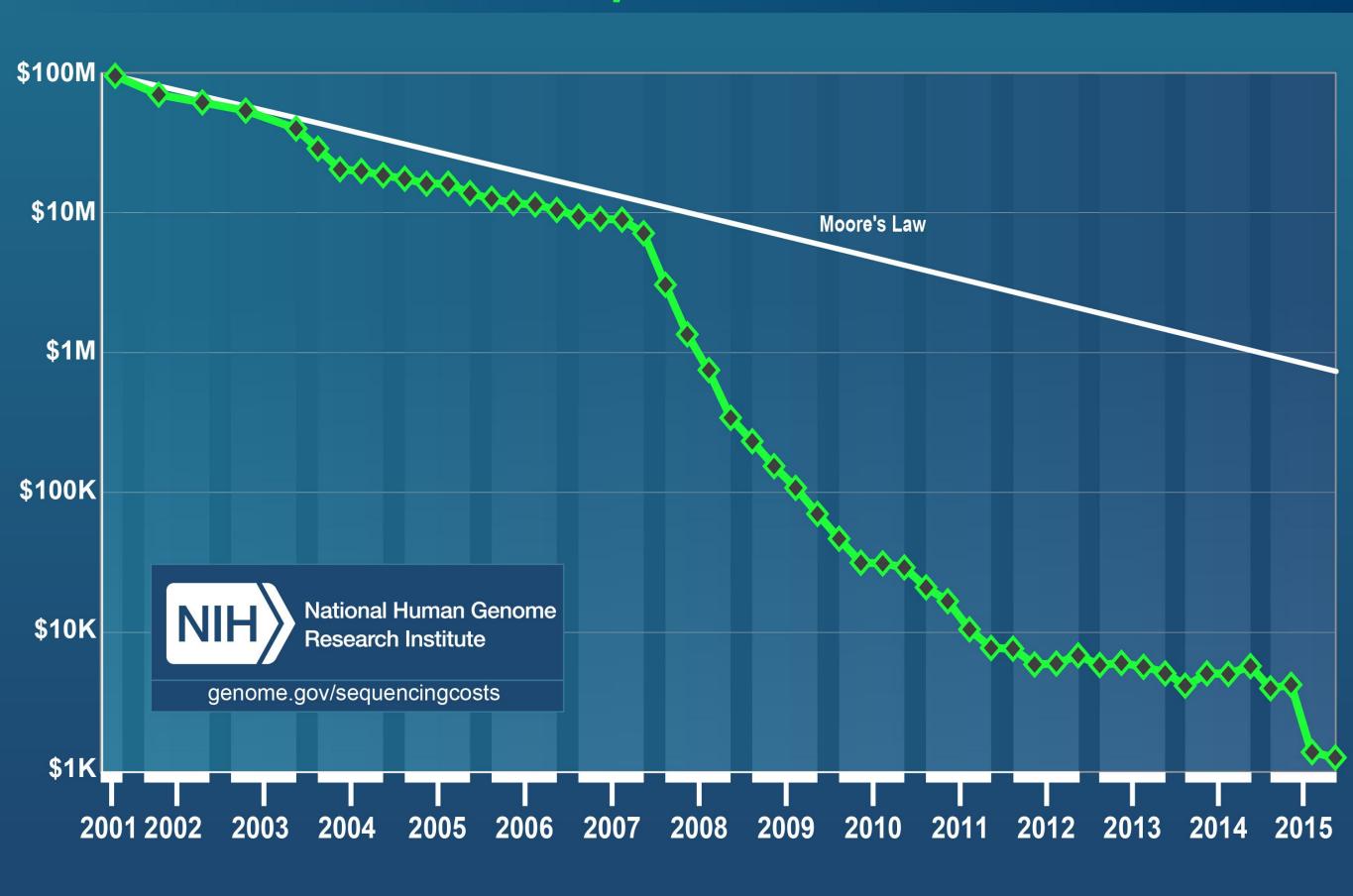
#### Plotting and Visualization with R

https://r4ds.had.co.nz/data-visualisation.html

https://is.gd/ggplot

#### Cost per Genome



#### Cost per Human Genome





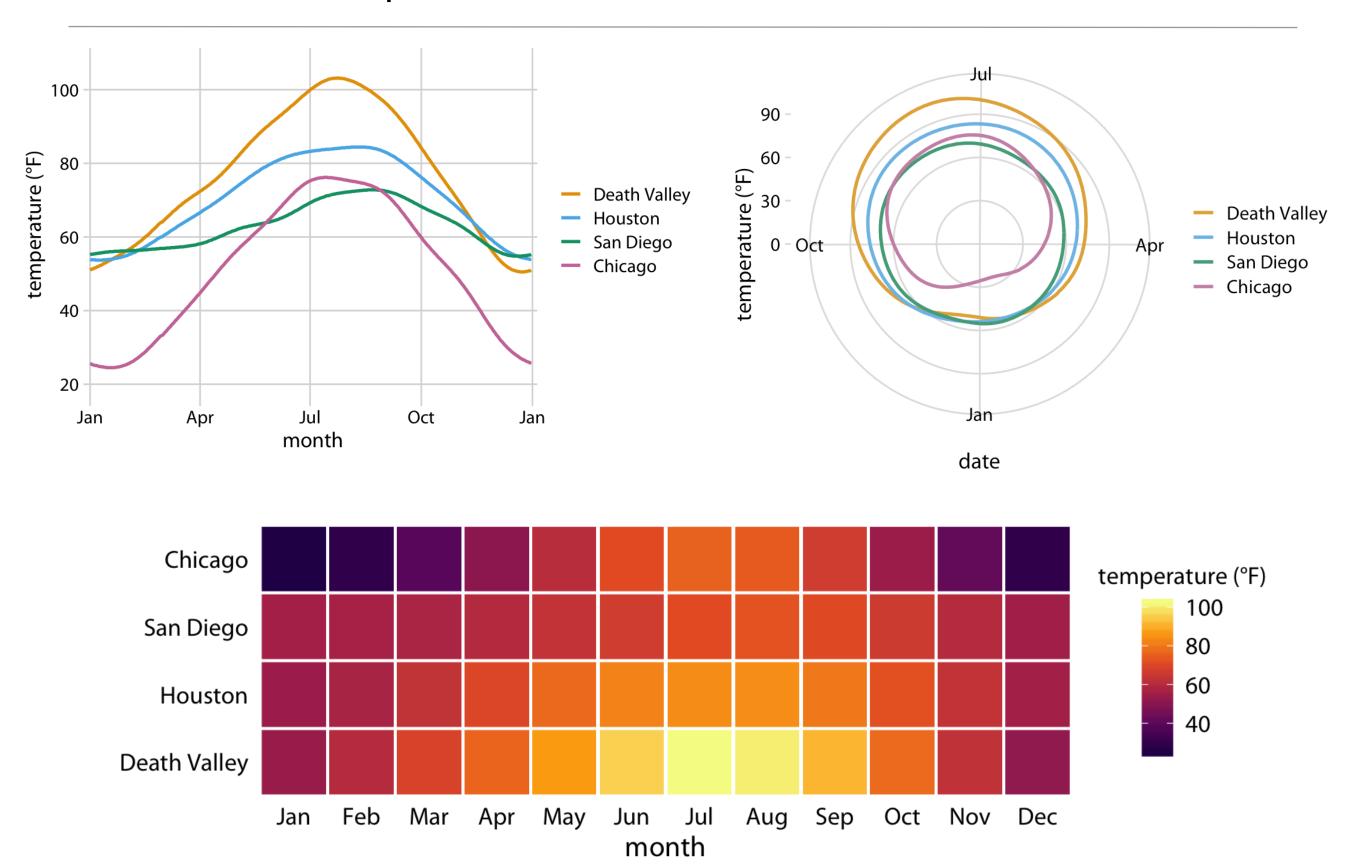




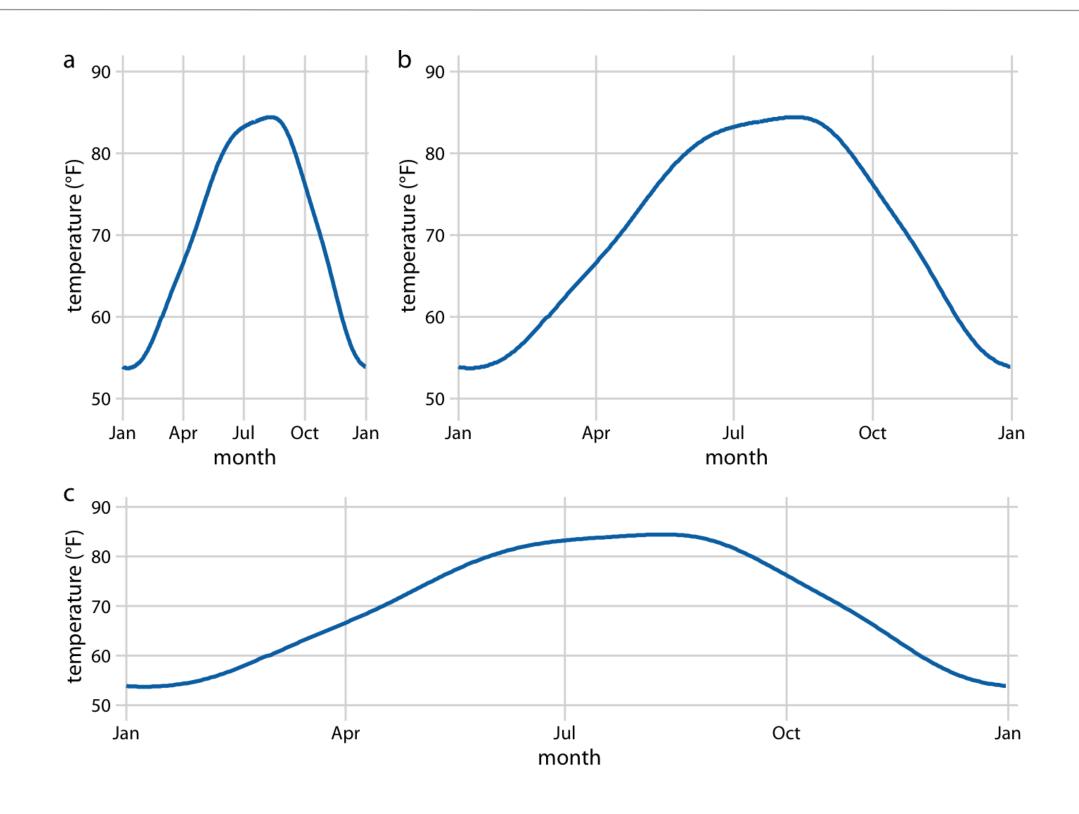
Fundamentals of Data visualization



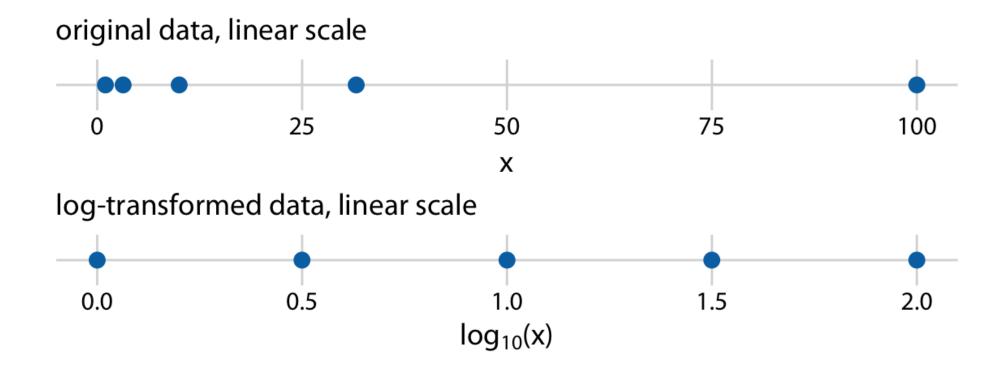
#### Alternative representations



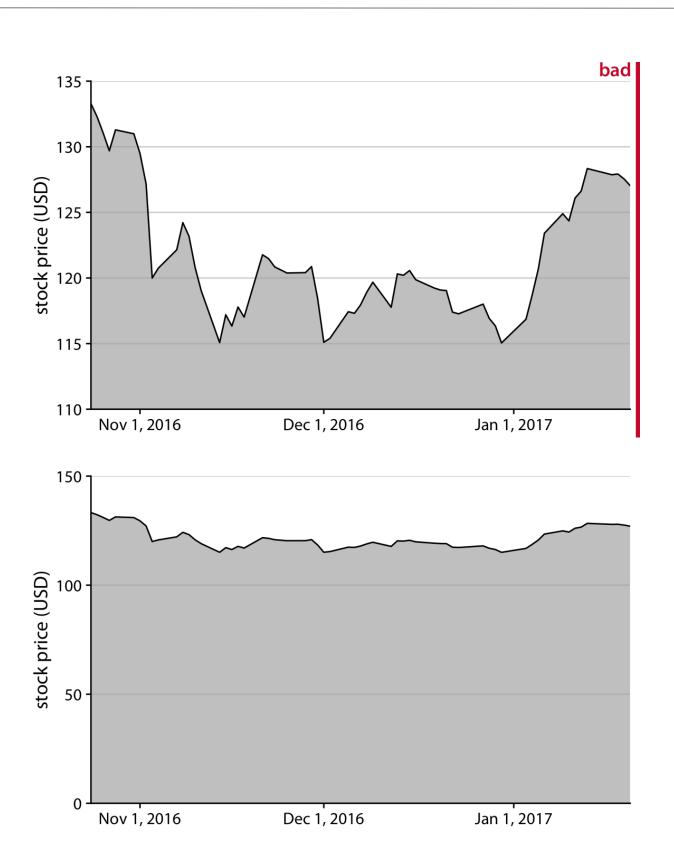
## Aspect ratio



# Scaling axes



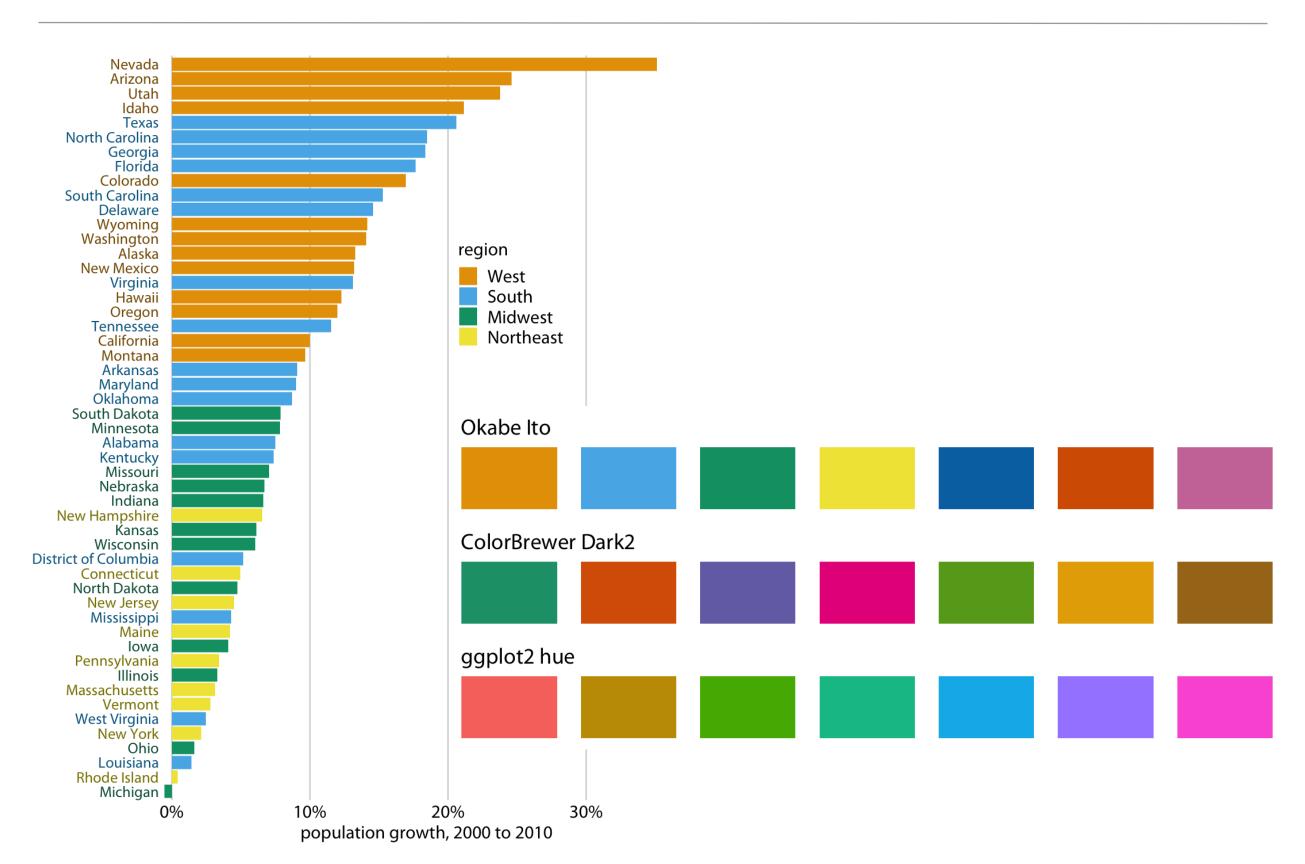
# Scaling axes



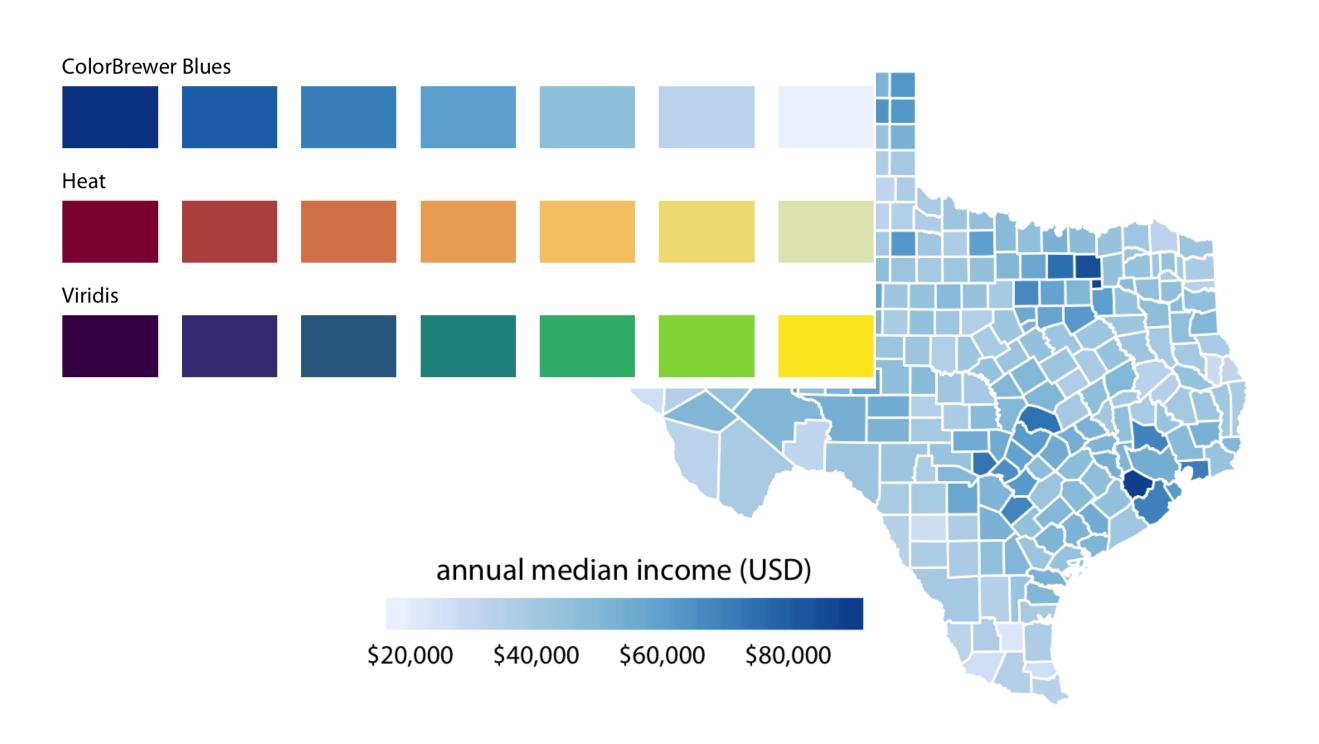
#### Color scales – three uses

- 1) To distinguish between groups
- 2) To represent data values
- 3) To highlight

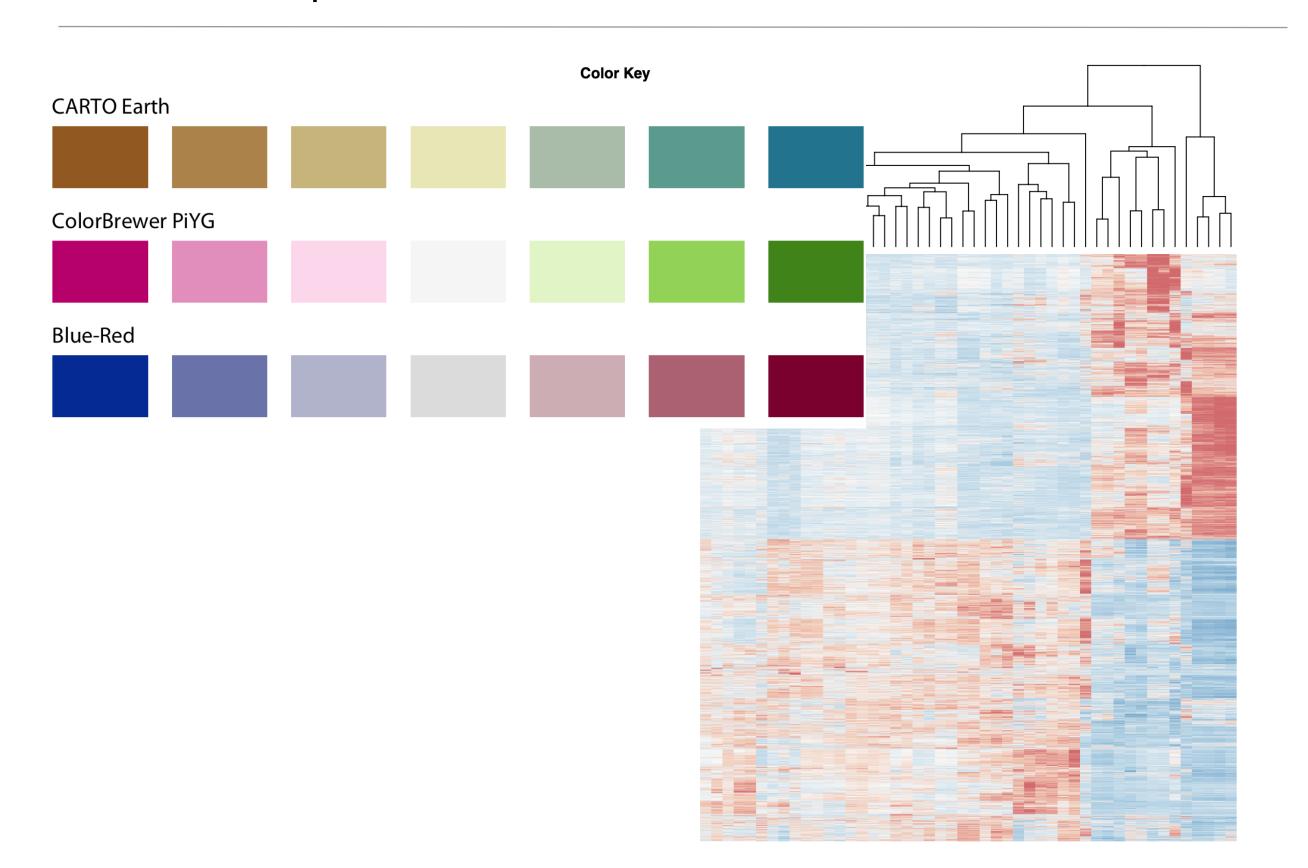
## Color to distinguish



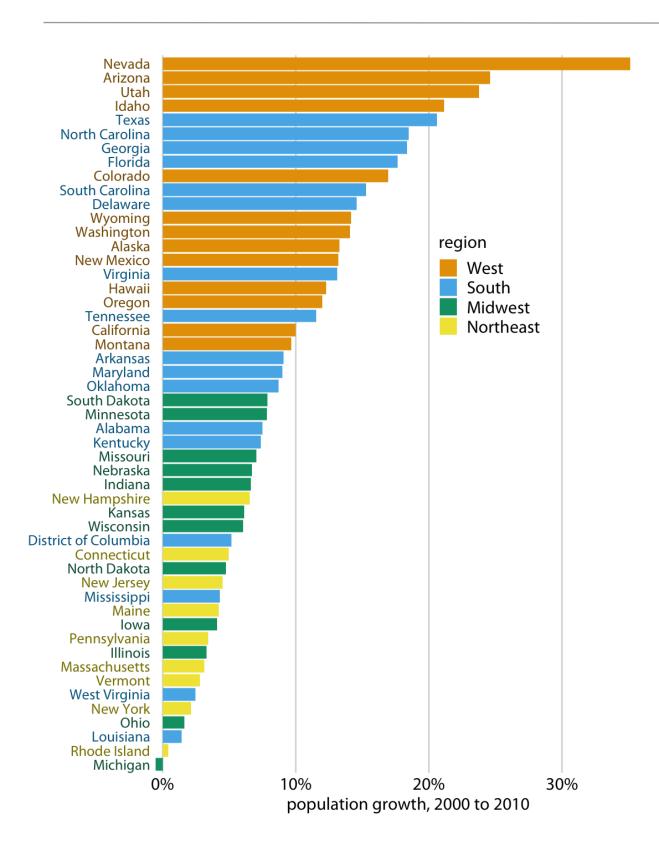
## Color to represent data values



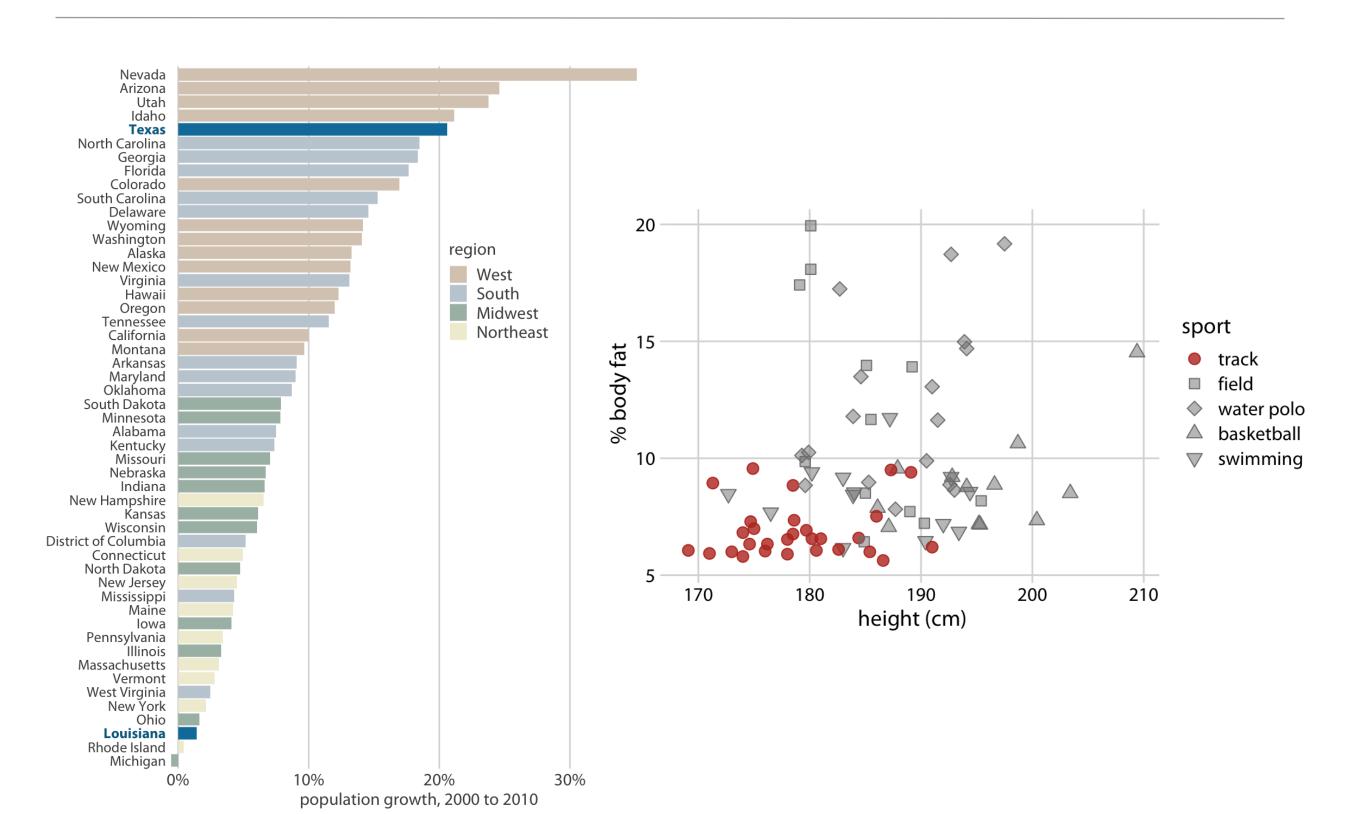
## Color to represent data values

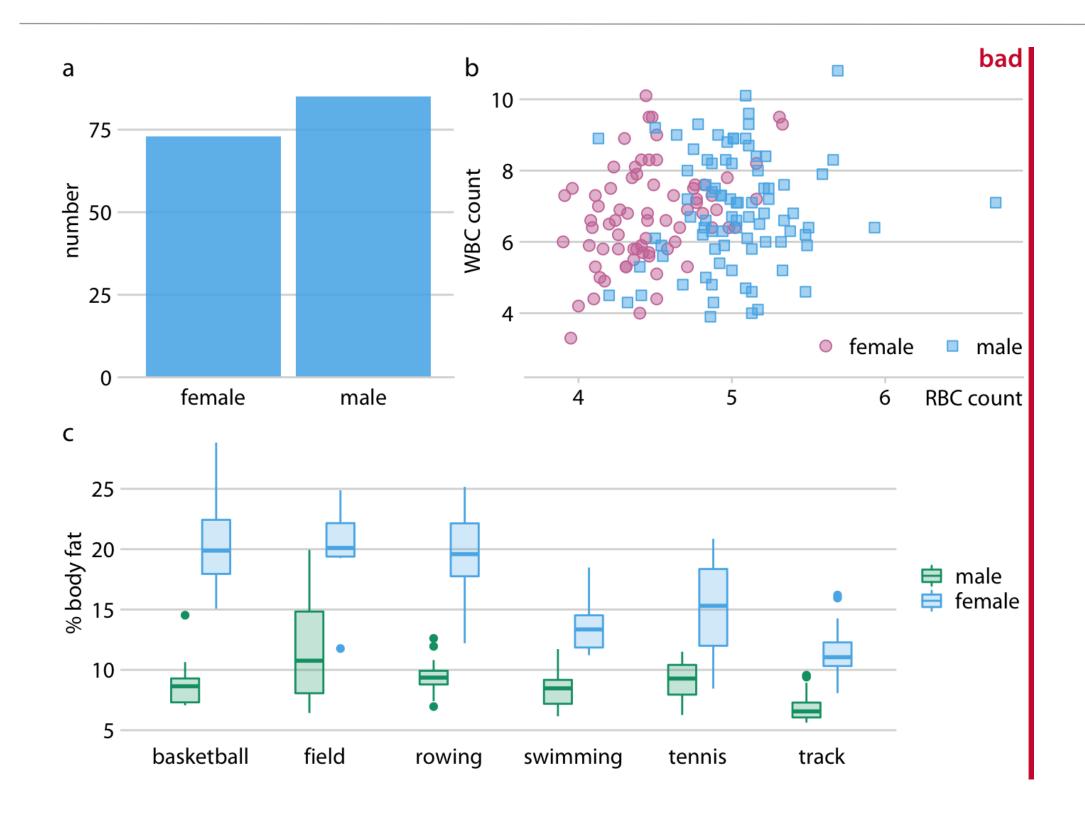


## Color to highlight

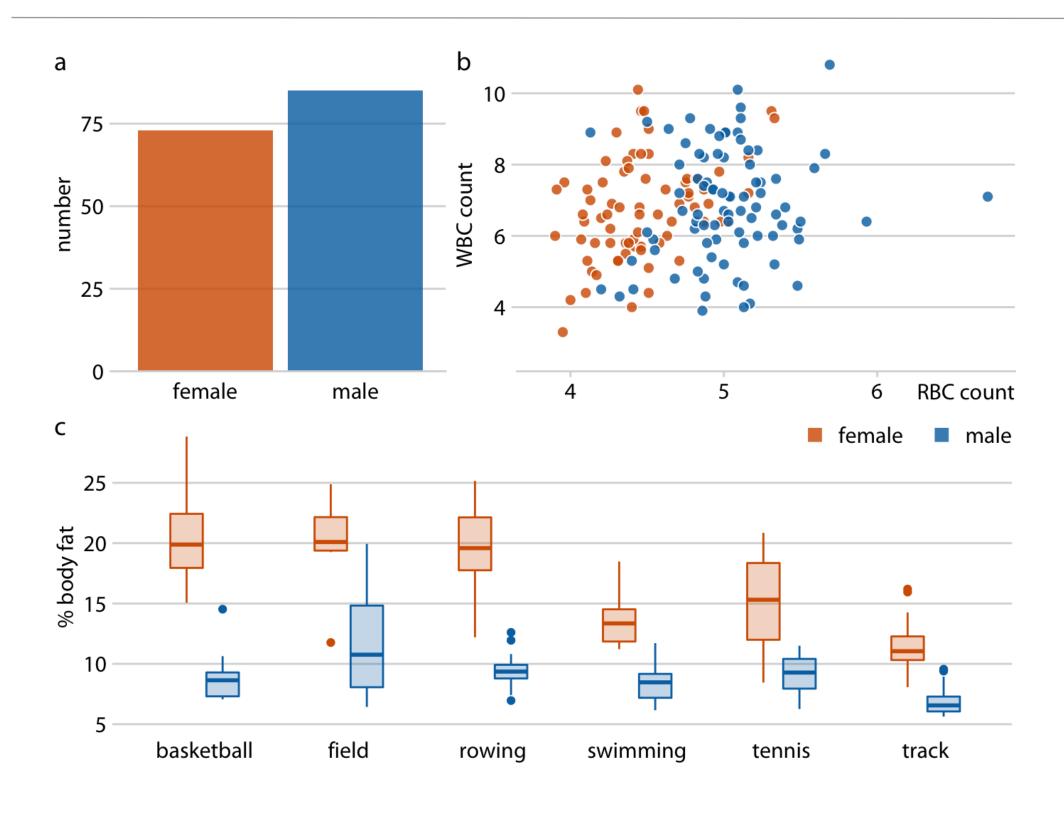


### Color to highlight

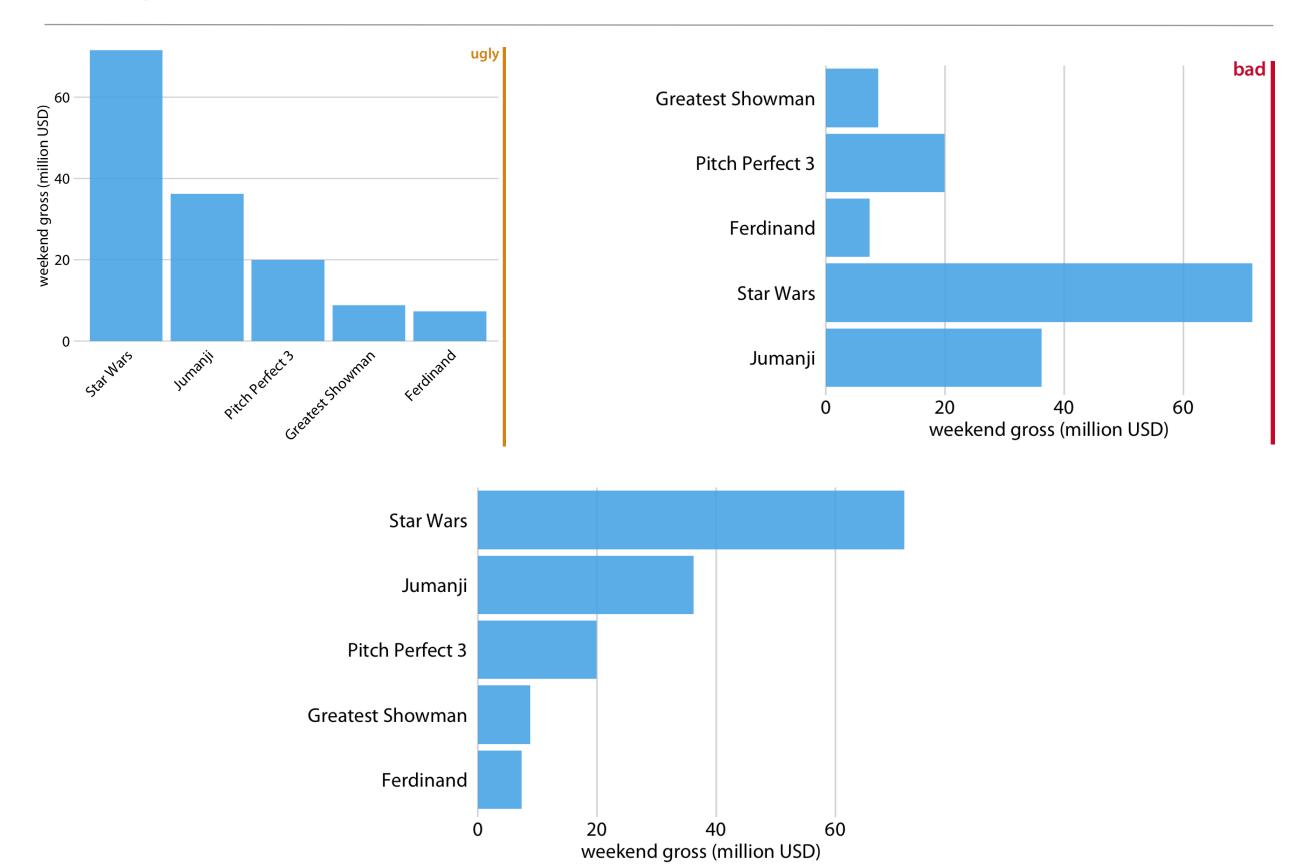




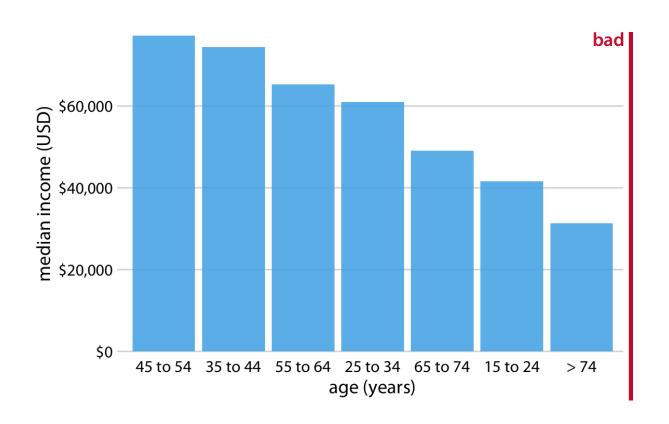
### Be consistent across panels and figures

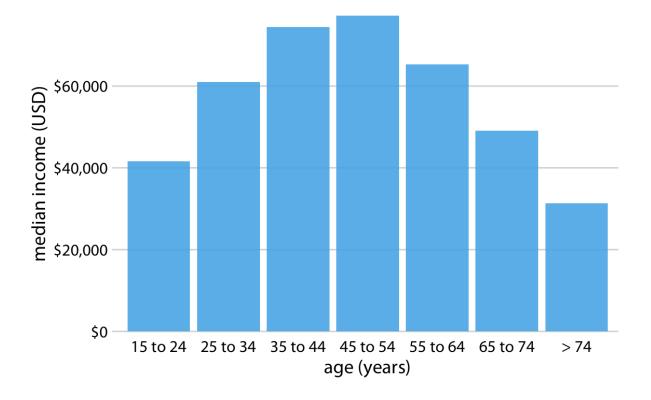


## Bar plots

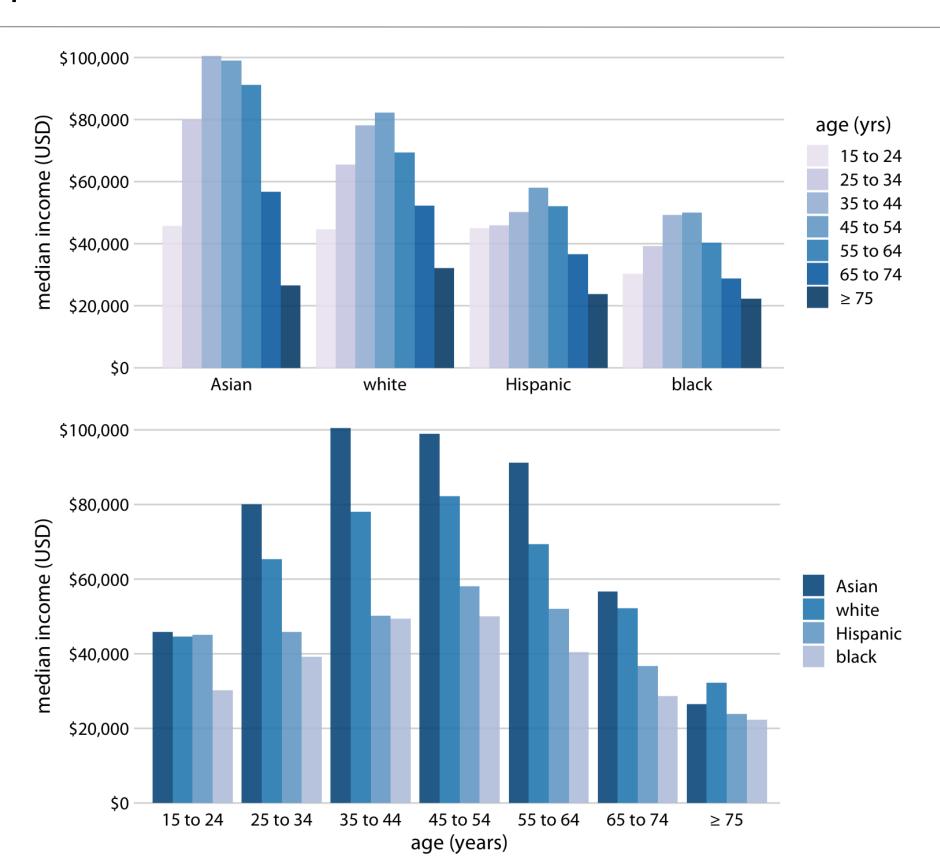


## Bar plots

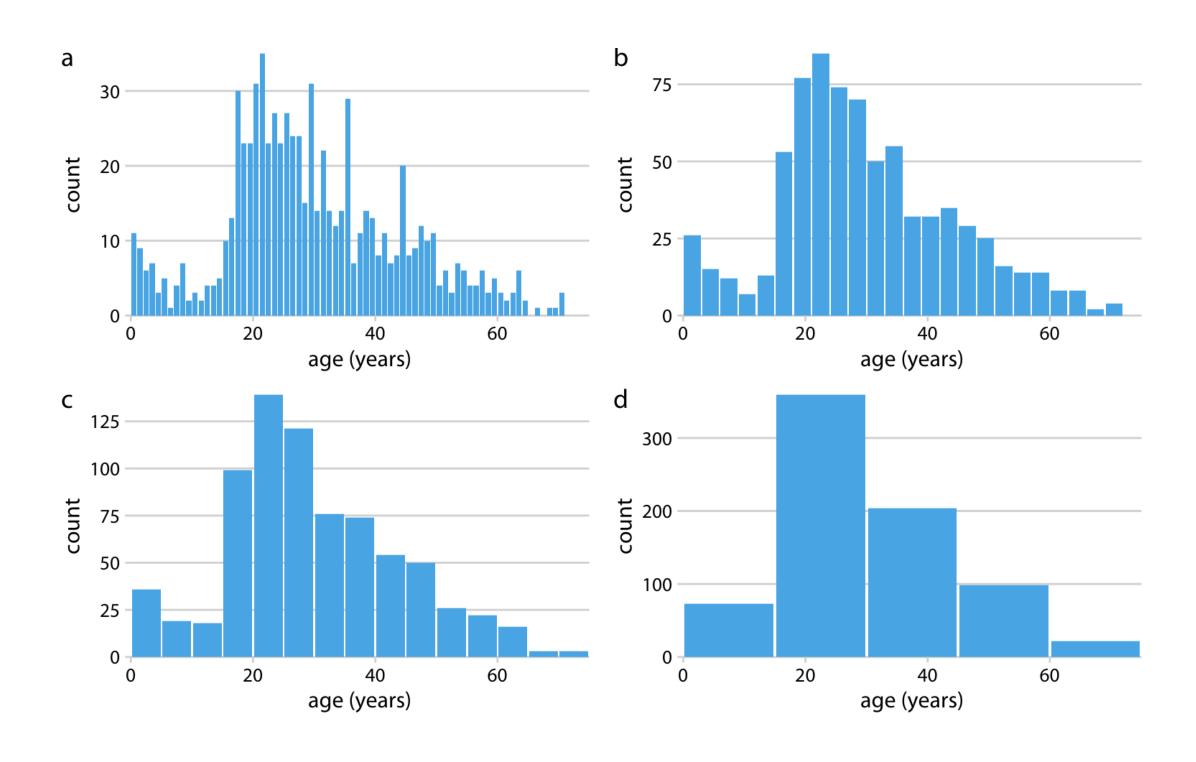




### Grouped Bar Plots



## Granularity of your data



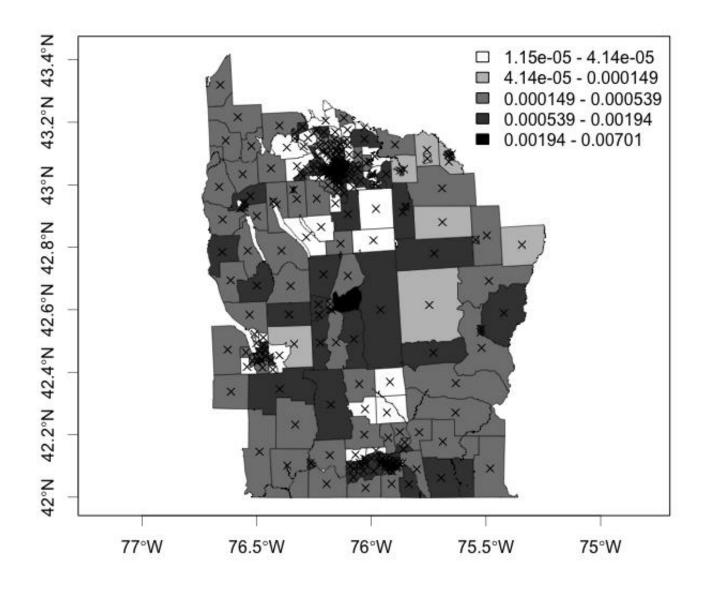
Advanced and Other Applications in R

#### Spatial Epidemiology and Maps

```
library(SpatialEpi)

data(NYleukemia)
sp.obj <- NYleukemia$spatial.polygon
centroids <- latlong2grid(NYleukemia$geo[, 2:3])
population <- NYleukemia$data$population
cases <- NYleukemia$data$cases

plotmap(cases/population, sp.obj, log=TRUE, nclr=5)
points(grid2latlong(centroids), pch=4)
```



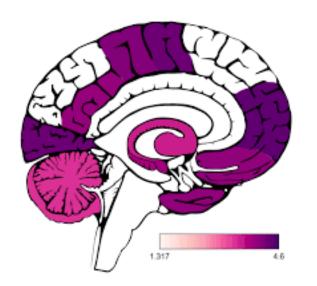
### Anatomical Mapping

#### CerebroViz

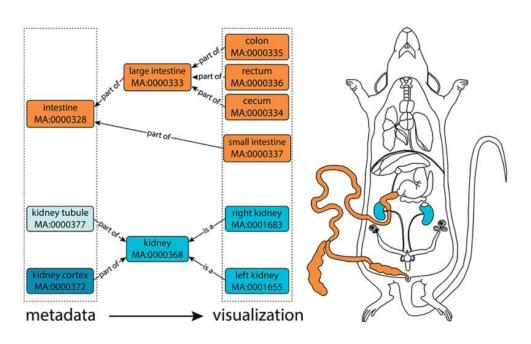
https://github.com/ethanbahl/cerebroViz

COMICS

https://github.com/y-popov/COMICS

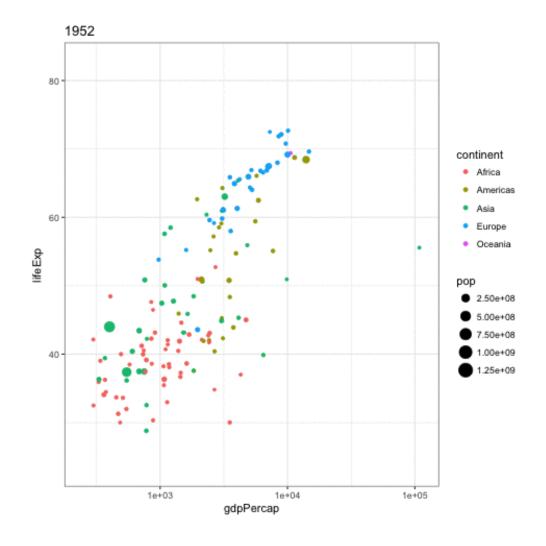


library("cerebroViz") data(cerebroEx) head(cerebroEx)[, c(1:7)]



# Animation / Interactivity

Indicator Name	2011	2012	2013	2014	2015	2016	Average	Improvement
Prevalence of Obesity	19.1	23.6	23.3	20.5	24.0	23.2	22.28	-21.47
Prevalence of Tobacco Use	17.4	15.0	15.3	12.2	16.6	16.7	15.53	4.02
Prevalence of Cardiovascular Disease	5.0	4.9	1.5	4.4	4.9	6.2	4.48	-24.00
Prevalence of Diabetes	8.0	7.2	9.3	7.2	7.5	10.4	8.27	-30.00



## Expose yourself to lots of ideas!

