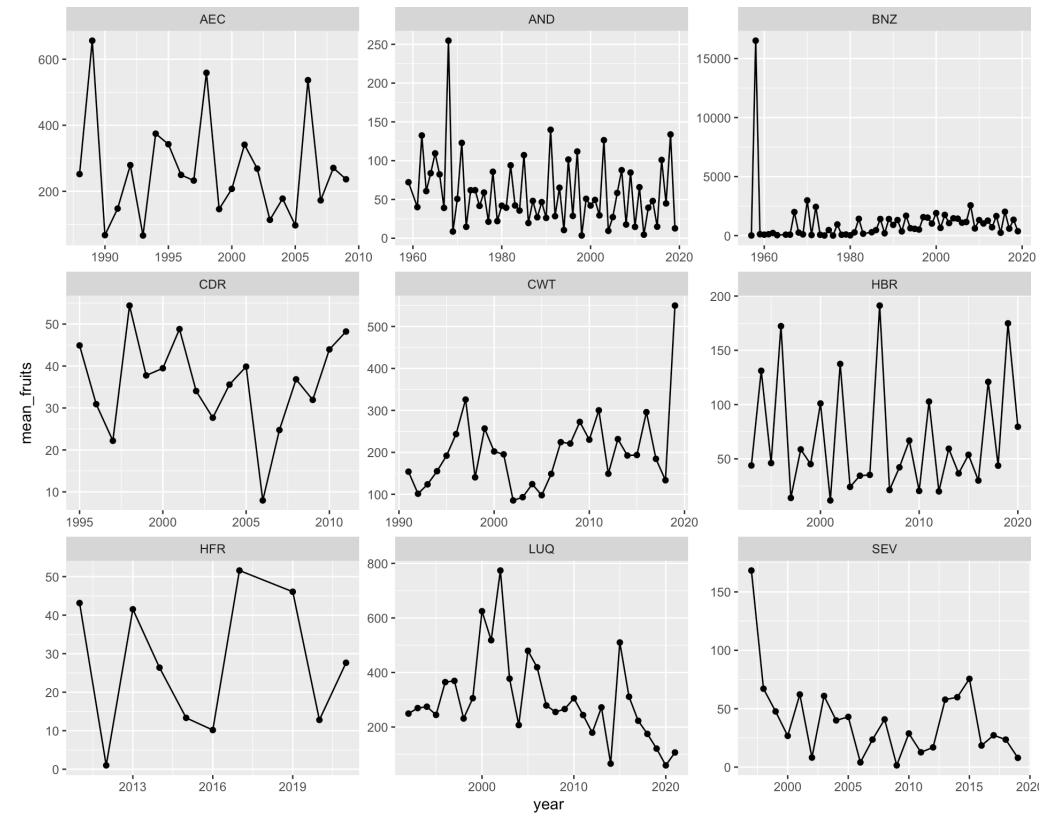
- <https://r-charts.com/color-palettes/>
- library(paletter) - <https://github.com/EmilHvitfeldt/paletteer>
- <https://r-graph-gallery.com/color-palette-finder>
- <https://medialab.github.io/iwanthue/>
- <https://projects.susielu.com/viz-palette>
- <https://colorbrewer2.org/#type=sequential&scheme=YIGnBu&n=3>

# Facets

- `facet_wrap()`
- `facet_grid()`

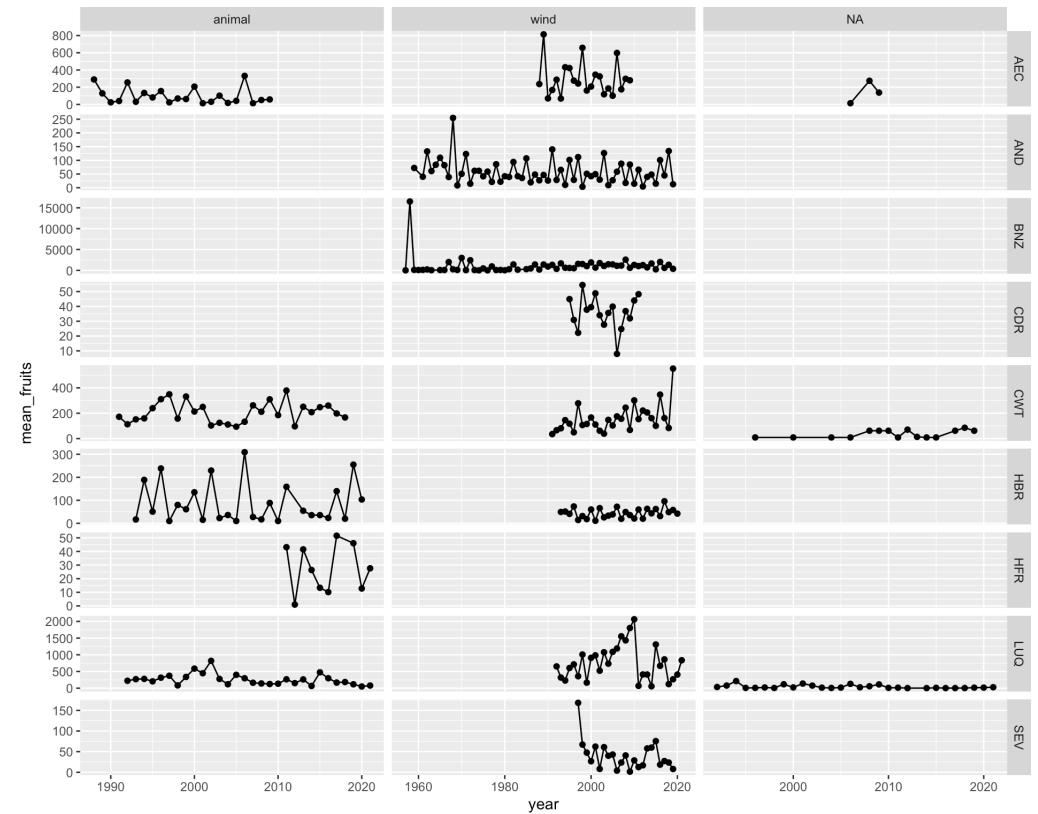
# Facets

```
dt |>  
  group_by(year, site) |>  
  summarise(mean_fruits = mean(fruits,  
                                na.rm = TRUE)) |>  
  ggplot(aes(x = year,  
             y = mean_fruits)) +  
  geom_point() +  
  geom_line() +  
  facet_wrap(~site, scales = "free")
```



# Facets

```
dt |>  
  group_by(year, site, pollinator_code)  
  summarise(mean_fruits = mean(fruits,  
                                na.rm = TRUE)) |>  
  ggplot(aes(x = year,  
             y = mean_fruits)) +  
  geom_point() +  
  geom_line() +  
  facet_grid(site~pollinator_code,  
             scales = "free")
```



# Tipos de Geoms

```
[1] "geom_abline"           "geom_area"          "geom_bar"  
[4] "geom_bin_2d"          "geom_bin2d"         "geom_blank"  
[7] "geom_boxplot"         "geom_col"           "geom_contour"  
[10] "geom_contour_filled"  "geom_count"         "geom_crossbar"  
[13] "geom_curve"           "geom_density"        "geom_density_2d"  
[16] "geom_density_2d_filled" "geom_density2d"    "geom_density2d_filled"  
[19] "geom_dotplot"         "geom_errorbar"      "geom_errorbarh"  
[22] "geom_freqpoly"        "geom_function"     "geom_hex"  
[25] "geom_histogram"       "geom_hline"         "geom_jitter"  
[28] "geom_label"           "geom_line"          "geom_linerange"  
[31] "geom_map"              "geom_path"          "geom_point"  
[34] "geom_pointrange"      "geom_polygon"       "geom_qq"  
[37] "geom_qq_line"         "geom_quantile"     "geom_raster"
```

# Tipos de Geoms

## Geoms

Use a geom function to represent data points, use the geom's aesthetic properties to represent variables.  
Each function returns a layer.

### GRAPHICAL PRIMITIVES

```
a <- ggplot(economics, aes(date, unemploy))
b <- ggplot(seals, aes(x = long, y = lat))
```

- a + geom\_blank()** and **a + expand\_limits()**  
Ensure limits include values across all plots.
- b + geom\_curve(aes(yend = lat + 1, xend = long + 1), curvature = 1)** - x, yend, alpha, angle, color, curvature, linetype, size
- a + geom\_path(lineend = "butt", linejoin = "round", linemitre = 1)** - x, y, alpha, color, group, linetype, size
- a + geom\_polygon(aes(alpha = 50))** - x, y, alpha, color, fill, group, subgroup, linetype, size
- b + geom\_rect(aes(xmin = long, ymin = lat, xmax = long + 1, ymax = lat + 1))** - xmax, xmin, ymax, ymin, alpha, color, fill, linetype, size
- a + geom\_ribbon(aes(ymin = unemploy - 900, ymax = unemploy + 900))** - x, ymax, ymin, alpha, color, fill, group, linetype, size

### LINE SEGMENTS

common aesthetics: x, y, alpha, color, linetype, size

```
b + geom_abline(aes(intercept = 0, slope = 1))
b + geom_hline(aes(yintercept = lat))
b + geom_vline(aes(xintercept = long))

b + geom_segment(aes(yend = lat + 1, xend = long + 1))
b + geom_spoke(aes(angle = 1:1155, radius = 1))
```

### ONE VARIABLE continuous

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

c + geom_area(stat = "bin")
x, y, alpha, color, fill, linetype, size

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

c + geom_dotplot()
x, y, alpha, color, fill, linetype, size

c + geom_freqpoly()
x, y, alpha, color, group, linetype, size

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

c2 + geom_qq(aes(sample = hwy))
x, y, alpha, color, fill, linetype, size, weight
```

### discrete

```
d <- ggplot(mpg, aes(fl))
d + geom_bar()
x, alpha, color, fill, linetype, size, weight
```

### TWO VARIABLES

#### both continuous

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

e + geom_label(aes(label = cty), nudge_x = 1, nudge_y = 1)
x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust

e + geom_point()
x, y, alpha, color, fill, shape, size, stroke

e + geom_quantile()
x, y, alpha, color, group, linetype, size, weight

e + geom_rug(sides = "bl")
x, y, alpha, color, linetype, size

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)
x, y, label, alpha, angle, color, family, fontface, hjust, lineheight, size, vjust
```

#### one discrete, one continuous

```
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
```

#### both discrete

```
g <- ggplot(diamonds, aes(cut, color))

g + geom_count()
x, y, alpha, color, fill, shape, size, stroke

e + geom_jitter(height = 2, width = 2)
x, y, alpha, color, fill, shape, size
```

### 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, color, group, linetype, size, weight

l + geom_contour_filled(aes(fill = z))
x, y, alpha, color, fill, group, linetype, size, subgroup
```



### continuous bivariate distribution

```
h <- ggplot(diamonds, aes(carat, price))

h + geom_bin2d(binwidth = c(0.25, 500))
x, y, alpha, color, fill, linetype, size, weight

h + geom_density_2d()
x, y, alpha, color, group, linetype, size

h + geom_hex()
x, y, alpha, color, fill, size
```

### continuous function

```
i <- ggplot(economics, aes(date, unemploy))

i + geom_area()
x, y, alpha, color, fill, linetype, size

i + geom_line()
x, y, alpha, color, group, linetype, size

i + geom_step(direction = "hv")
x, y, alpha, color, group, linetype, size
```

### visualizing error

```
jf <- data.frame(grp = c("A", "B"), fit = 4:5, se = 1:2)
jf <- ggplot(jf, 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, y, max, min, 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
```

### maps

Draw the appropriate geometric object depending on the simple features present in the data. aes() arguments:  
map\_id, alpha, color, fill, linetype, linewidth.

```
nc <- sf::st_read(system.file("shape/nc.shp", package = "sf"))

ggplot(nc) +
  geom_sf(aes(fill = AREA))
```

# Subset de los datos

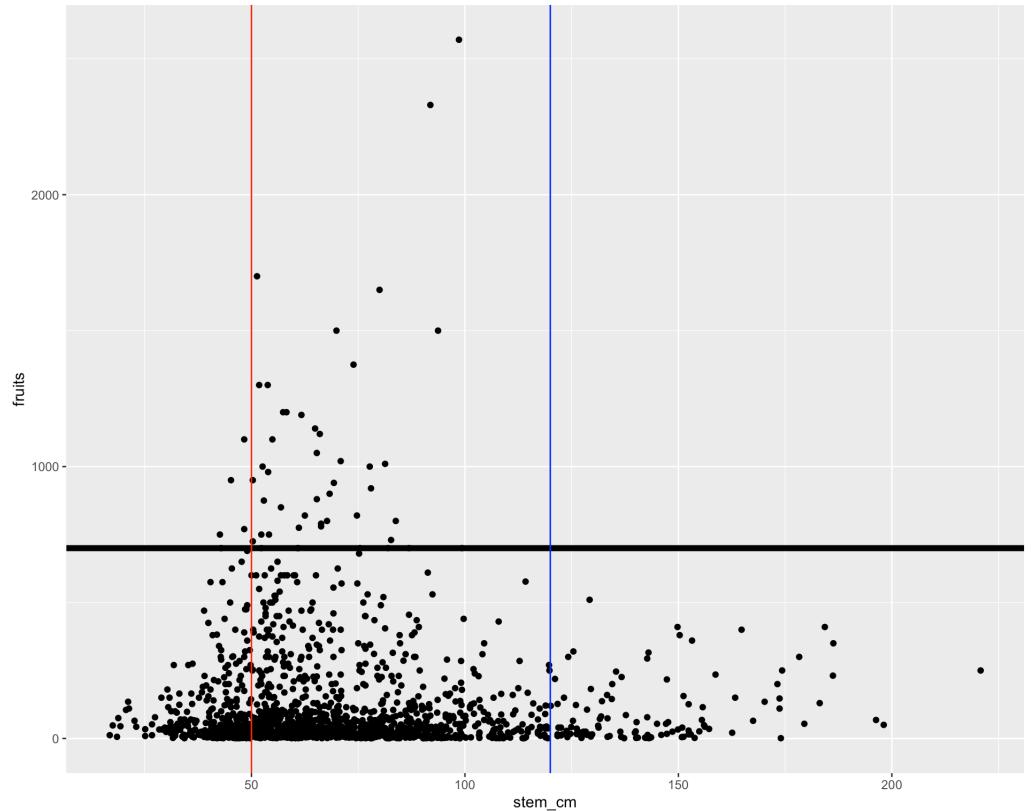
Seleccionar datos con información para diámetro de tronco. Sólo hay información para el sitio llamado “AND”.

```
dt_diam <- dt |> filter(!is.na(stem_cm))
```

```
filter: removed 80,777 rows (98%), 1,503 rows remaining
```

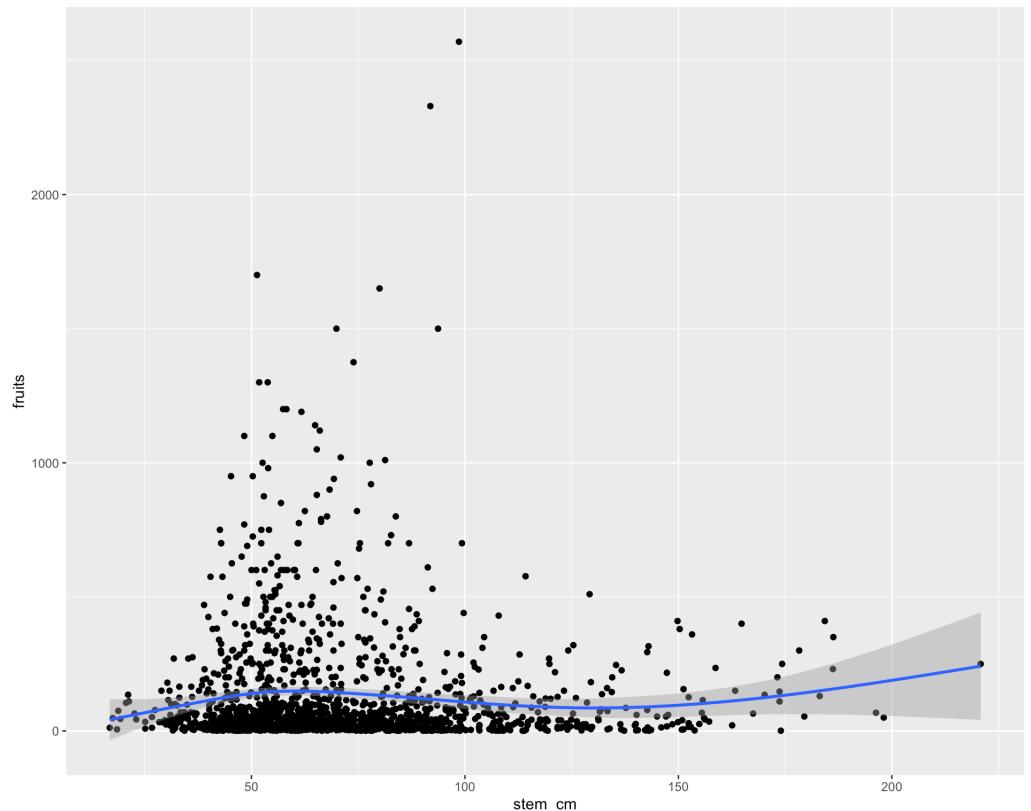
# Geoms - Líneas verticales y horizontales

```
ggplot(dt_diam,  
       aes(x = stem_cm, y = fruits)) +  
  geom_point() +  
  geom_hline(yintercept = 700,  
             size = 2) +  
  geom_vline(xintercept = c(50, 120),  
             color = c("red", "blue"))
```



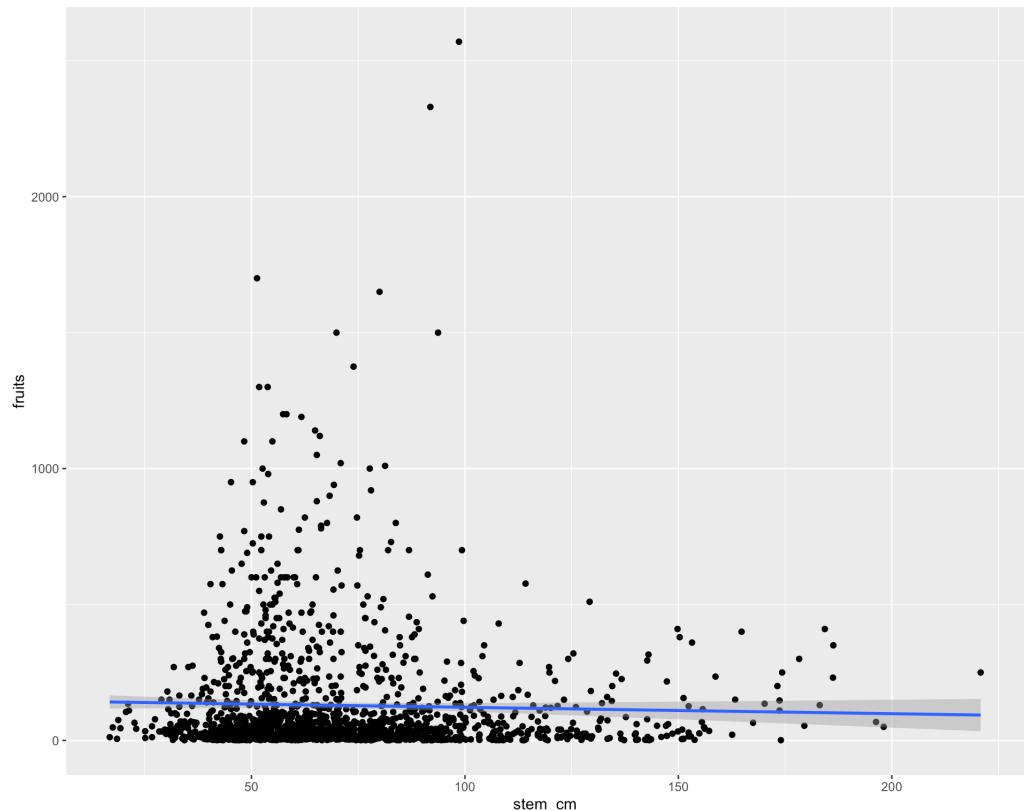
# Geoms - Líneas de tendencia

```
ggplot(dt_diam,  
       aes(x = stem_cm, y = fruits)) +  
  geom_point() +  
  geom_smooth()
```



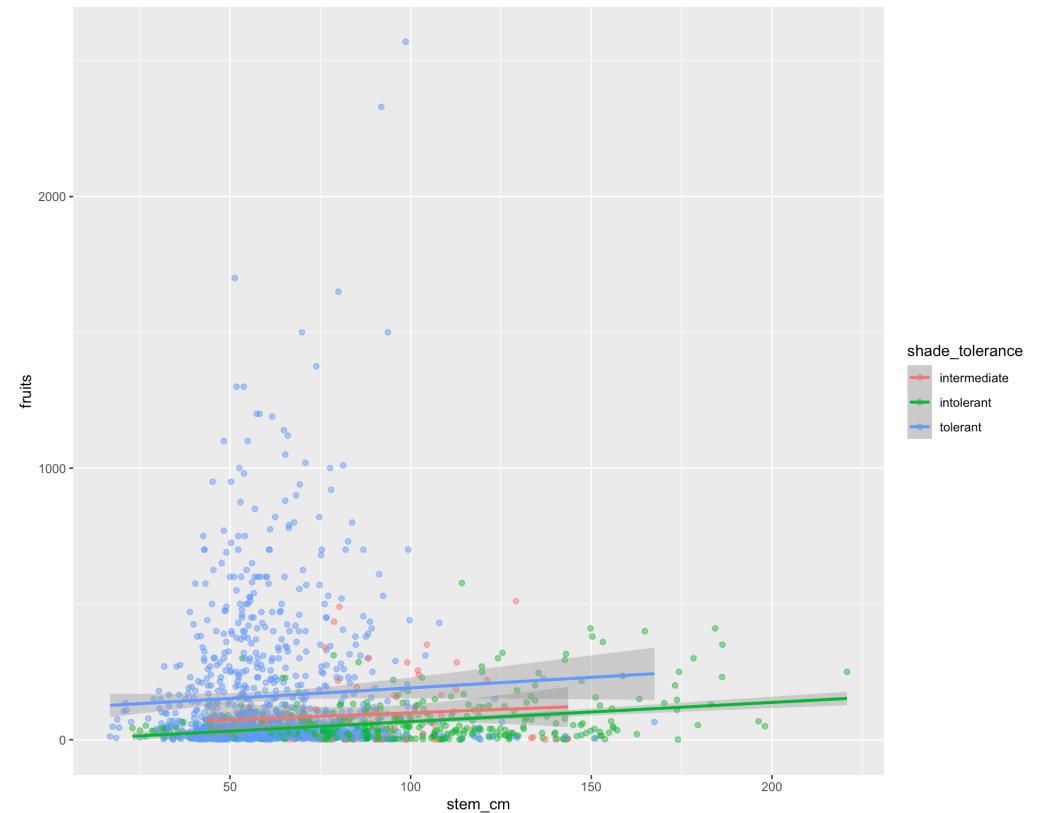
# Geoms - Líneas de tendencia

```
ggplot(dt_diam,  
       aes(x = stem_cm, y = fruits)) +  
  geom_point() +  
  geom_smooth(method = "lm")
```



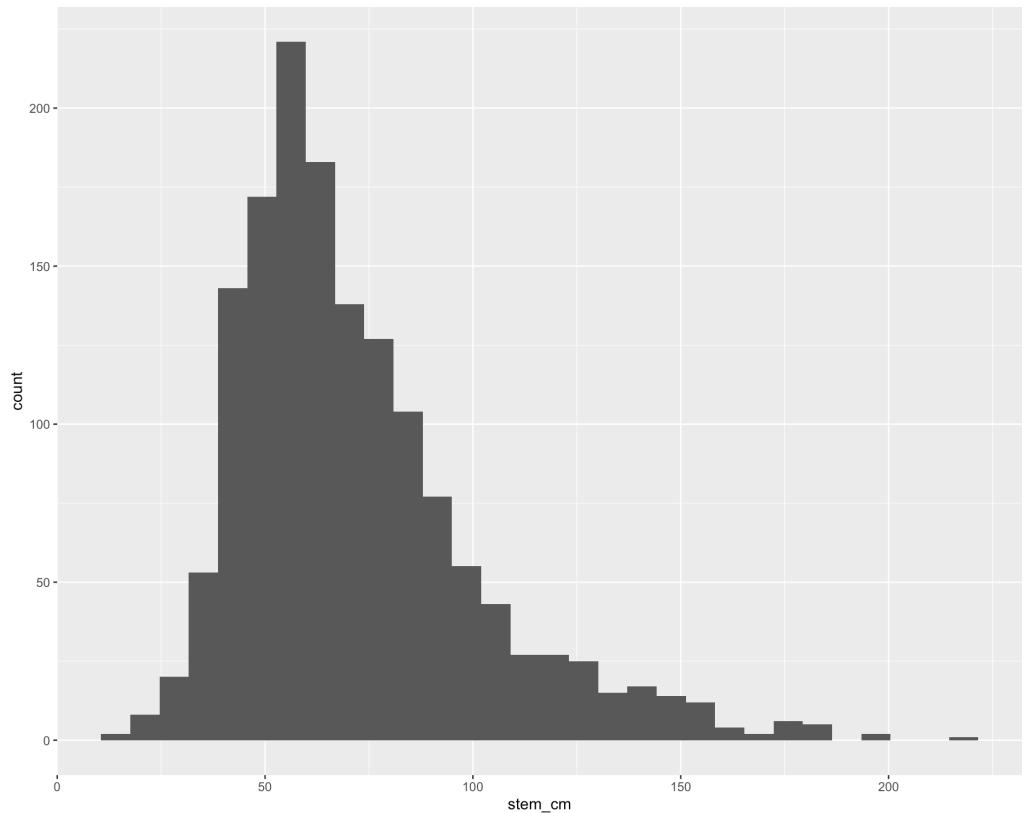
# Geoms - Líneas de tendencia

```
ggplot(dt_diam,  
       aes(x = stem_cm, y = fruits,  
            color = shade_tolerance)) +  
  geom_point(alpha = 0.5) +  
  geom_smooth(method = "lm")
```



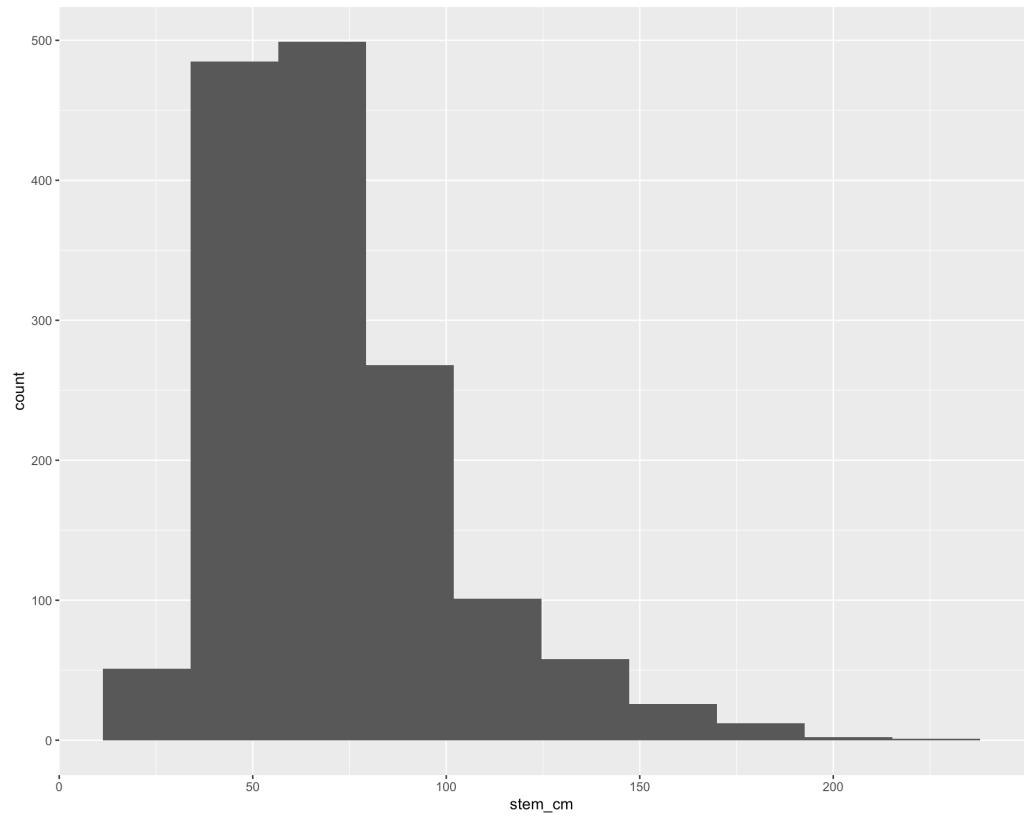
# Geoms - Histograma

```
ggplot(dt_diam,  
       aes(x = stem_cm)) +  
  geom_histogram()
```



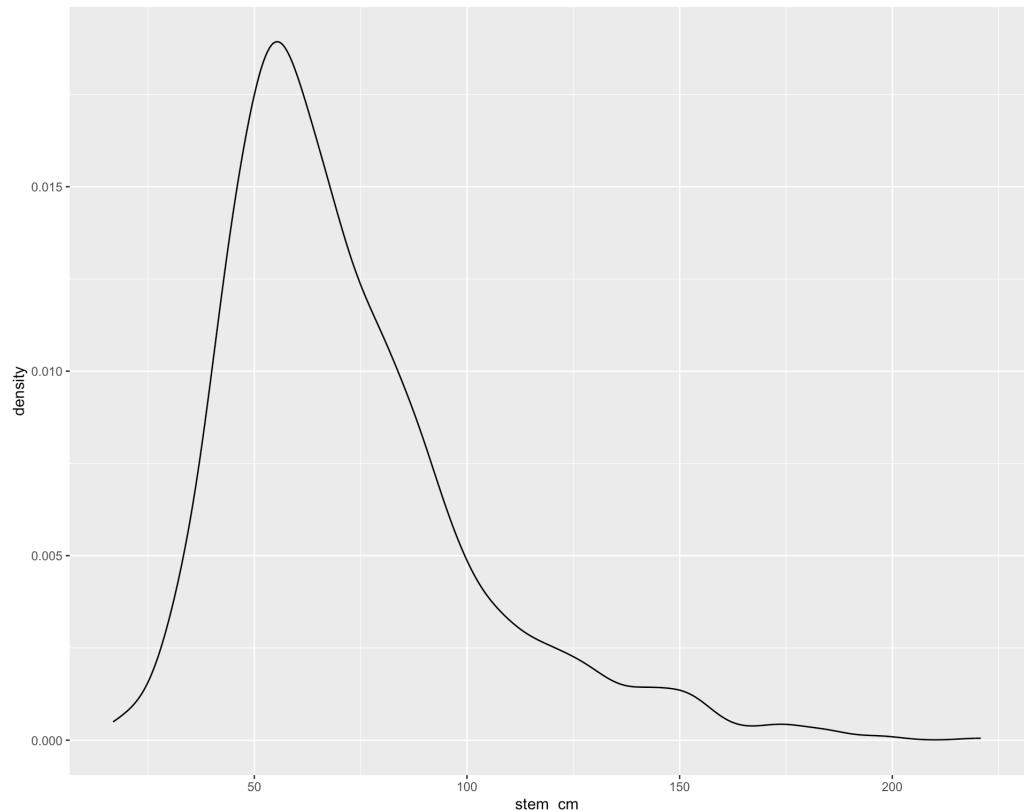
# Geoms - Histograma

```
ggplot(dt_diam,  
       aes(x = stem_cm)) +  
  geom_histogram(bins = 10)
```



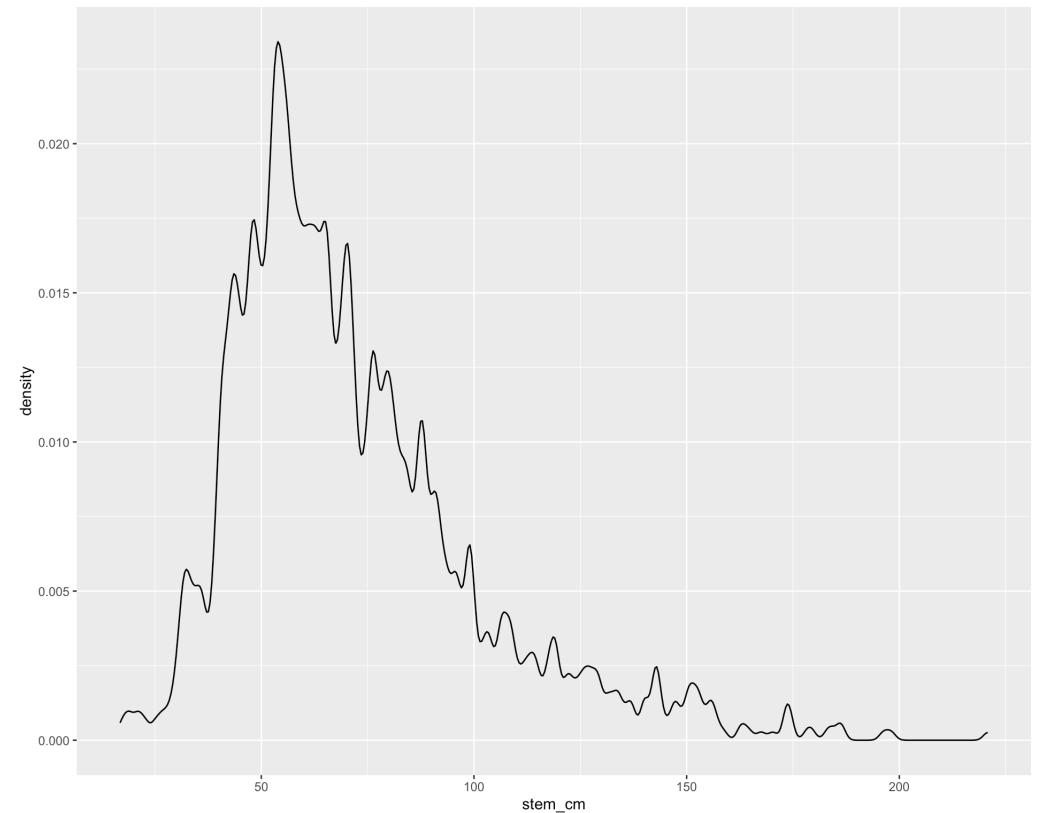
# Geoms - Densidad

```
ggplot(dt_diam,  
       aes(x = stem_cm)) +  
  geom_density()
```



# Geoms - Densidad

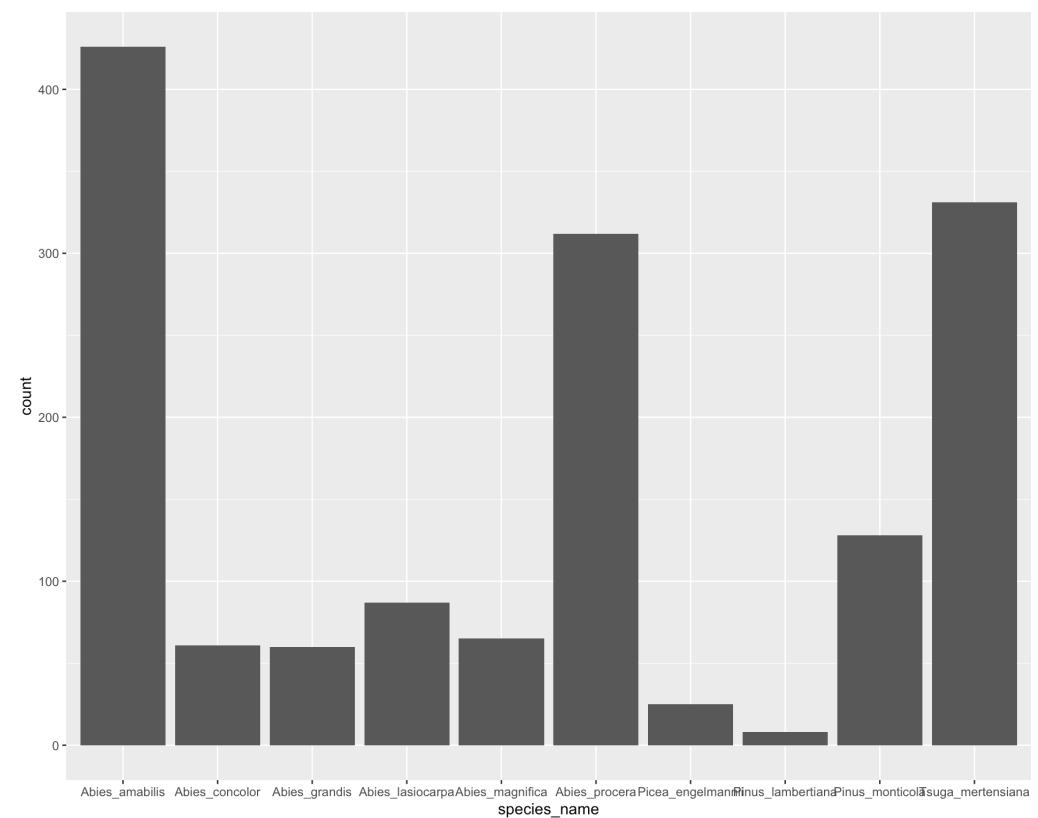
```
ggplot(dt_diam,  
       aes(x = stem_cm)) +  
  geom_density(bw = 1)
```



# Geoms - Barras

Contar número de casos: `stat = "count"`

```
ggplot(dt_diam,  
       aes(x = species_name)) +  
  geom_bar(stat = "count")
```

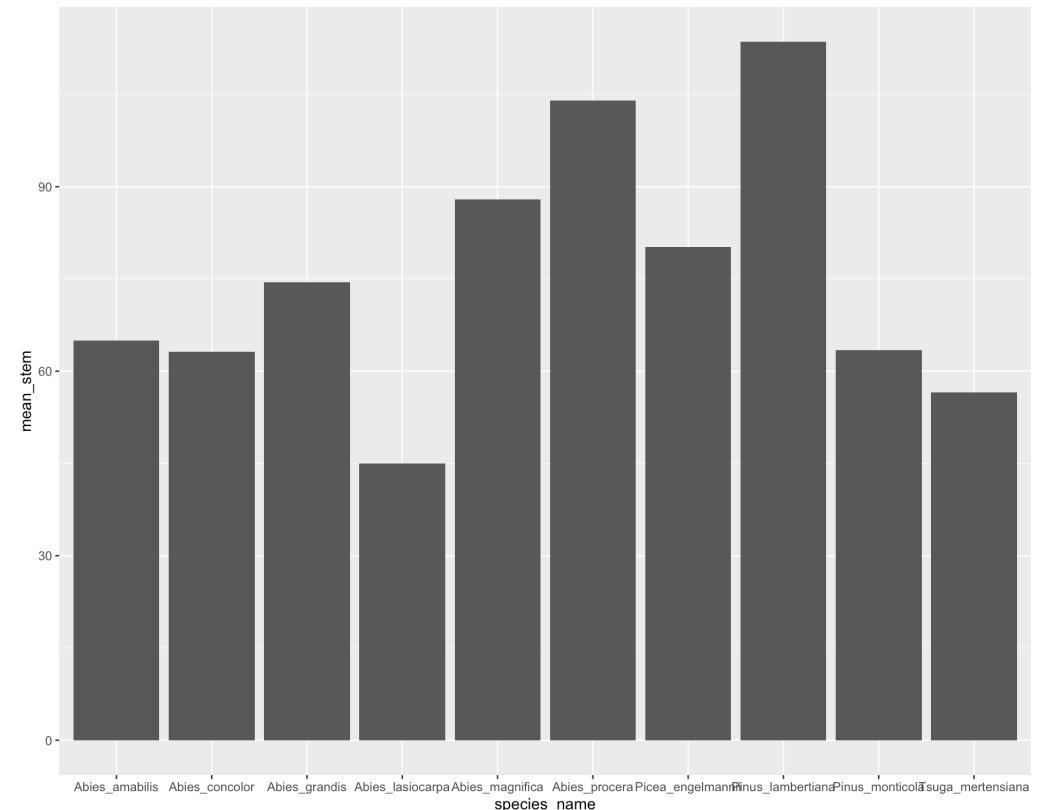


# Geoms - Barras

Para datos resumidos - *alternativa 1:*

Usar valores ya calculados: `stat = "identity"`

```
dt_diam |>  
  group_by(species_name) |>  
  summarise(mean_stem = mean(stem_cm)) |>  
  ggplot(aes(x = species_name,  
              y = mean_stem)) +  
  geom_bar(stat = "identity")
```

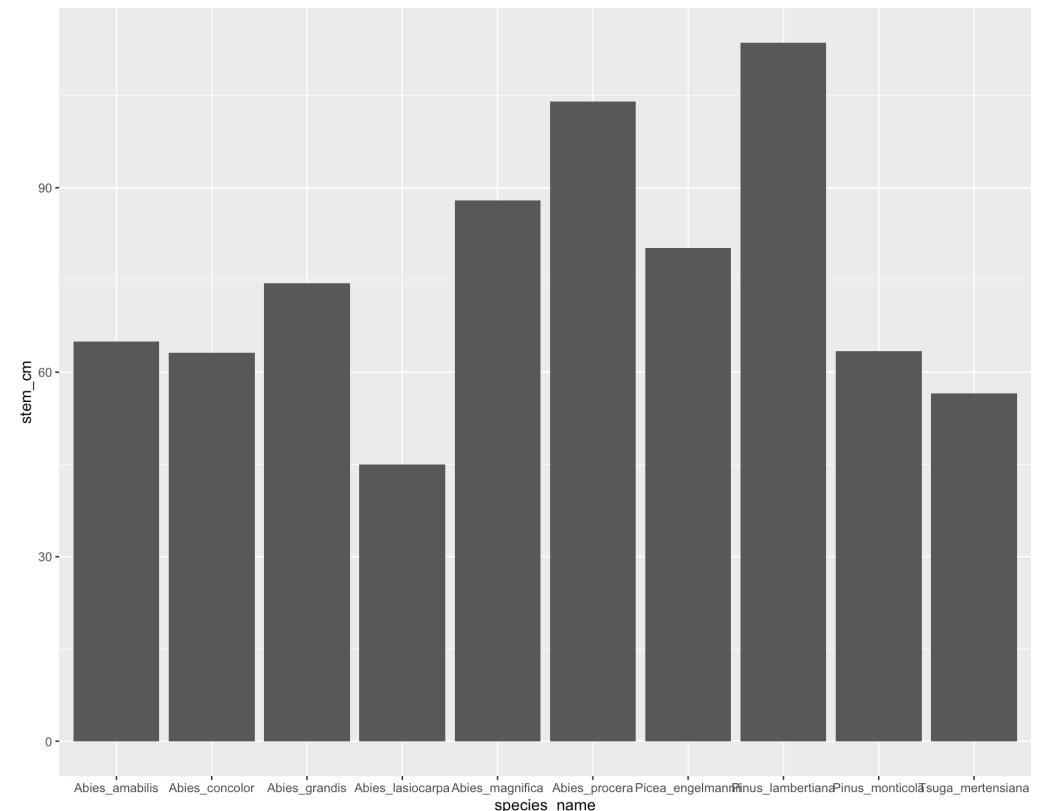


# Geoms - Barras

Para datos resumidos - *alternativa 2:*

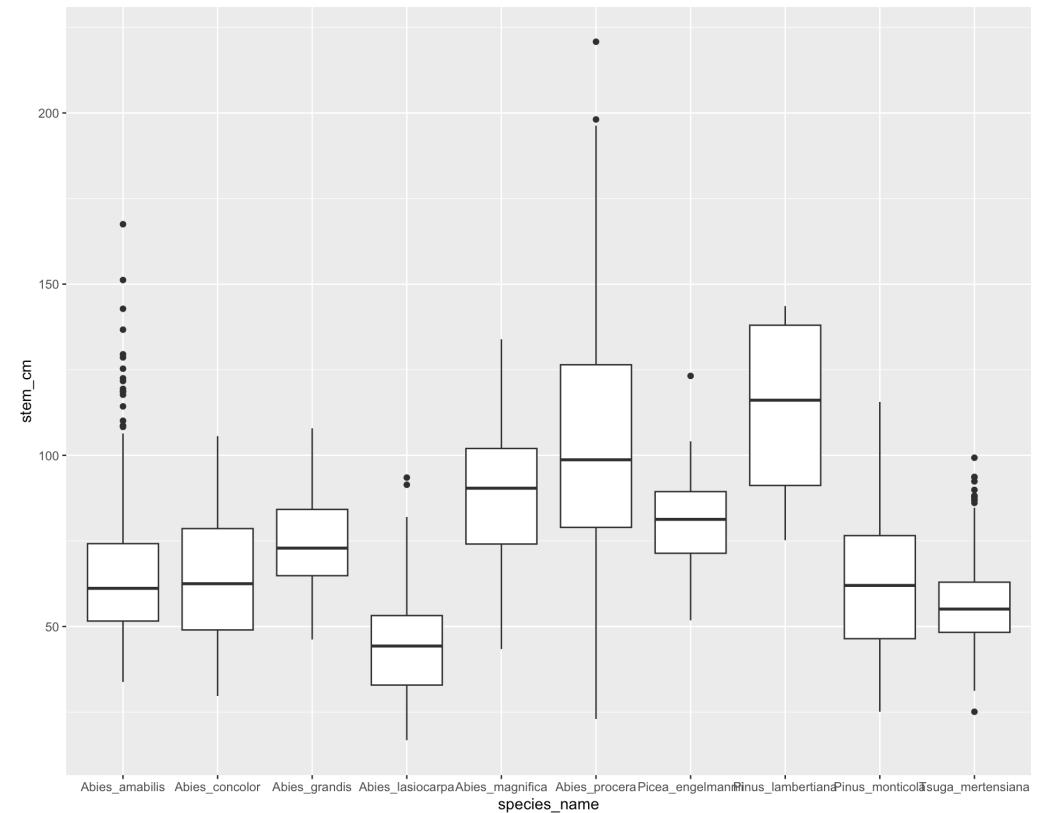
Calcular valores dentro de ggplot: **stat = "summary"**

```
ggplot(dt_diam,  
       aes(x = species_name,  
            y = stem_cm)) +  
  geom_bar(stat = "summary",  
           fun = "mean")
```



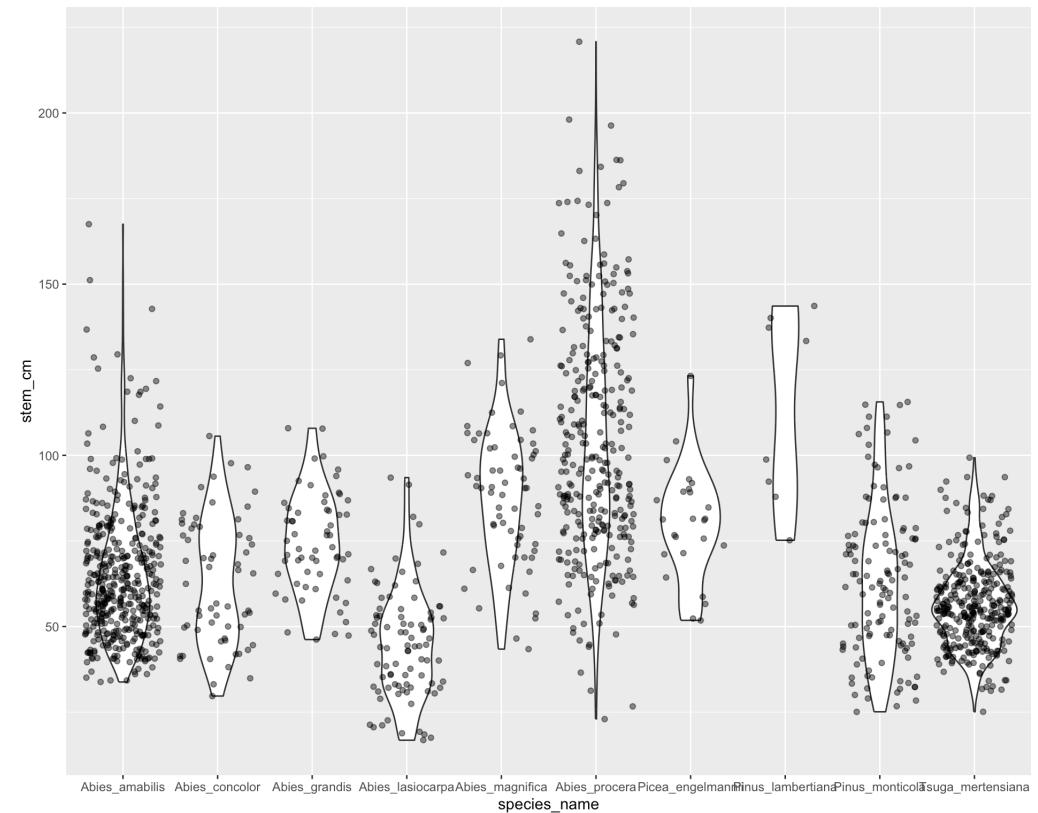
# Geoms - Boxplot

```
ggplot(dt_diam,  
       aes(x = species_name,  
            y = stem_cm)) +  
  geom_boxplot()
```



# Geoms - Violin y puntos

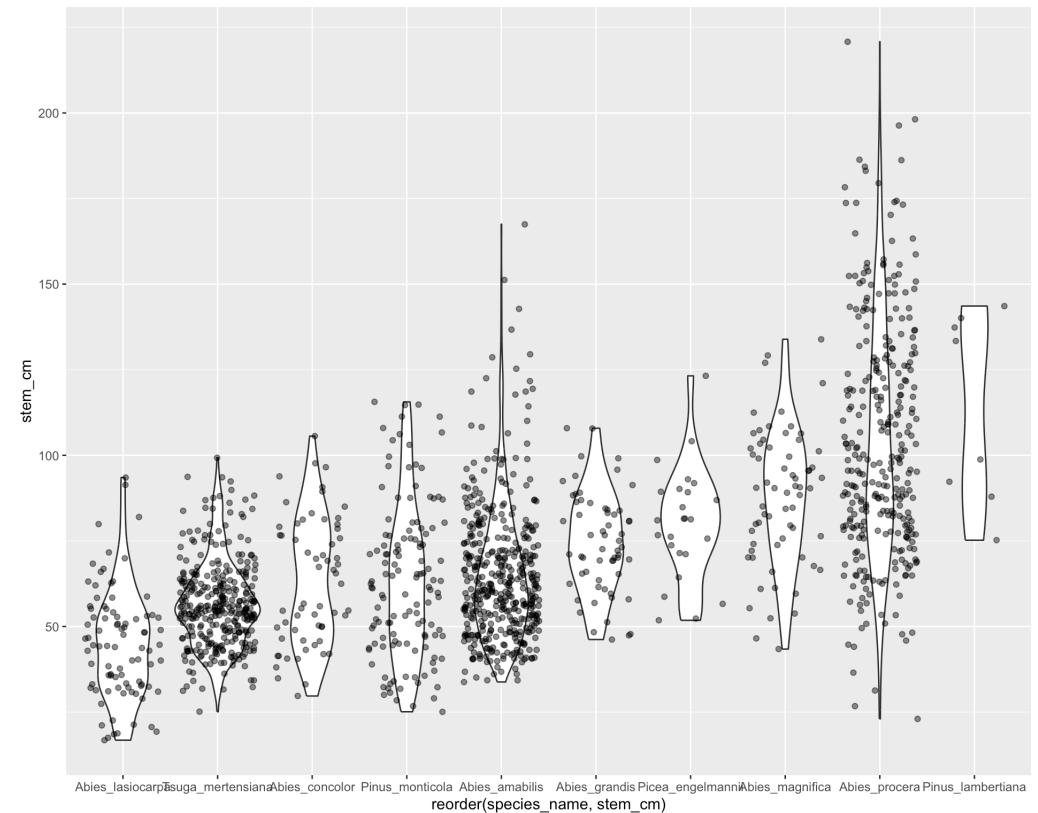
```
ggplot(dt_diam,  
       aes(x = species_name,  
            y = stem_cm)) +  
  geom_violin() +  
  geom_jitter(alpha = 0.5)
```



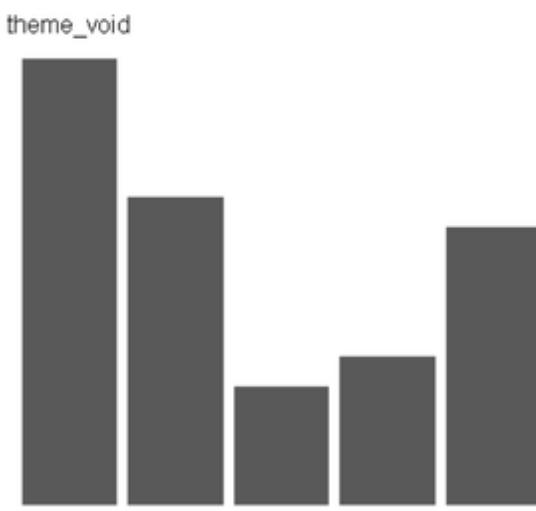
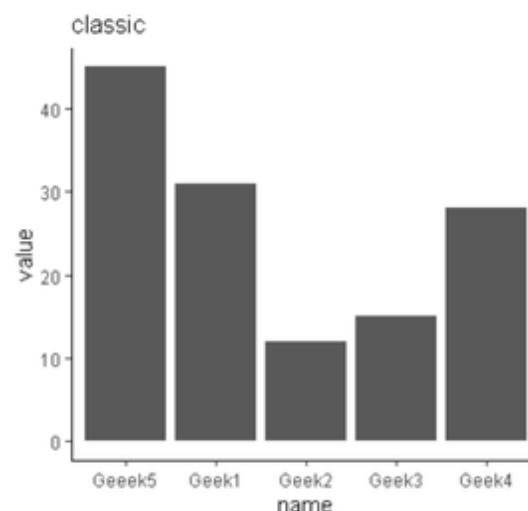
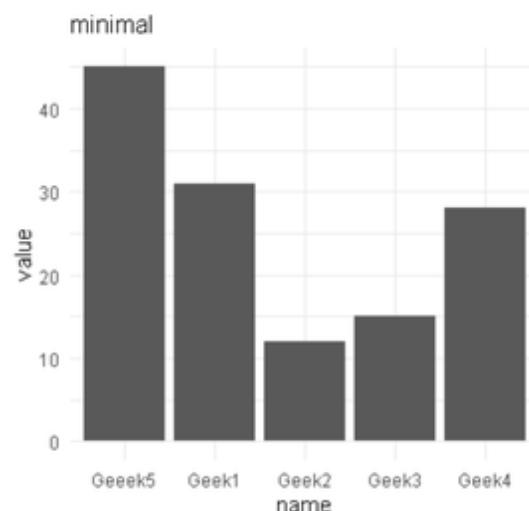
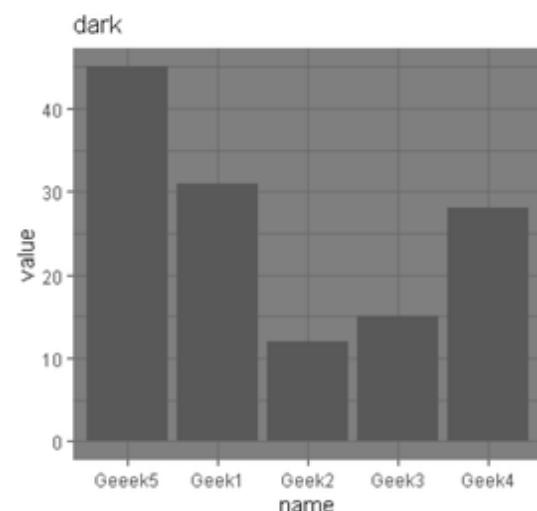
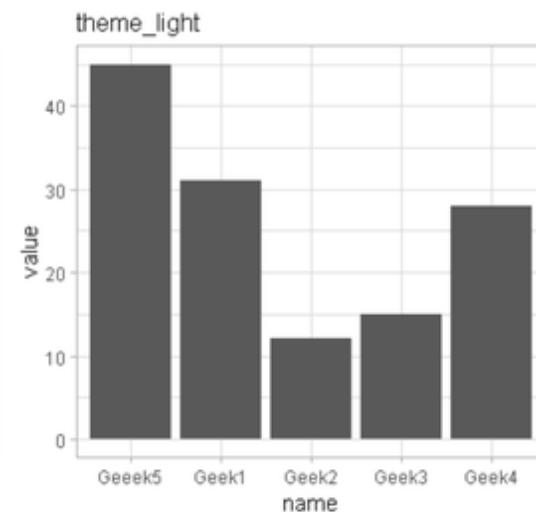
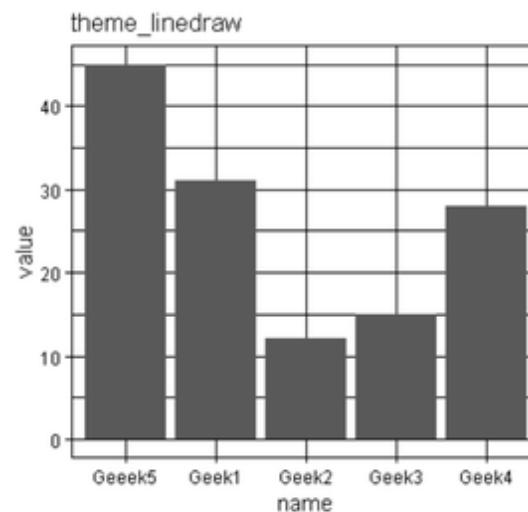
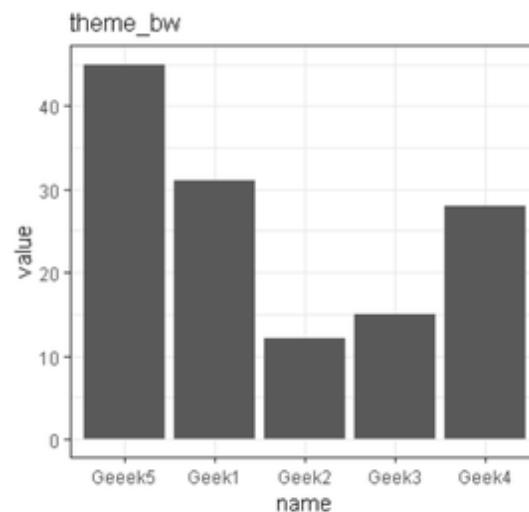
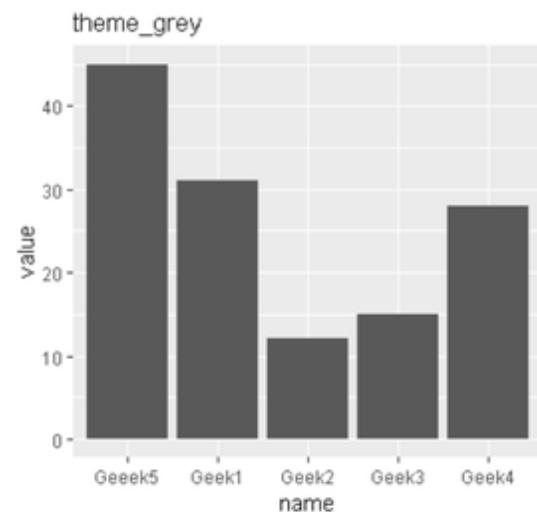
# Geoms - Violin y puntos

Ordernar eje de la X en base a valores del eje de la Y

```
ggplot(dt_diam,  
       aes(x = reorder(species_name, stem_cm),  
            y = stem_cm)) +  
  geom_violin() +  
  geom_jitter(alpha = 0.5)
```

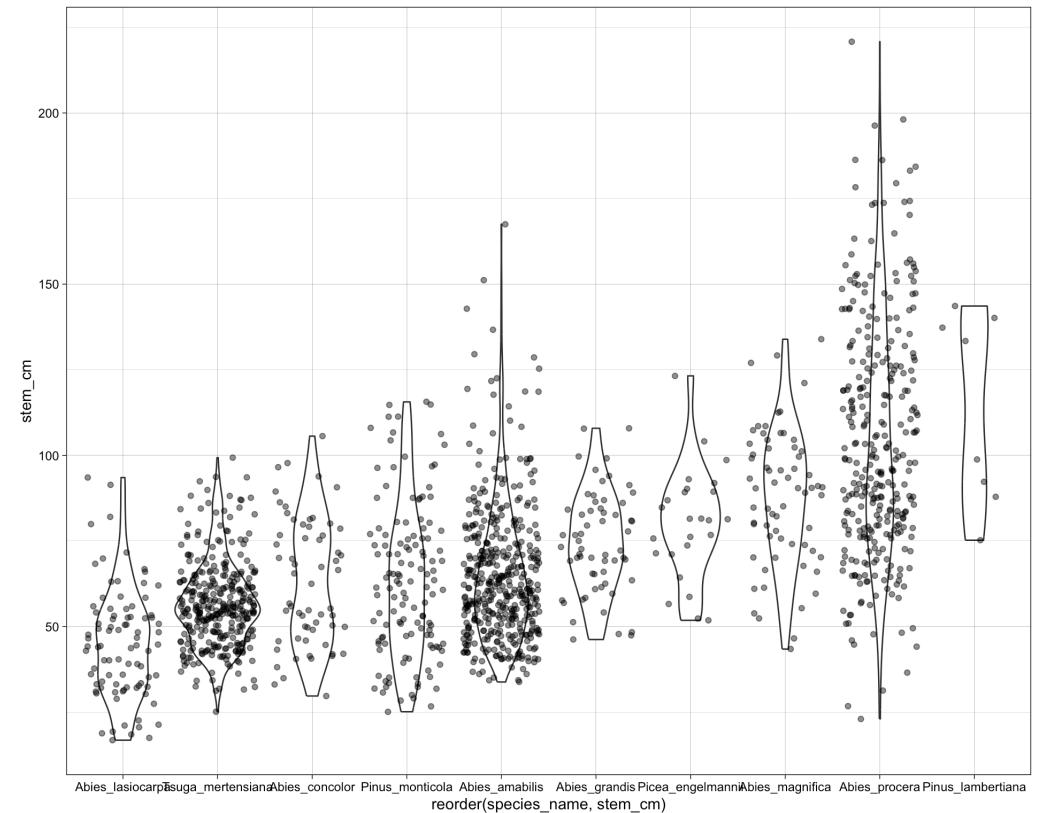


# Themes



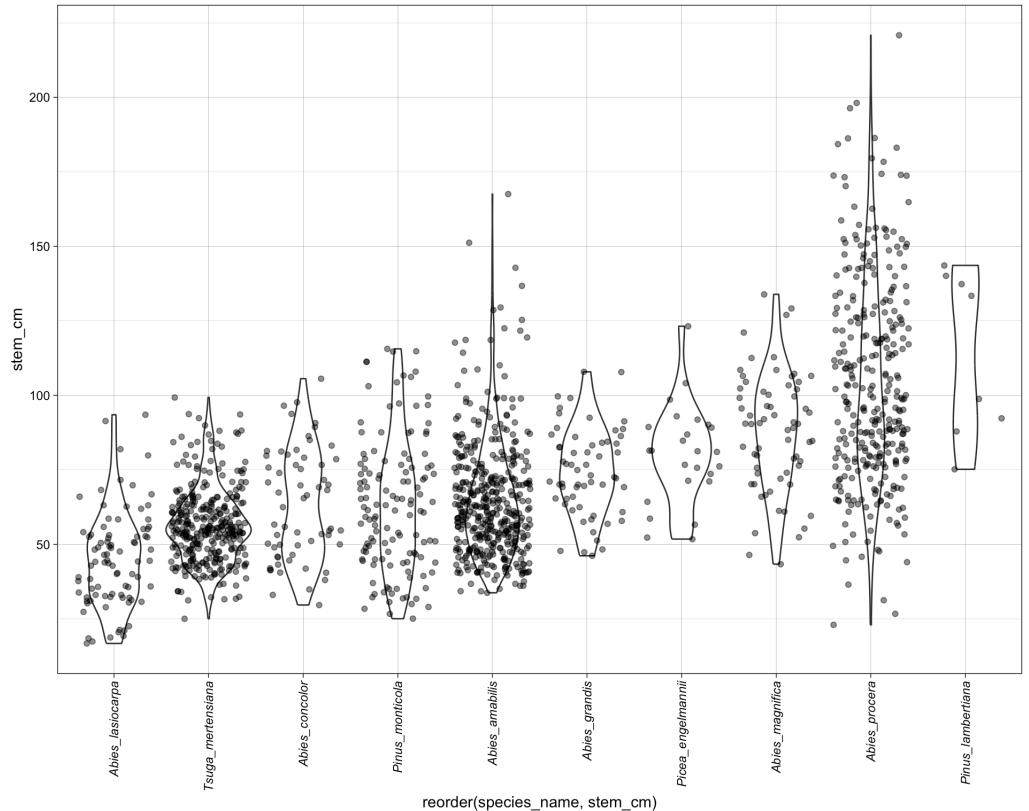
# Themes

```
ggplot(dt_diam,  
       aes(x = reorder(species_name, stem_cm),  
            y = stem_cm)) +  
  geom_violin() +  
  geom_jitter(alpha = 0.5) +  
  theme_linedraw()
```



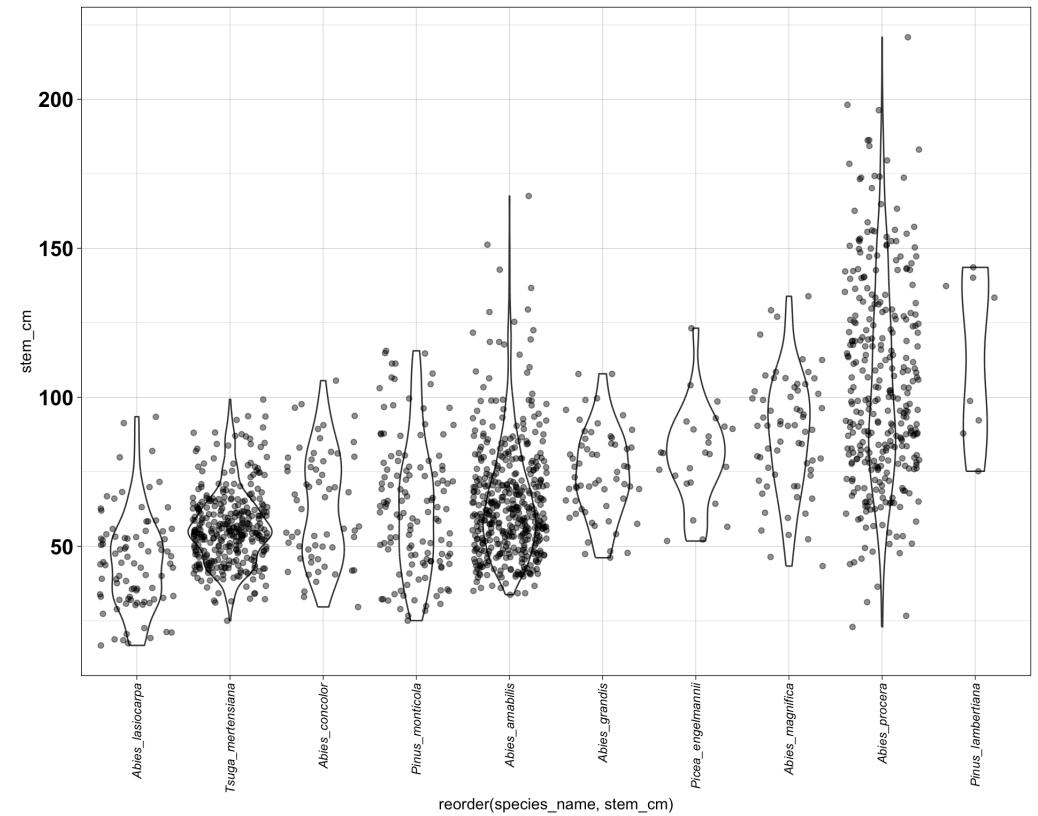
# Themes

```
ggplot(dt_diam,  
       aes(x = reorder(species_name, stem_cm),  
            y = stem_cm)) +  
  geom_violin() +  
  geom_jitter(alpha = 0.5) +  
  theme_linedraw() +  
  theme(axis.text.x = element_text(  
    angle = 90,  
    hjust = 1, vjust = 0.5,  
    face = "italic"))
```



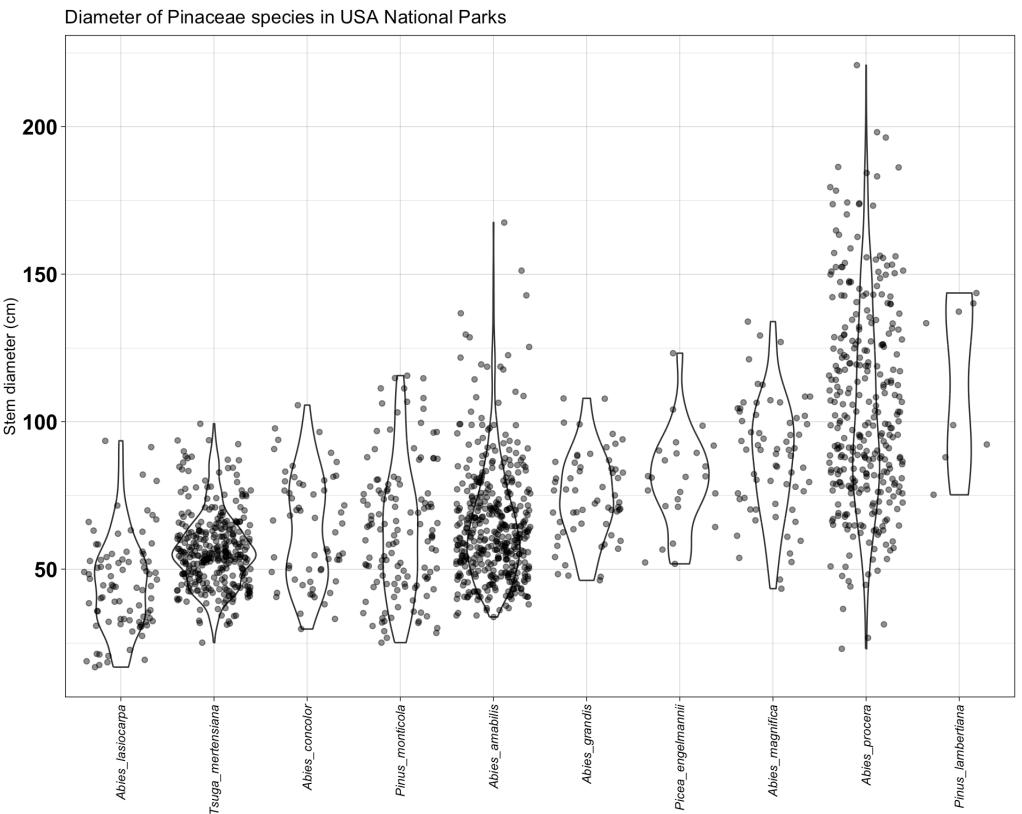
# Themes

```
ggplot(dt_diam,  
       aes(x = reorder(species_name, stem_cm),  
            y = stem_cm)) +  
  geom_violin() +  
  geom_jitter(alpha = 0.5) +  
  theme_linedraw() +  
  theme(axis.text.x = element_text(  
    angle = 90,  
    hjust = 1, vjust = 0.5,  
    face = "italic"),  
    axis.text.y = element_text(  
    size = 15, face = "bold"))
```



# Themes

```
ggplot(dt_diam,  
       aes(x = reorder(species_name, stem_cm),  
            y = stem_cm)) +  
  geom_violin() +  
  geom_jitter(alpha = 0.5) +  
  theme_linedraw() +  
  theme(axis.text.x = element_text(  
    angle = 90,  
    hjust = 1, vjust = 0.5,  
    face = "italic"),  
    axis.text.y = element_text(  
      size = 15, face = "bold")) +  
  labs(x = NULL,  
       y = "Stem diameter (cm)",  
       title = "Diameter of Pinaceae species")
```



# Themes

Para fijar un tema que se aplique a todos los gráficos:

```
theme_set(theme_minimal())
```

Existen muchos paquetes con temas predeterminados. Muchos también vienen con especificaciones para las escalas de los aesthetics (scales). Ejemplos:

```
library(hrbrthemes)
library(ggthemes)
library(ggpmolological)
library(tvthemer)
library(ggtech)
library(ggthemr)
library(ggsci)
```

`library(ggThemeAssist)` es una addin de RStudio que ayuda a cambiar la apariencia de los temas. <https://github.com/calligross/ggthemeassist>

# Composición de figuras

library(patchwork)



# Composición de figuras

Nombrar los plots como objetos

```
library(patchwork)

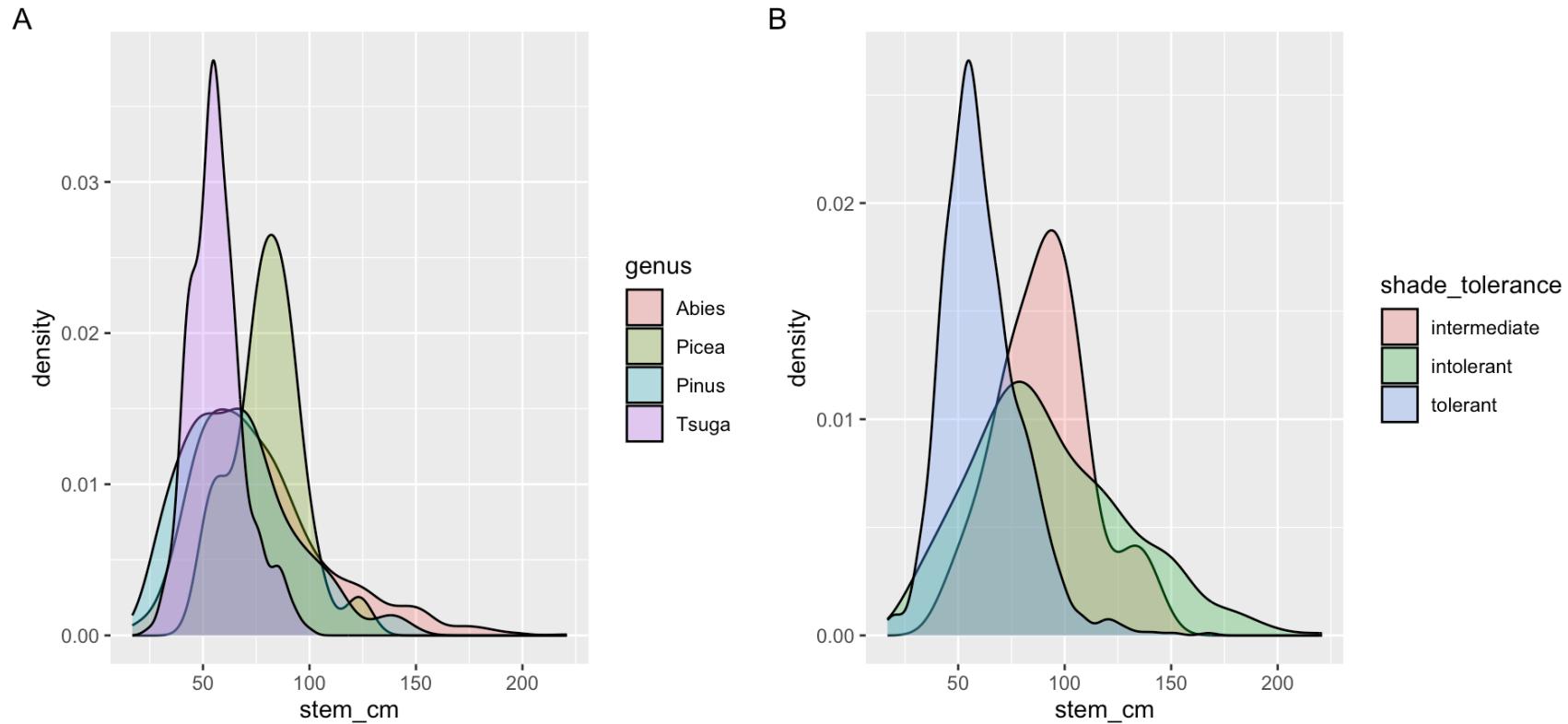
p1 <- ggplot(dt_diam,
    aes(x = stem_cm,
        fill = genus)) +
geom_density(alpha = 0.3)

p2 <- ggplot(dt,
    aes(x = stem_cm,
        fill = shade_tolerance)) +
geom_density(alpha = 0.3)

p1 + p2 +
plot_annotation(
title = "Diameter distribution",
tag_levels = 'A')
```

# Composición de figuras

Diameter distribution



# Guardar gráficos

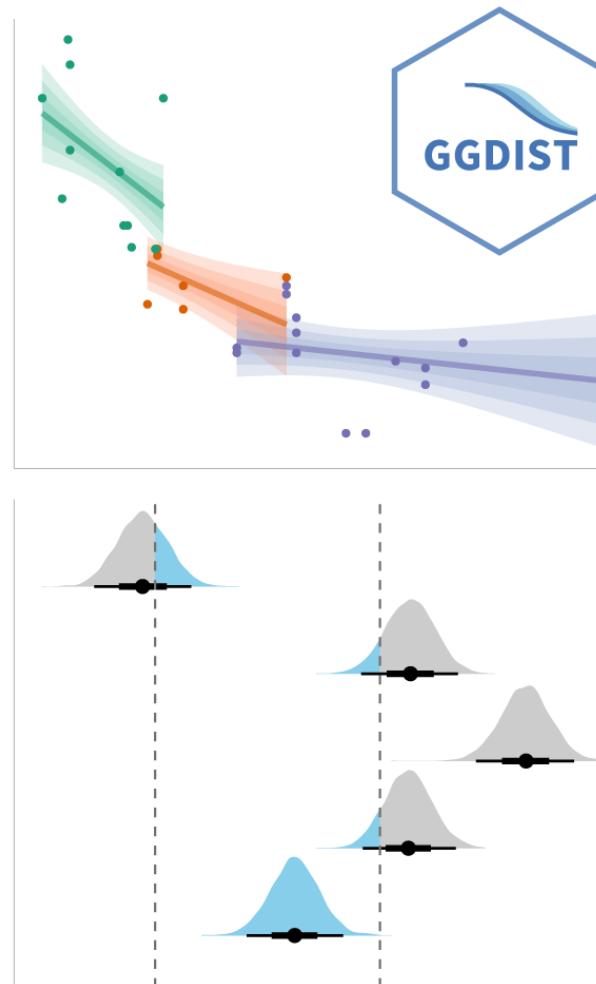
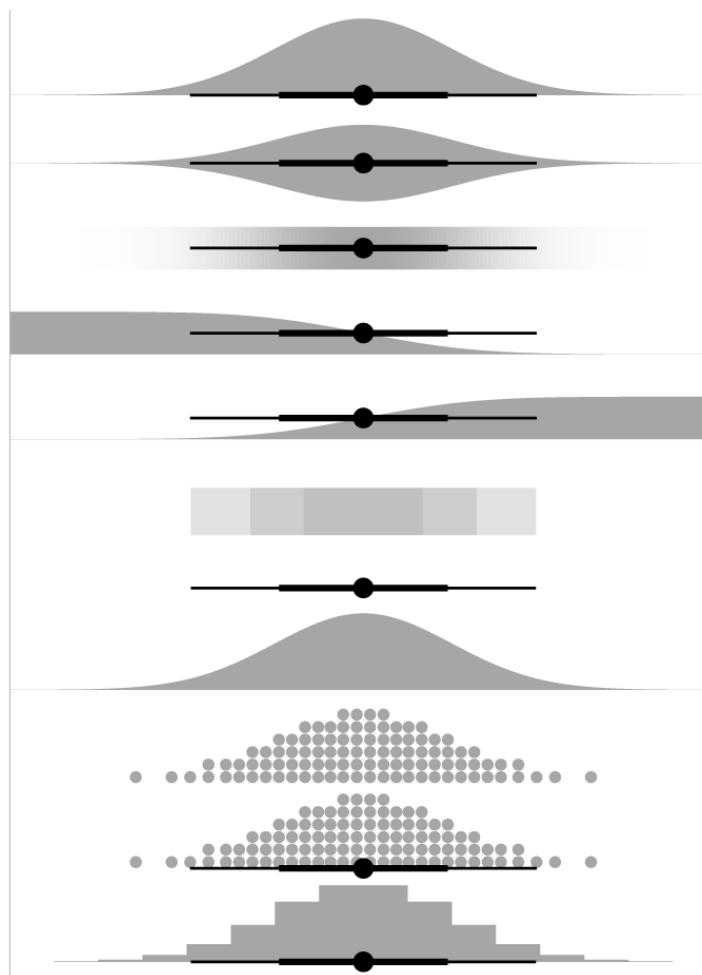
```
ggsave(p1, width = 20, units = "cm",
       filename = here("img/figure_1.pdf"))
```

Argumentos:

- width
- height
- units = ("in", "cm", "mm", "px")
- device = ("png", "pdf", "jpeg", "tiff", "svg", ...)

# Otros paquetes útiles

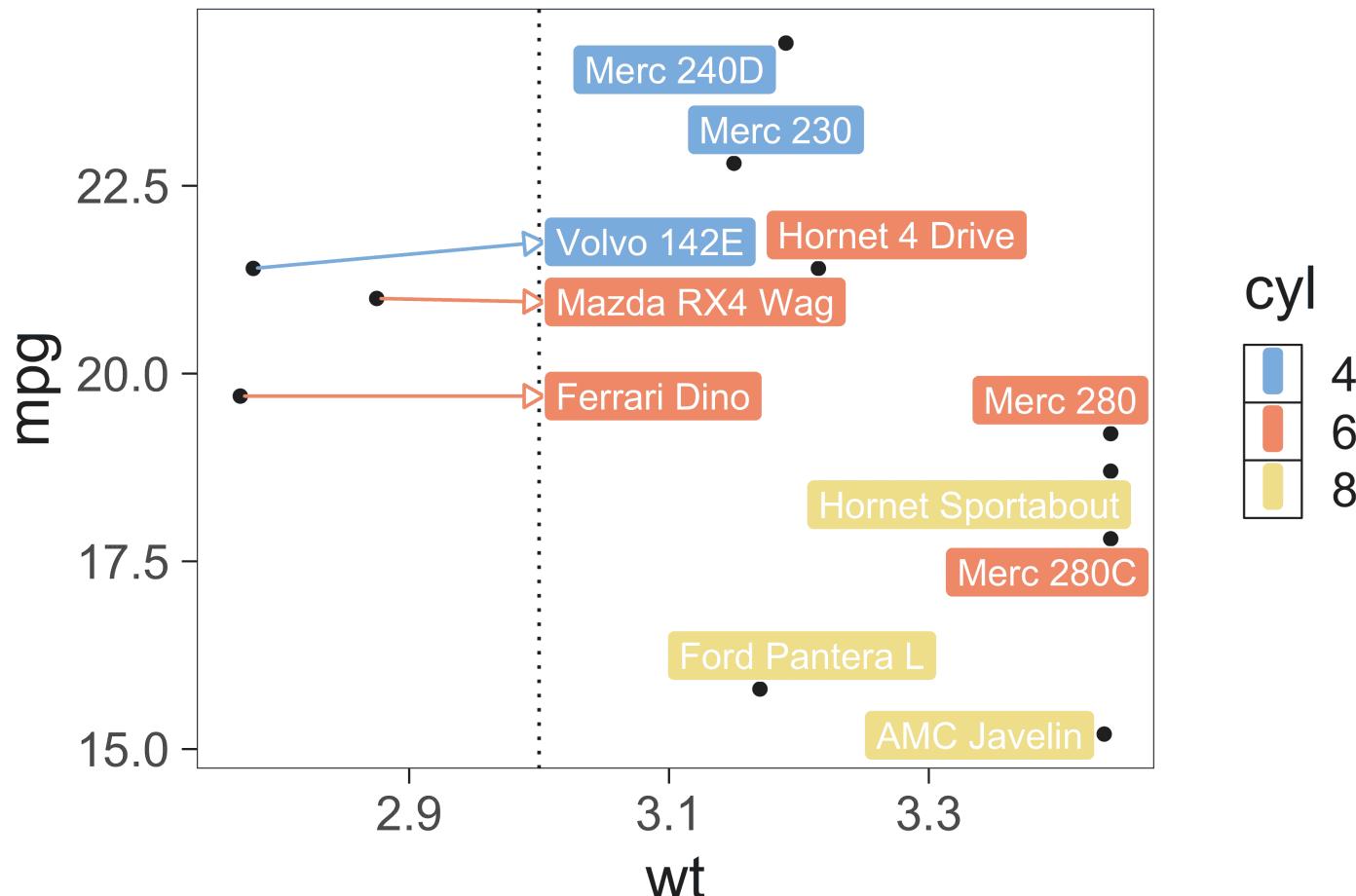
`library(ggdist)` - graficar datos con distribuciones amplias (ej. Bayes posteriores).



<https://mjskay.github.io/ggdist/>

# Otros paquetes útiles

library(ggrepel) - etiquetar datos

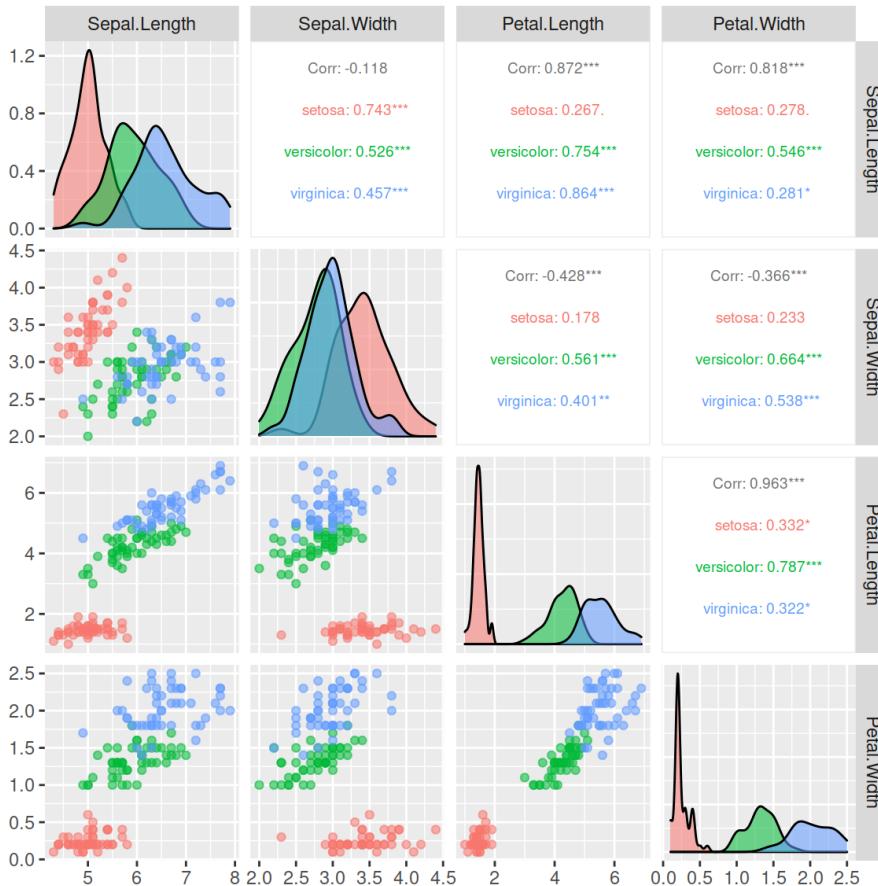


<https://cran.r-project.org/web/packages/ggrepel/vignettes/ggrepel.html>

# Otros paquetes útiles

library(GGally) - exploración de datos y relación entre variables.

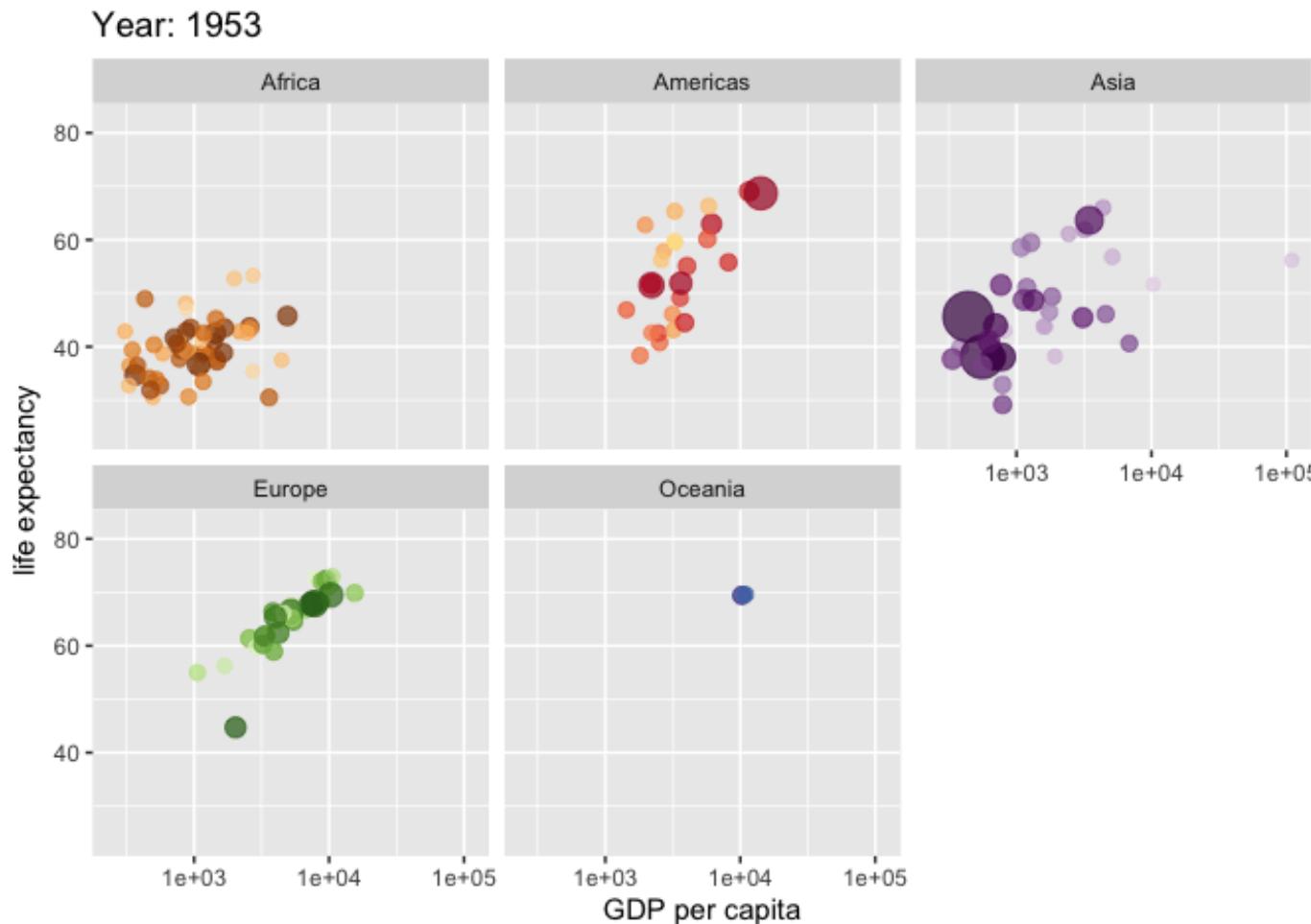
```
GGally::ggpairs(dataset)
```



<https://github.com/ggobi/ggally>

# Otros paquetes útiles

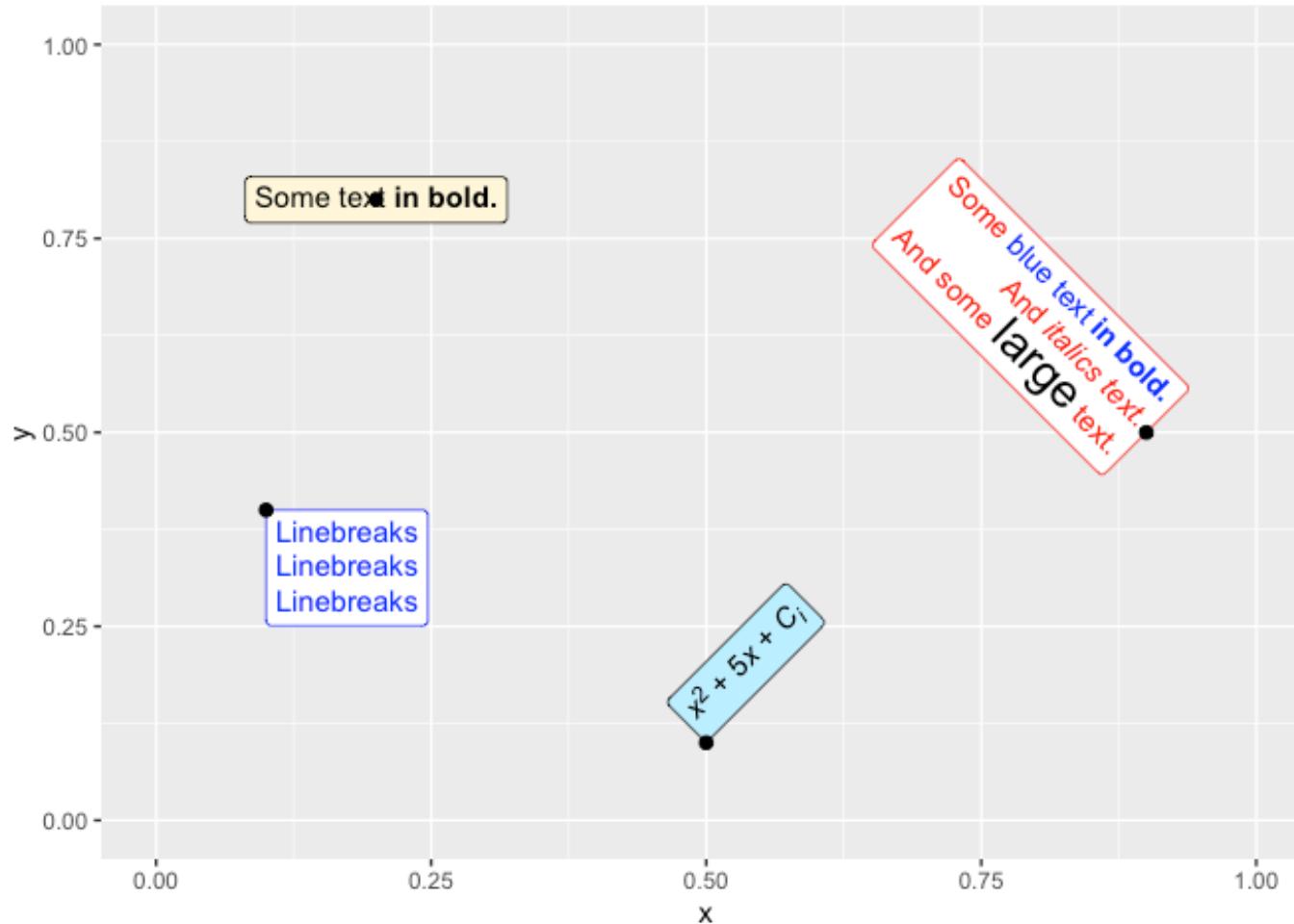
library(gganimate) - animar gráficos



<https://gganimate.com/>

# Otros paquetes útiles

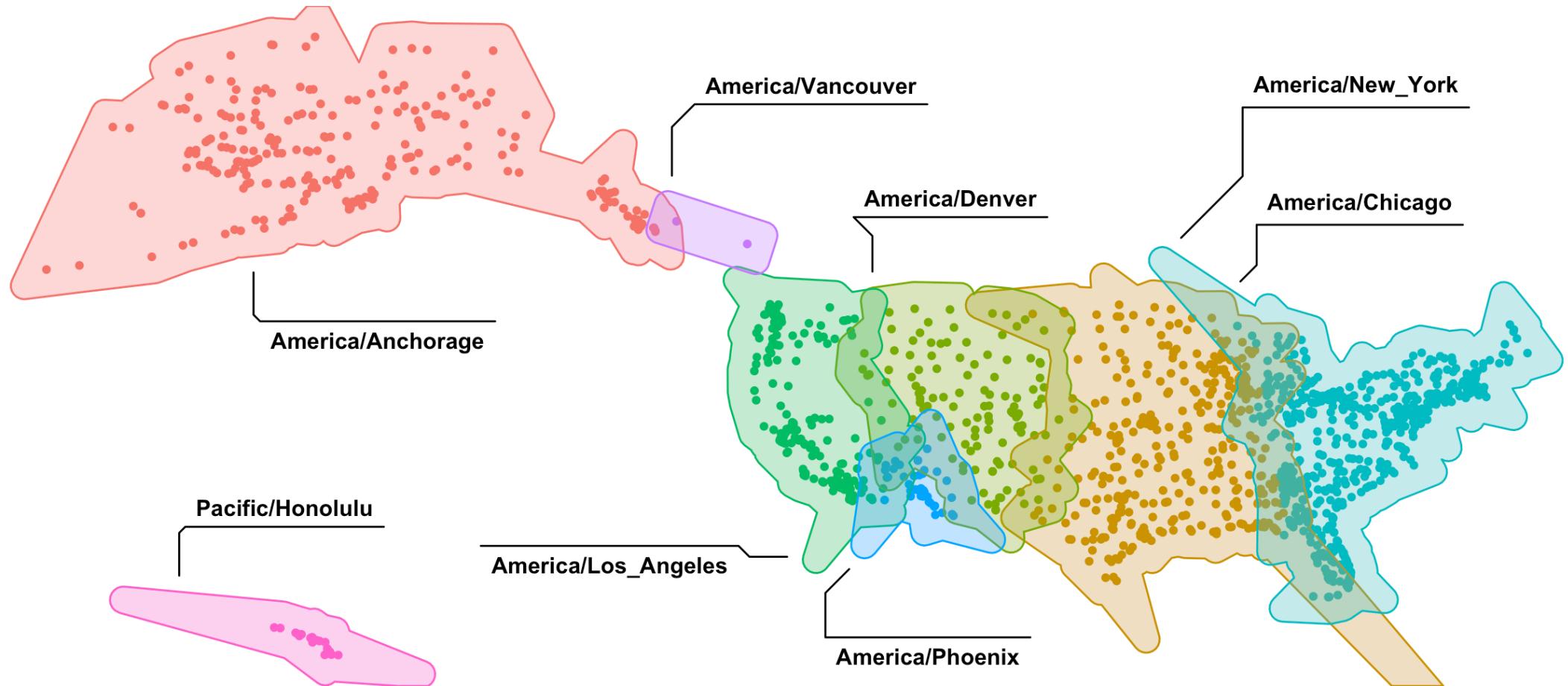
library(ggtext) -



<https://github.com/wilkelab/ggtext>

# Otros paquetes útiles

library(ggforce) - resaltar características de los datos.



<https://github.com/thomasp85/ggforce>

# Más extensiones de GGplot

<https://github.com/erikgahner/awesome-ggplot2>

<https://exts.ggplot2.tidyverse.org/gallery/>

# Recursos

- Referencia base a ggplot2
- ggplot2 book - Hadley Wickham
- R for Data Science Book - 3. Data visualization
- R Graphics Cookbook, 2nd edition - Winston Chang
- Fundamentals of Data Visualization - Claus O. Wilke
- RStudio CheatSheets - “*Data visualization with ggplot2*”

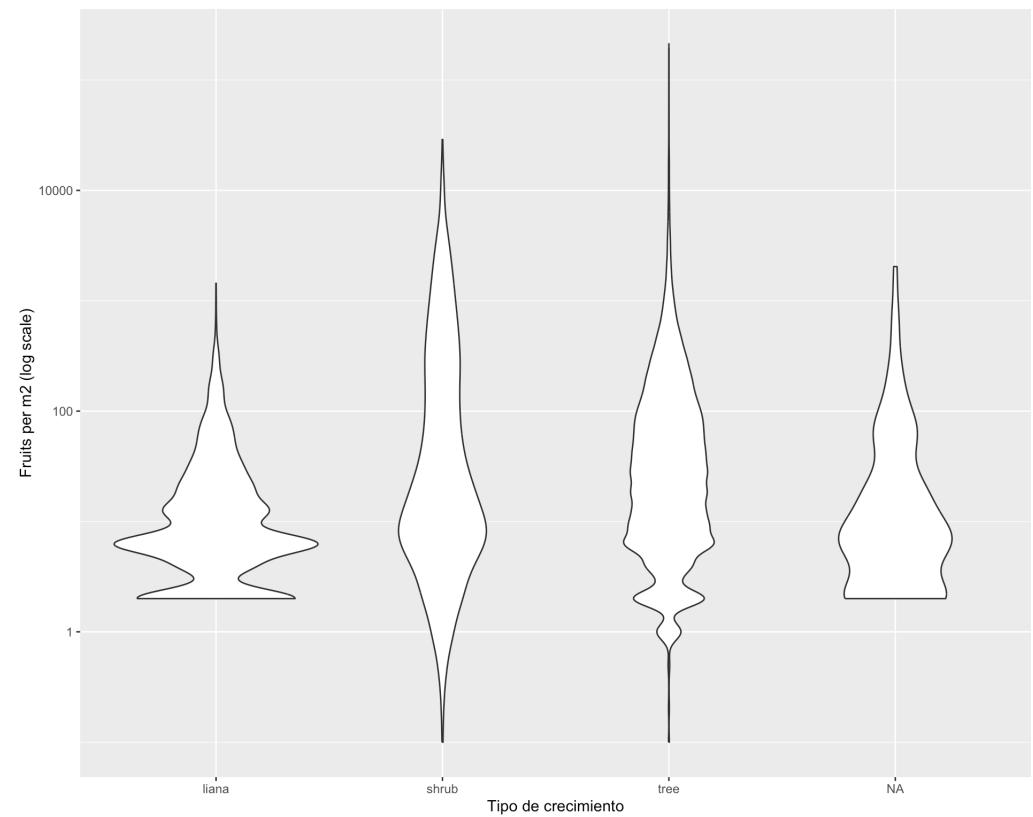
# Ejercicio 1

Con el dataset completo (dt) genera un violin plot del número de frutos por m<sup>2</sup> (fruits) en escala logarítmica para los distintos tipos de crecimiento (growth\_form)

# Ejercicio 1

Con el dataset completo (dt) genera un violin plot del número de frutos por m<sup>2</sup> (fruits) en escala logarítmica para los distintos tipos de crecimiento (growth\_form)

```
ggplot(dt,
       aes(x = growth_form,
            y = fruits)) +
  geom_violin() +
  labs(x = "Tipo de crecimiento",
       y = "Fruits per m2 (log scale)") -
  scale_y_log10()
```



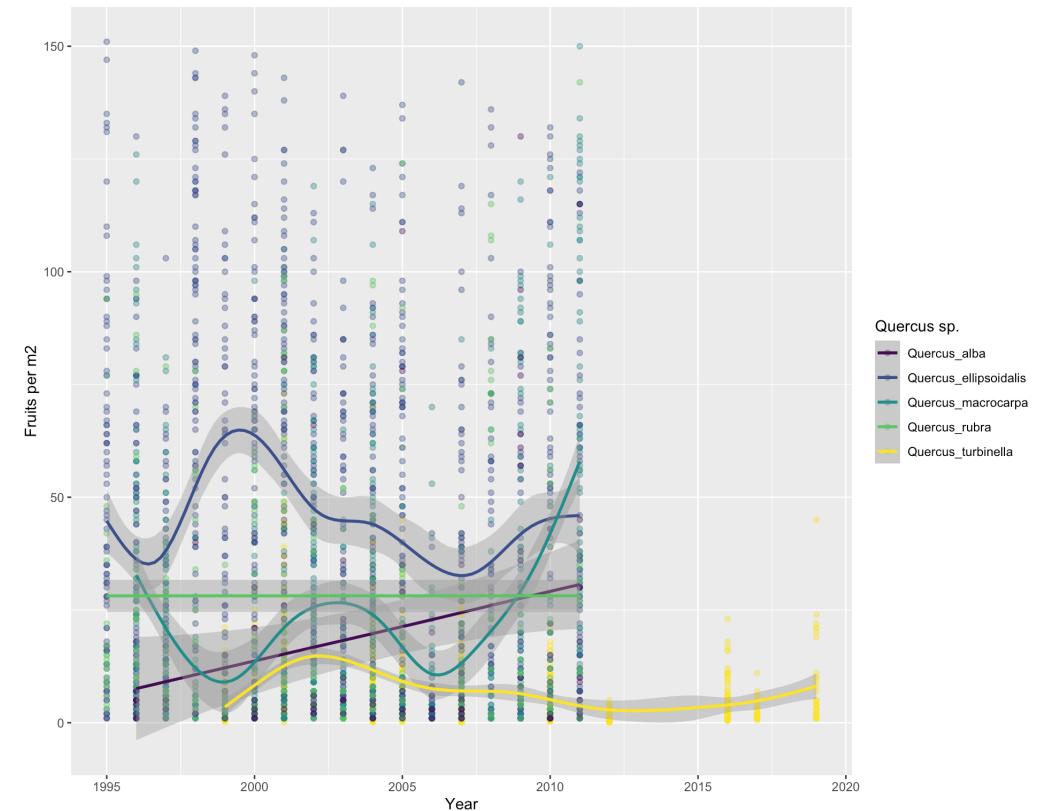
## Ejercicio 2

Para las distintas especies del género *Quercus*, sacar una tendencia del número de frutos por m<sup>2</sup> a lo largo de los años.

# Ejercicio 2

Para las distintas especies del género *Quercus*, sacar una tendencia del número de frutos por m<sup>2</sup> a lo largo de los años.

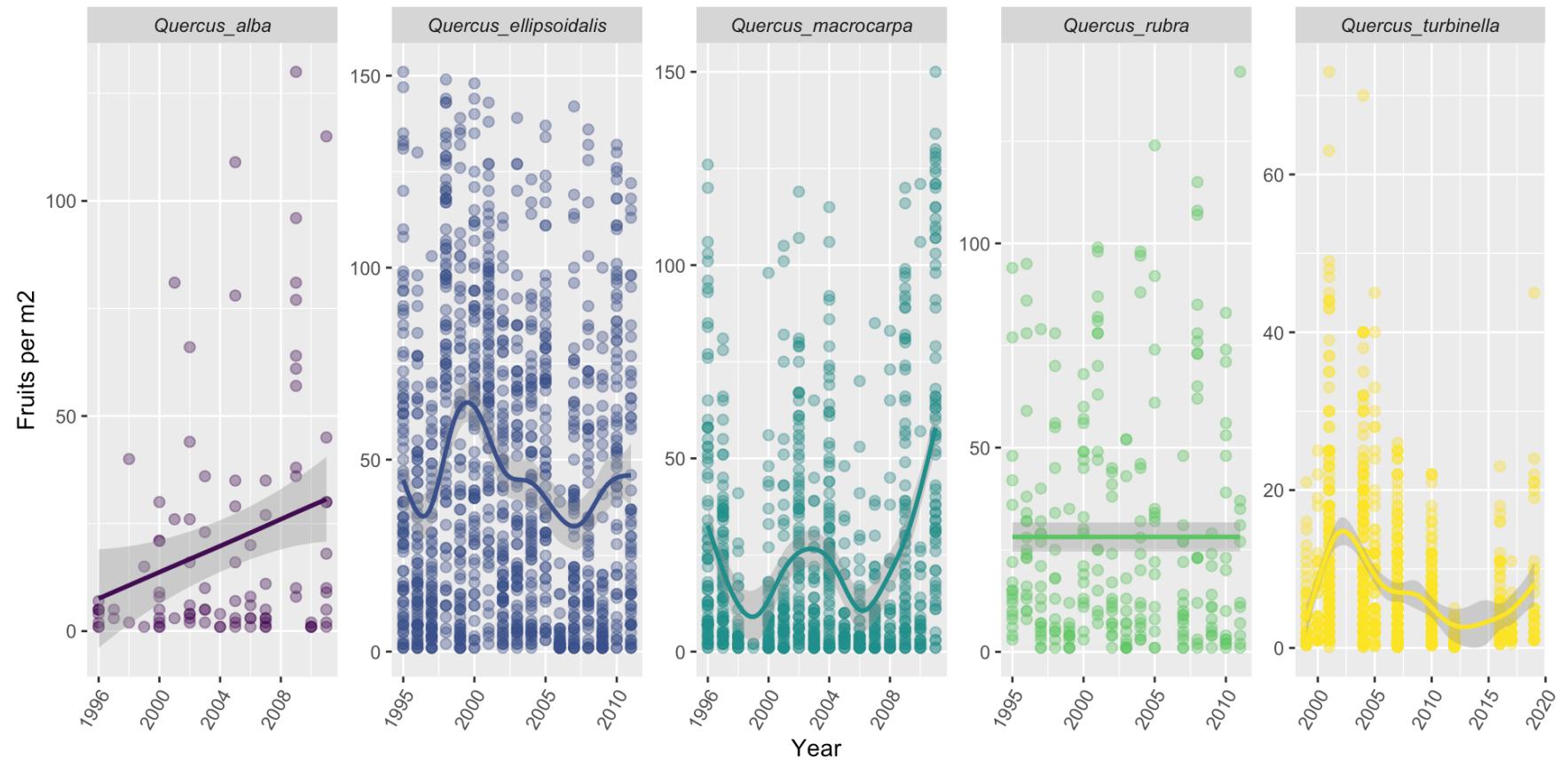
```
dt |> filter(genus == "Quercus") |>  
  ggplot(aes(x = year,  
             y = fruits,  
             color = species_name)) +  
  geom_point(alpha = 0.4) +  
  geom_smooth() +  
  scale_color_viridis_d() +  
  labs(x = "Year",  
       y = "Fruits per m2",  
       color = "Quercus sp.")
```



# Ejercicio 2

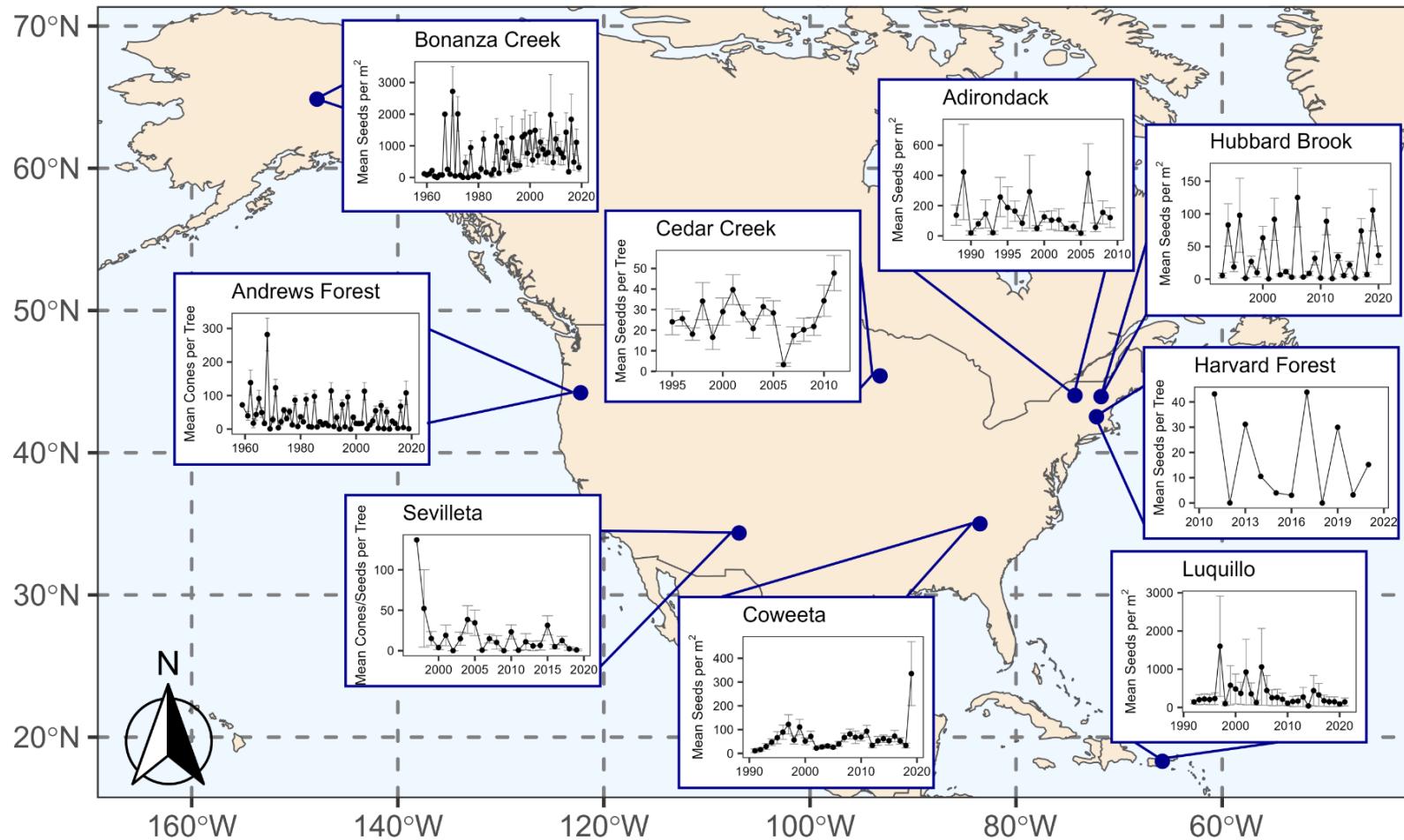
```
dt |> filter(genus == "Quercus") |>
  ggplot(aes(x = year,
             y = fruits,
             color = species_name)) +
  geom_point(alpha = 0.4, size = 2) +
  geom_smooth() +
  scale_color_viridis_d() +
  facet_wrap(~species_name, scales = "free", nrow = 1) +
  theme(legend.position = "none",
        axis.text.x = element_text(angle = 60, hjust = 1),
        strip.text = element_text(face = "italic")) +
  labs(x = "Year",
       y = "Fruits per m2)
```

# Ejercicio 2



# Ejercicio 3

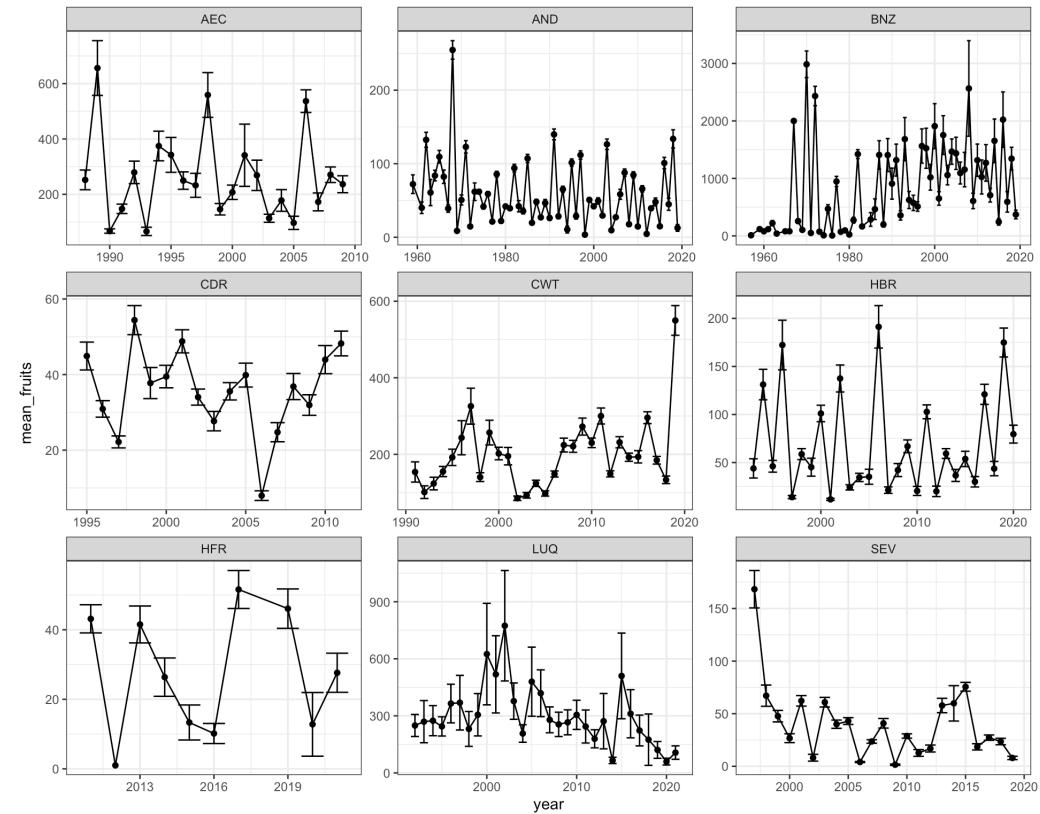
Replica las gráficas de esta figura (lo más similar posible):



# Ejercicio 3

Alternativa 1 - Sacar primero la media y la SE del número de frutos y luego graficar

```
dt |>  
  #remove outlier year  
  filter(!(site == "BNZ" & year == 1958))  
  group_by(site, year) |>  
  summarise(mean_fruits = mean(fruits, na.rm = TRUE),  
            se_fruits = sd(fruits, na.rm = TRUE)/sqrt(n() - 1))  
  ggplot(aes(x = year, y = mean_fruits)) +  
    geom_point() +  
    geom_errorbar(aes(ymin = mean_fruits - se_fruits,  
                      ymax = mean_fruits + se_fruits)) +  
    geom_line() +  
    facet_wrap(~site, scales = "free") +  
    theme_bw()
```



# Ejercicio 3

Alternativa 2 - Sacar la media y la SE de los frutos dentro de ggplot

```
dt |>  
  #remove outlier year  
  filter(!(site == "BNZ" & year == 1958))  
  ggplot(aes(x = year, y = fruits)) +  
    geom_point(stat = "summary", fun = "mean") +  
    geom_line(stat = "summary", fun = "mean") +  
    stat_summary(  
      fun.data = "mean_se",  
      geom = "errorbar",  
      width = 0.2) +  
    facet_wrap(~site, scales = "free") +  
    theme_bw()
```

