

Advanced Visualizations

Statistical Computing & Programming

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Supplementary materials

Companion videos

- Using `ggcorrplot()` and `geom_parliment()`
- Package `patchwork`
- Creating animations
- Creating interactive plots

Additional resources

- Top 50 `ggplot2` visualizations
- Extend `ggplot2` by creating your own stat, geom, and theme

ggplot2 extensions

Packages

For these slides we will use the following packages.

```
library(tidyverse)
library(gapminder)    # some data
library(ggcorrplot)   # correlogram plots
library(ggpol)         # parliament plots and more
library(patchwork)     # combining plots
library(gganimate)     # animations
library(ggiraph)       # interactive plots
```

Install any CRAN packages you do not have with

```
install.packages("package_name"). Package patchwork needs to be installed  
by running devtools::install_github("thomasp85/patchwork").
```

Code not shown for plots is available in the presentation notes. Press P.

Data: Flint water crisis

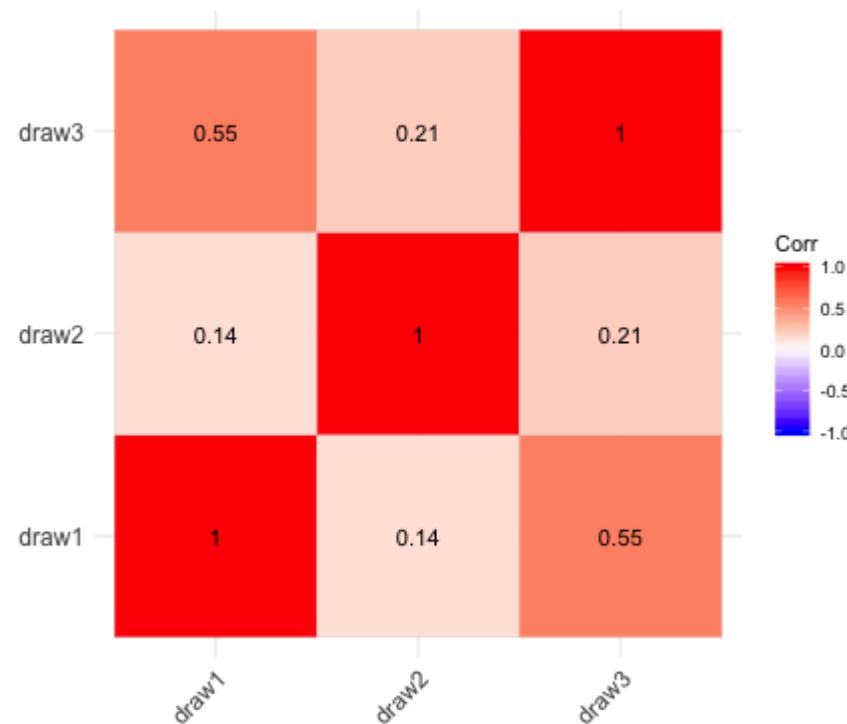
```
flint <- read_csv("http://www2.stat.duke.edu/~sms185/data/health/flint.csv")
```

```
#> # A tibble: 271 x 6
#>   id    zip  ward draw1  draw2  draw3
#>   <dbl> <dbl> <dbl>  <dbl>  <dbl>  <dbl>
#> 1 1     48504     6  0.344  0.226  0.145
#> 2 2     48507     9  8.13   10.8   2.76
#> 3 4     48504     1  1.11   0.11   0.123
#> 4 5     48507     8  8.01   7.45   3.38
#> 5 6     48505     3  1.95   0.048  0.035
#> 6 7     48507     9  7.2    1.4    0.2
#> 7 8     48507     9  40.6   9.73   6.13
#> 8 9     48503     5  1.1    2.5    0.1
#> 9 12    48507     9  10.6   1.04   1.29
#> 10 13    48505     3  6.2    4.2    2.3
#> # ... with 261 more rows
```

Correlogram: ggcorrplot

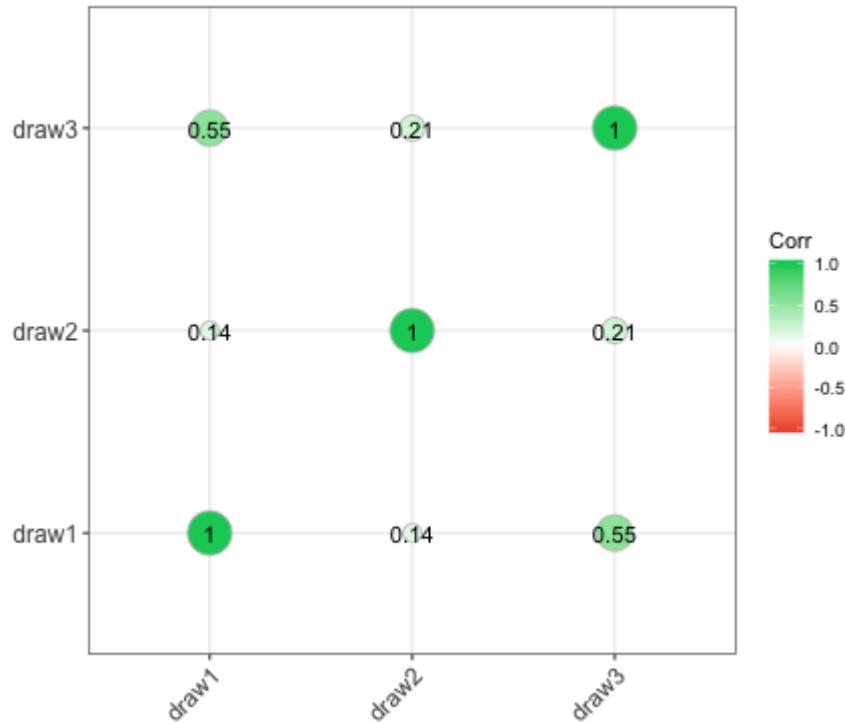
Full matrix

```
corr_mat <- round(cor(flint[, c("draw1", "draw2", "draw3")]), 2)  
ggcorrplot(corr = corr_mat, lab = TRUE)
```



Full matrix

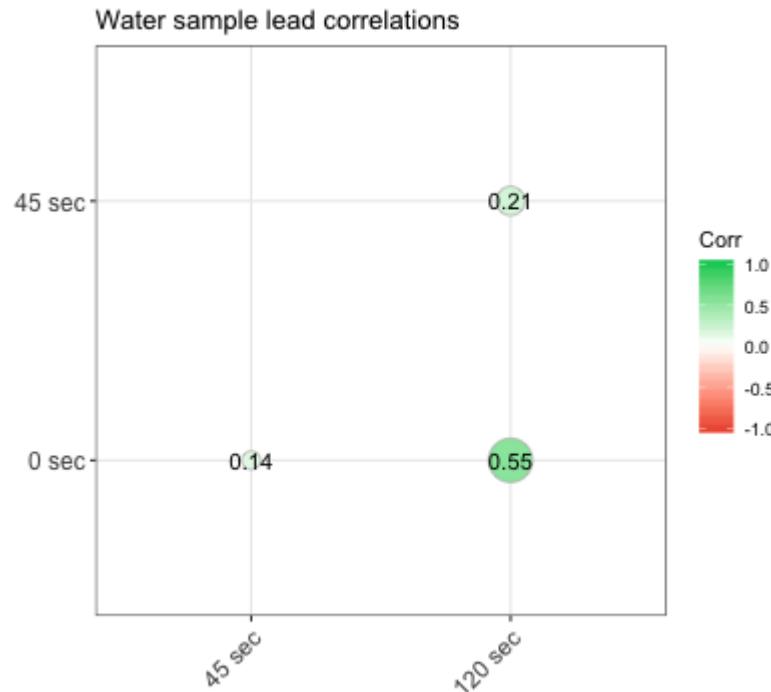
```
ggcorrplot(corr = corr_mat, method = "circle", type = "full", lab = TRUE,  
          colors = c("tomato2", "white", "springgreen3"),  
          ggtheme = theme_bw)
```



Lower triangular

```
lbl <- c("0 sec", "45 sec", "120 sec")

ggcorrplot(corr = corr_mat, method = "circle", type = "lower", lab = TRUE,
           colors = c("tomato2", "white", "springgreen3"), ggtheme = theme_bw) +
  labs(title = "Water sample lead correlations") +
  scale_x_discrete(labels = lbl[2:3]) +
  scale_y_discrete(labels = lbl[1:2])
```



Parliment plots: ggplot

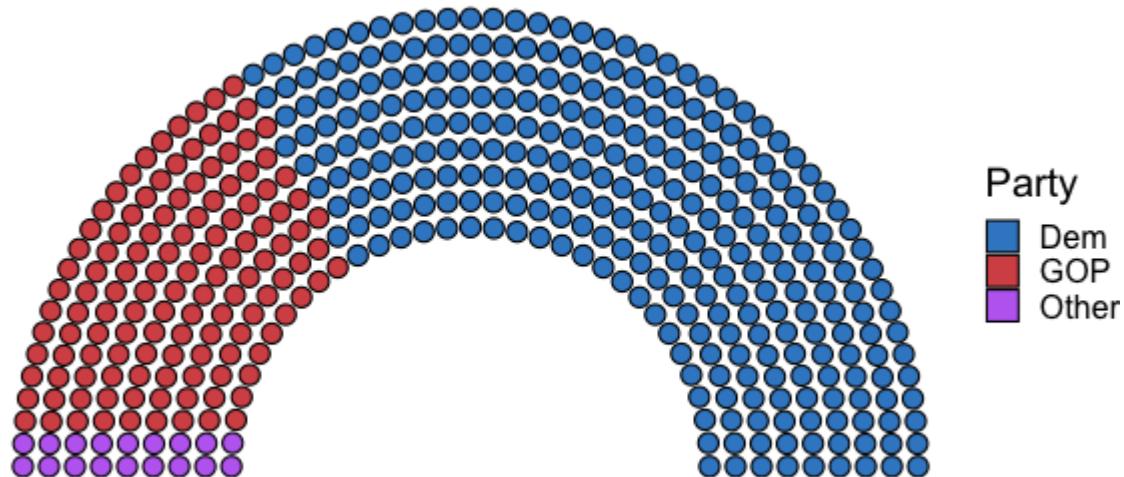
Data: Congressional seats

```
congress <- read_csv("http://www2.stat.duke.edu/~sms185/data/politics/cor  
congress
```

```
#> # A tibble: 432 x 5  
#>   year_start year_end party  branch  seats  
#>       <dbl>     <dbl> <chr>   <chr>    <dbl>  
#> 1 1913      1915 dem    house    290  
#> 2 1913      1915 dem    senate    51  
#> 3 1913      1915 gop    house    127  
#> 4 1913      1915 gop    senate    44  
#> 5 1913      1915 other  house    18  
#> 6 1913      1915 other  senate    1  
#> 7 1913      1915 vacant house    NA  
#> 8 1913      1915 vacant senate   NA  
#> 9 1915      1917 dem    house    231  
#> 10 1915     1917 dem    senate    56  
#> # ... with 422 more rows
```

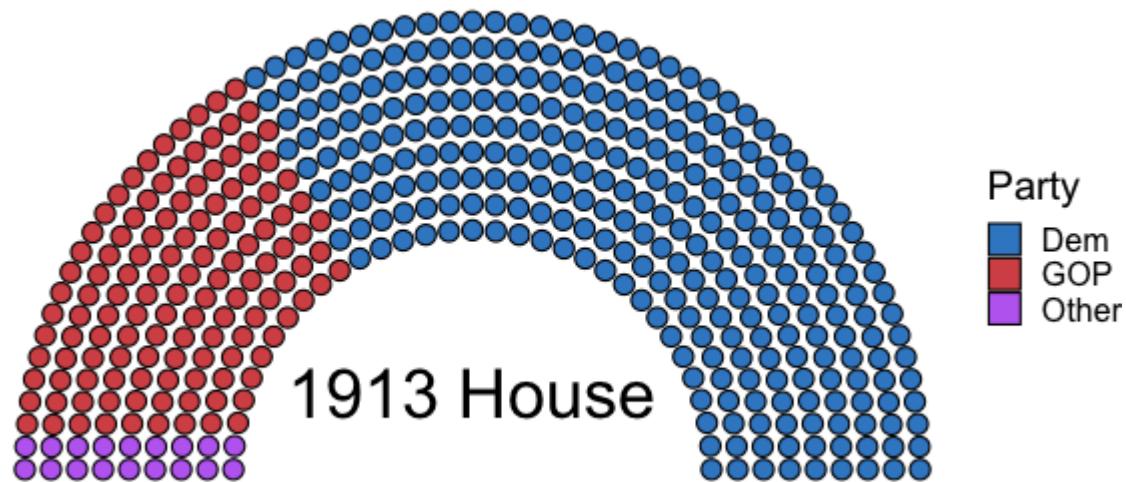
Parliment plot

```
ggplot(data = congress[congress$year_start == 1913 & congress$branch == "house", ]) +  
  geom_parliament(aes(seats = seats, fill = factor(party)), show.legend = TRUE, color = "black")  
  scale_fill_manual(values = c("#3A89CB", "#D65454", "#BF6FF0", "Grey"),  
                    labels = c("Dem", "GOP", "Other", "Vacant")) +  
  labs(fill = "Party") +  
  coord_fixed() +  
  theme_void(base_size = 20)
```



Annotation

```
ggplot(data = congress[congress$year_start == 1913 & congress$branch == "house", ]) +  
  geom_parliament(aes(seats = seats, fill = factor(party)), show.legend = TRUE, color = "black")  
  scale_fill_manual(values = c("#3A89CB", "#D65454", "#BF6FF0", "Grey"),  
                    labels = c("Dem", "GOP", "Other", "Vacant")) +  
  annotate("text", x = 0, y = .5, label = "1913 House", size = 12) +  
  labs(fill = "Party") +  
  coord_fixed() +  
  theme_void(base_size = 20)
```



Package ggpolt

- Package ggpolt supports a few other geom functions:
 - geom_arcbar(),
 - geom_bartext(),
 - geom_circle(),
 - geom_tshighlight(),
 - geom_boxjitter().
- See <https://github.com/erocoar/ggpolt>

Organizing plots: package patchwork

My function: plot_congress()

```
plot_congress <- function(data, year, leg_branch, legend = TRUE, text_size = 12) {  
  data %>%  
    filter(year_start == year, branch == leg_branch) %>%  
    ggplot() +  
    geom_parliament(aes(seats = seats, fill = factor(party)),  
                     show.legend = legend, color = "black") +  
    scale_fill_manual(values = c("#3A89CB", "#D65454", "#BF6FF0", "Grey"),  
                      labels = c("Dem", "GOP", "Other", "Vacant")) +  
    annotate("text", x = 0, y = .5, label = paste(year, leg_branch),  
            size = text_size) +  
    labs(fill = "Party") +  
    coord_fixed() +  
    theme_void(base_size = 20)  
}
```

Use package patchwork to organize multiple plots in a single window. No need to facet.

```
my_plot <- ggplot()  
class(my_plot)  
  
#> [1] "gg"       "ggplot"
```

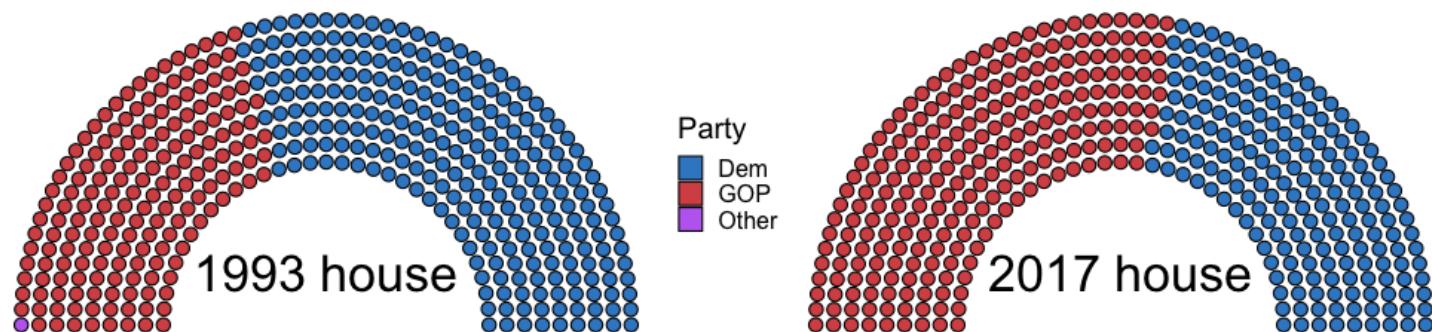
Plot creation

```
ph_1993 <- plot_congress(congress, 1993, "house")  
ph_2001 <- plot_congress(congress, 2001, "house", legend = FALSE)  
ph_2009 <- plot_congress(congress, 2009, "house", legend = FALSE)  
ph_2017 <- plot_congress(congress, 2017, "house", legend = FALSE)
```

Object ph_1993 has a legend, the rest do not.

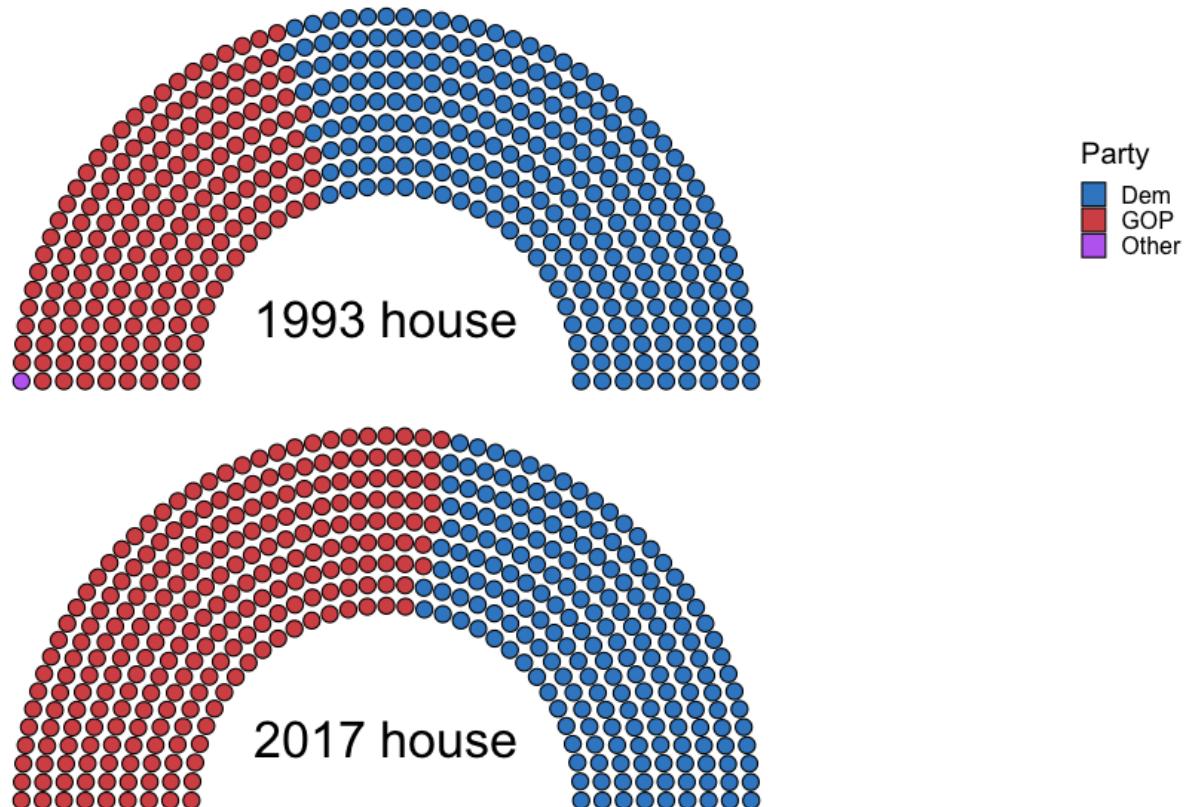
Horizontal patchwork

ph_1993 + ph_2017



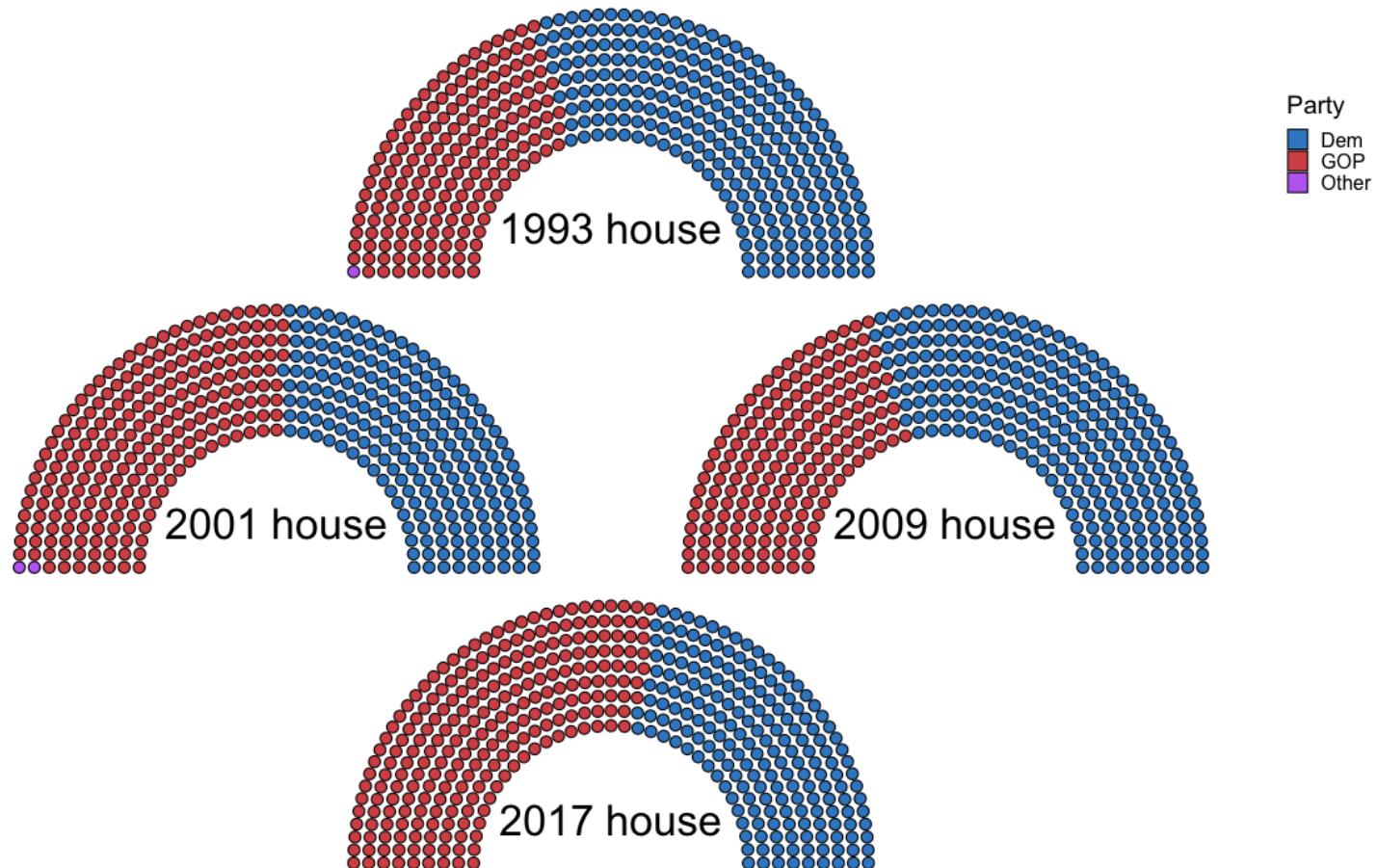
Vertical patchwork

```
ph_1993 + ph_2017 + plot_layout(ncol = 1)
```

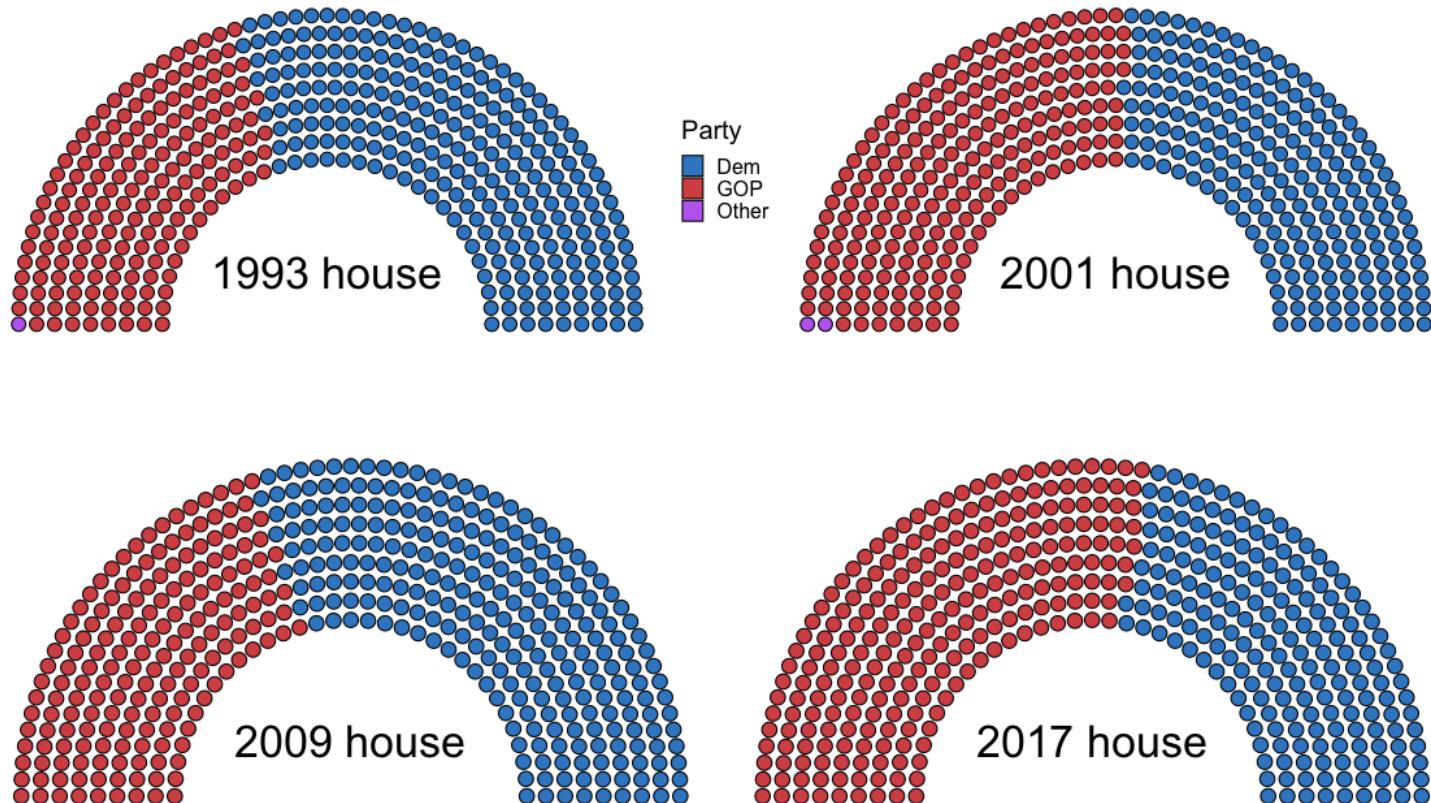


Group patchwork

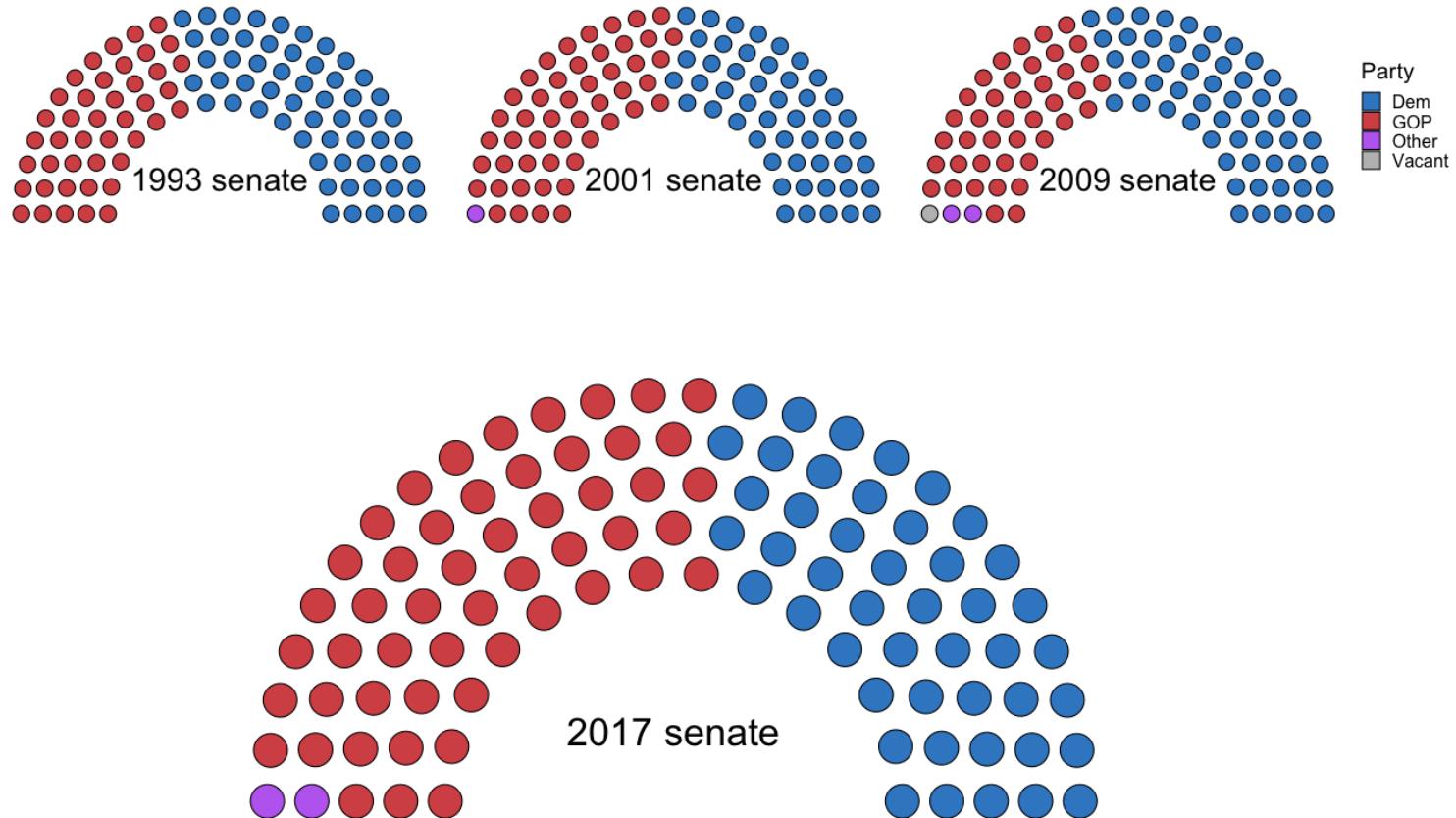
```
ph_1993 + (ph_2001 + ph_2009) + ph_2017 + plot_layout(ncol = 1)
```



(ph_1993 | ph_2001) / (ph_2009 | ph_2017)



(ps_1993 | ps_2001 | ps_2009) / ps_2017

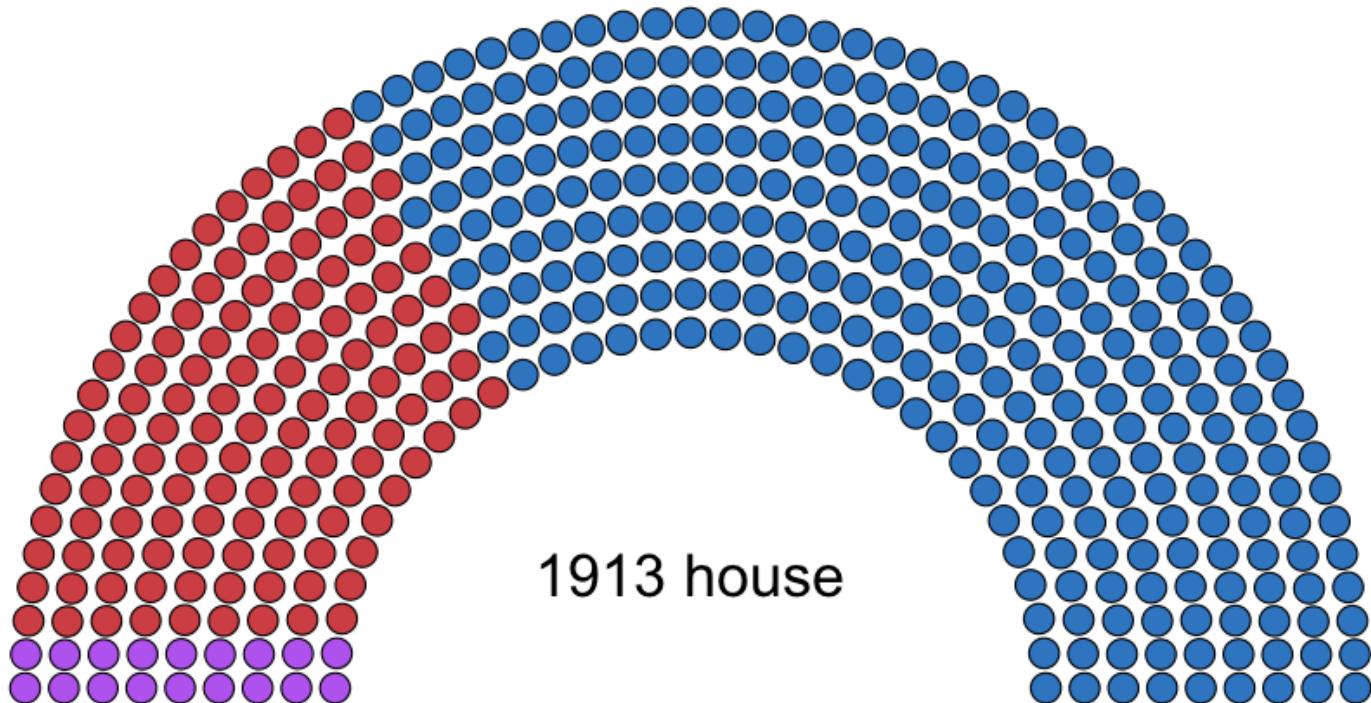


Package patchwork

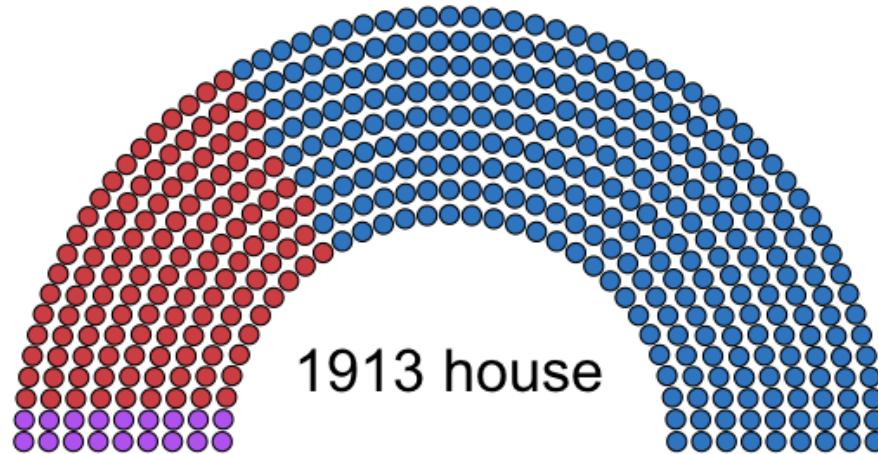
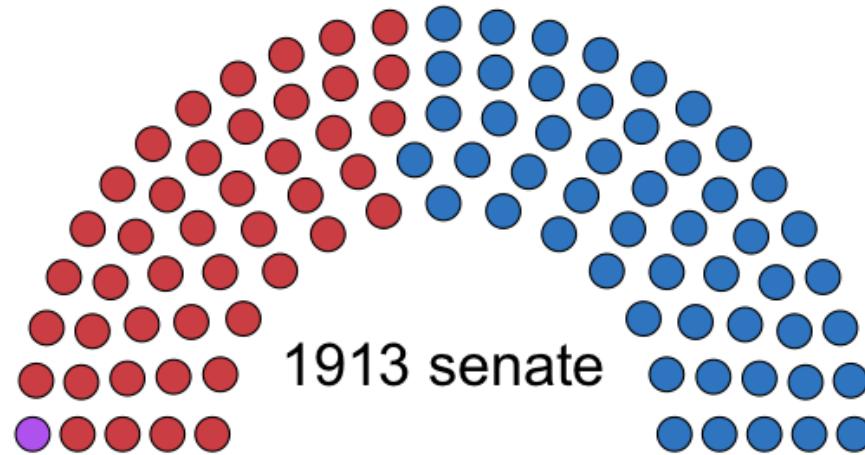
- Supports operators `+`, `-`, `|` (besides), `/` (over)
- Specify layouts and spacing with `plot_layout()`, `plot_spacer()`, respectively
- Add grouping with `{ }` or `()`
- Use `&` or `*` to add elements to all subplots, `*` only affects current nesting level
- See <https://github.com/thomasp85/patchwork>

GIF: gifski

Using gifski



Fast GIF with patchwork



Creating a GIF

1. Install `gifski` with `install.packages('gifski')`

2. Use chunk options

```
```{r animation.hook="gifski", interval = .75}  
```
```

3. Add code for plots in a loop

```
```{r animation.hook="gifski", interval = .75}  
for (i in seq(1913, 2019, 2)) {
 print({
 plot_congress(congress, year = i, leg_branch = "house", legend =
 })
}
```
```

4. To speed up future knits use chunk option `cache=TRUE`.

Animation: gganimate()

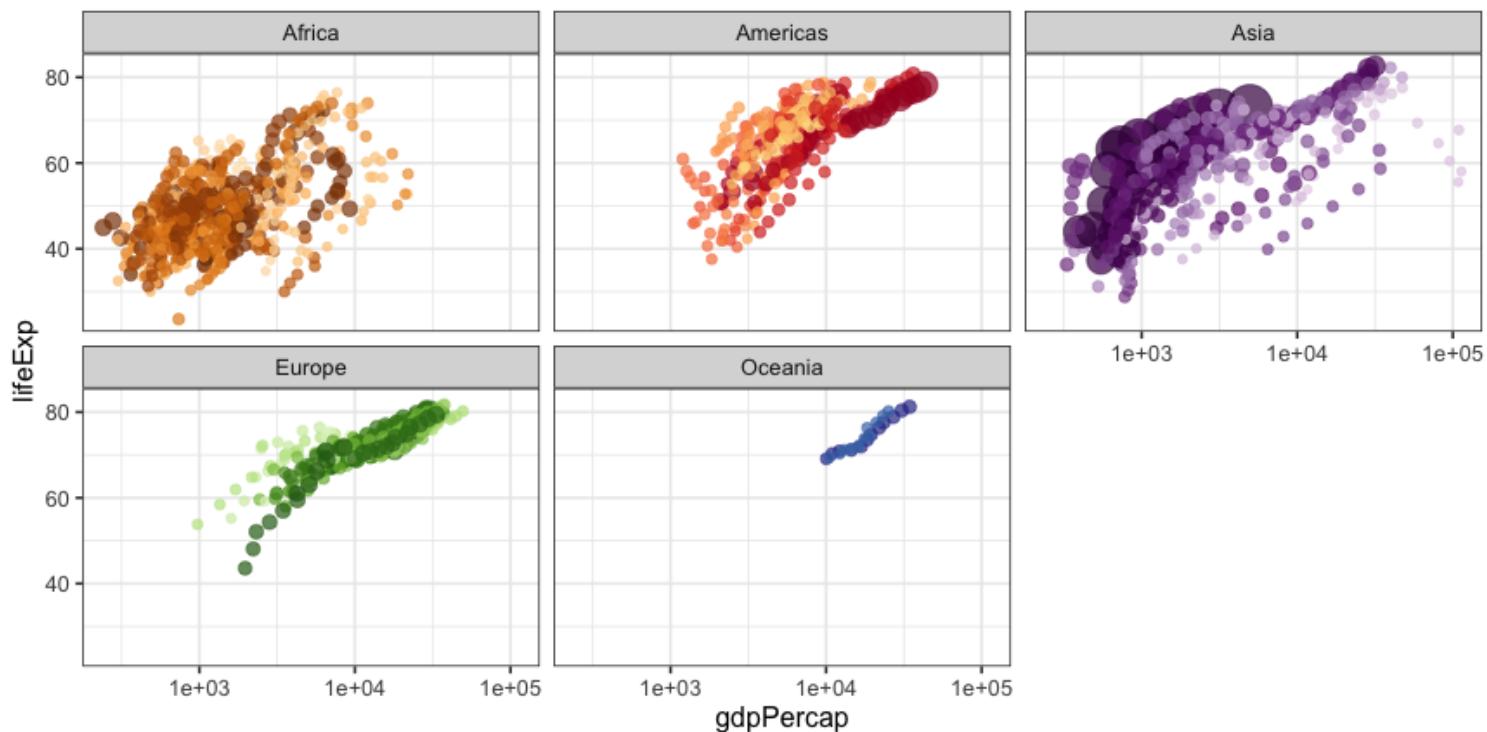
Data: gapminder

```
library(gapminder)  
gapminder
```

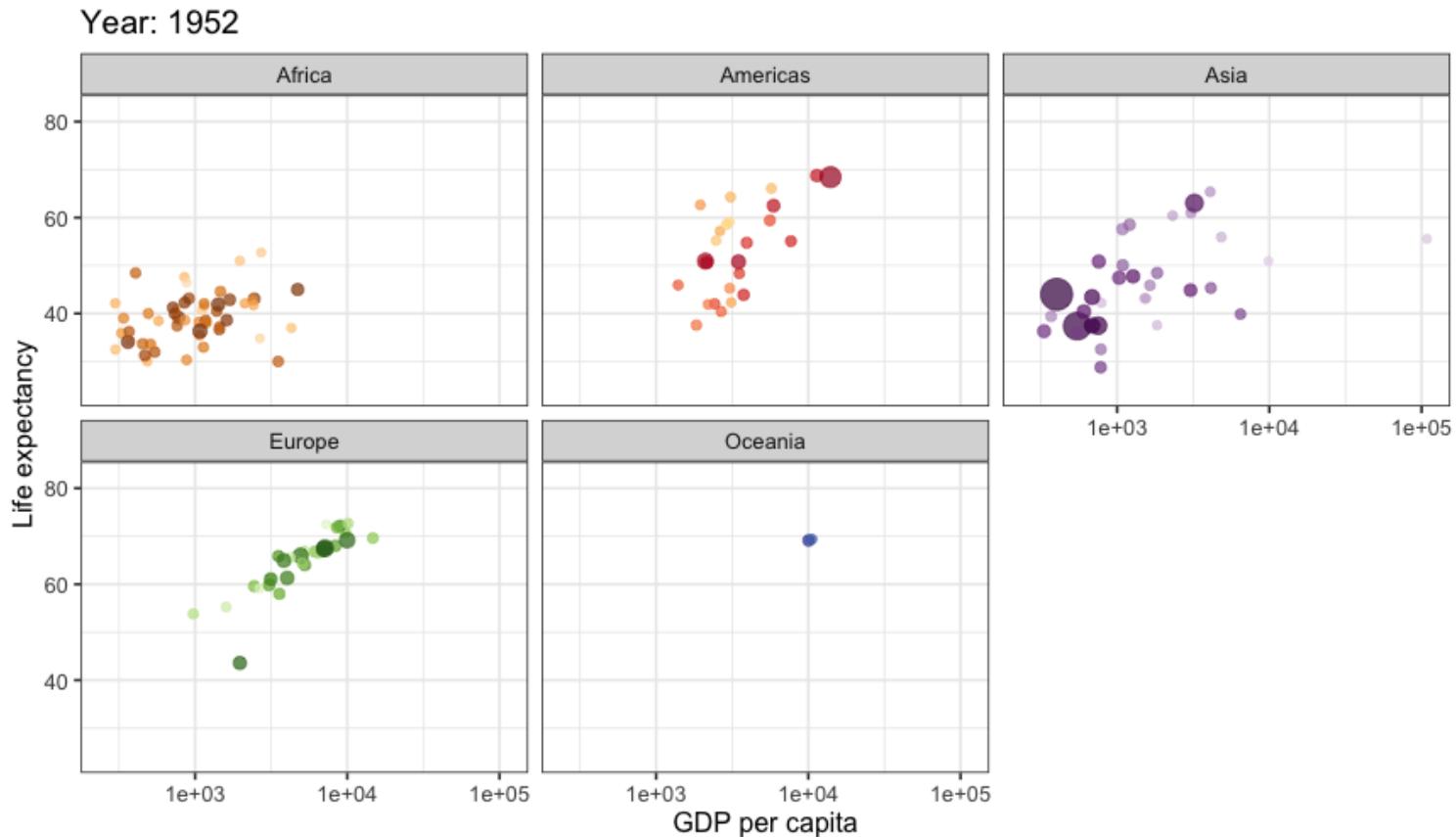
```
#> # A tibble: 1,704 x 6  
#>   country     continent   year lifeExp      pop gdpPercap  
#>   <fct>       <fct>     <int>    <dbl>    <int>      <dbl>  
#> 1 Afghanistan Asia      1952     28.8  8425333    779.  
#> 2 Afghanistan Asia      1957     30.3  9240934    821.  
#> 3 Afghanistan Asia      1962     32.0  10267083   853.  
#> 4 Afghanistan Asia      1967     34.0  11537966   836.  
#> 5 Afghanistan Asia      1972     36.1  13079460   740.  
#> 6 Afghanistan Asia      1977     38.4  14880372   786.  
#> 7 Afghanistan Asia      1982     39.9  12881816   978.  
#> 8 Afghanistan Asia      1987     40.8  13867957   852.  
#> 9 Afghanistan Asia      1992     41.7  16317921   649.  
#> 10 Afghanistan Asia     1997     41.8  22227415   635.  
#> # ... with 1,694 more rows
```

Nothing new

```
ggplot(gapminder, aes(x = gdpPercap, y = lifeExp, size = pop, colour = country)) +  
  geom_point(alpha = 0.7, show.legend = FALSE) +  
  scale_colour_manual(values = country_colors) +  
  scale_size(range = c(2, 12)) +  
  scale_x_log10() +  
  facet_wrap(~continent) +  
  theme_bw(base_size = 16)
```



Animate with `gganimate()`



What did we add?

Base plot

```
ggplot(gapminder, aes(x = gdpPercap, y = lifeExp, size = pop, colour = country)) +  
  geom_point(alpha = 0.7, show.legend = FALSE) +  
  scale_colour_manual(values = country_colors) +  
  scale_size(range = c(2, 12)) +  
  scale_x_log10() +  
  facet_wrap(~continent) +  
  theme_bw(base_size = 16)
```

Transform to animation

```
ggplot(gapminder, aes(x = gdpPercap, y = lifeExp, size = pop, colour = country)) +  
  geom_point(alpha = 0.7, show.legend = FALSE) +  
  scale_colour_manual(values = country_colors) +  
  scale_size(range = c(2, 12)) +  
  scale_x_log10() +  
  facet_wrap(~continent) +  
  theme_bw(base_size = 16) +  
  labs(title = 'Year: {frame_time}', x = 'GDP per capita', y = 'Life expectancy') +  
  transition_time(year) +  
  ease_aes('linear')
```

Package ganimate

- Core functions
 - `transition_*` () defines how the data should be spread out and how it relates to itself across time.
 - `view_*` () defines how the positional scales should change along the animation.
 - `shadow_*` () defines how data from other points in time should be presented in the given point in time.
 - `enter_*` () / `exit_*` () defines how new data should appear and how old data should disappear during the course of the animation.
 - `ease_aes` () defines how different aesthetics should be eased during transitions.
- Label variables
 - function dependent, use { } to access their values.
- See <https://ganimate.com>

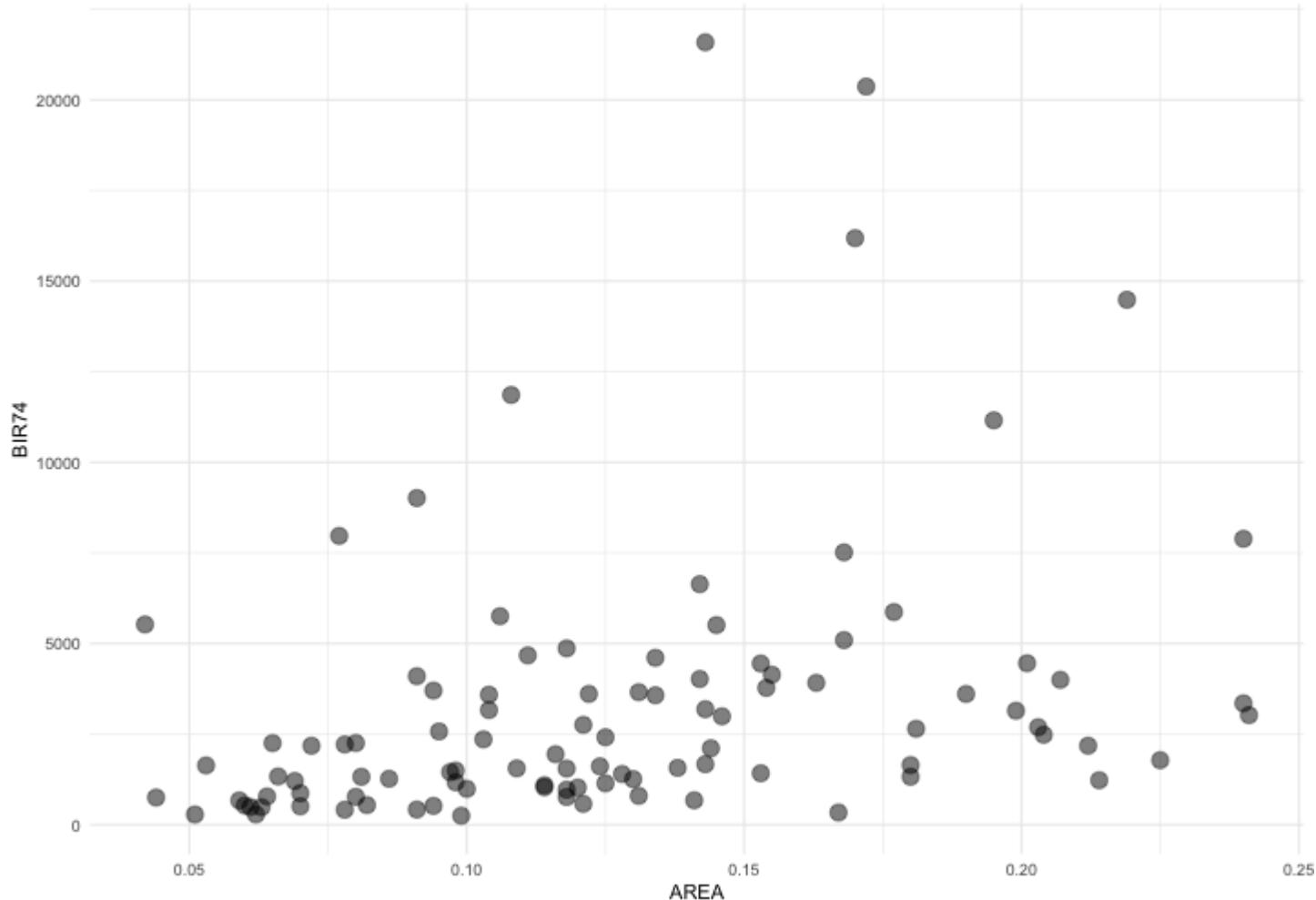
Interactive plots: ggiraph

Data: NC births and SID

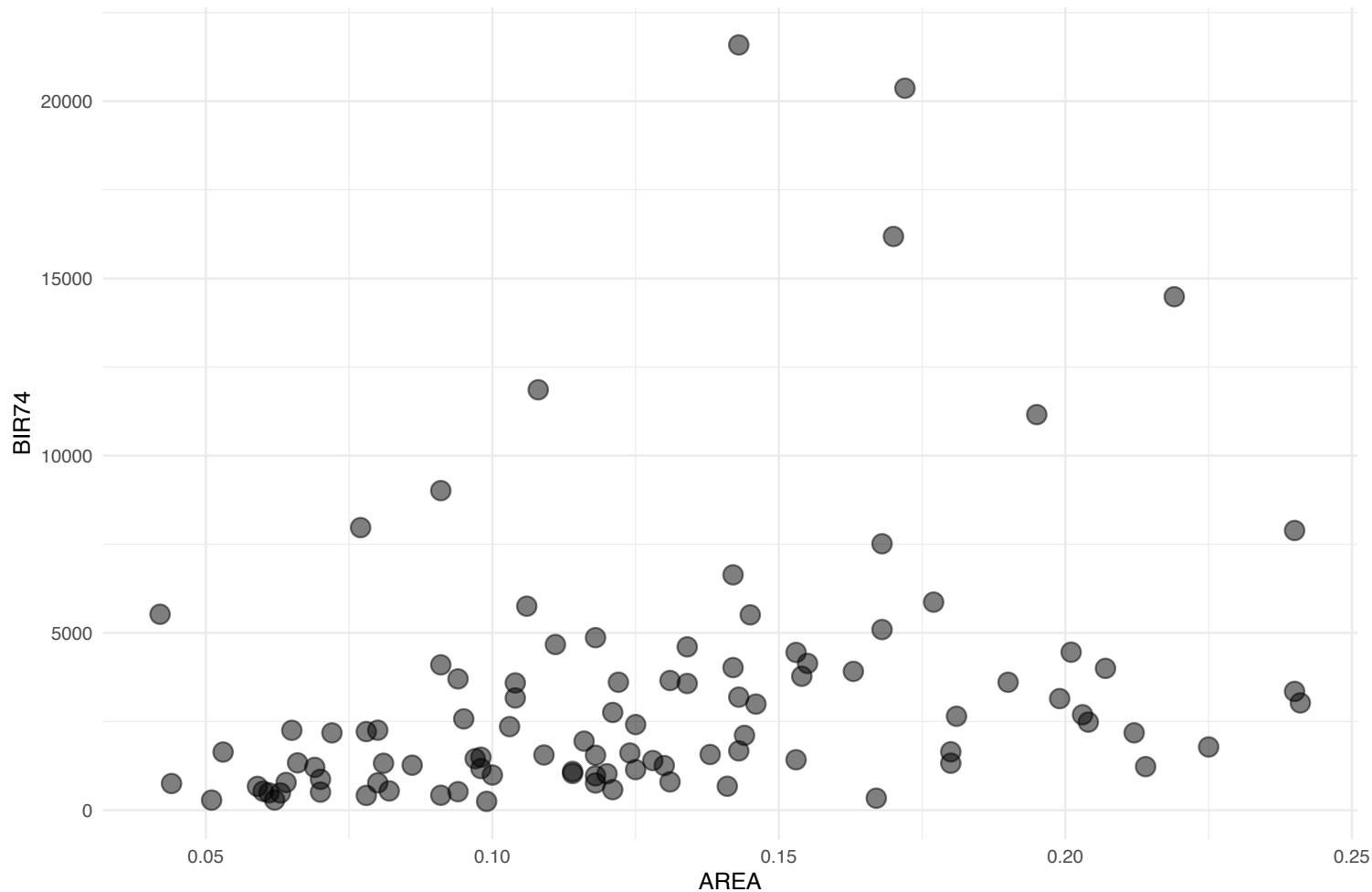
```
nc <- read_csv("http://www2.stat.duke.edu/~sms185/data/health/nc_birth_si  
nc
```

```
#> # A tibble: 100 x 4  
#>   NAME      AREA  BIR74  SID74  
#>   <chr>    <dbl> <dbl> <dbl>  
#> 1 Ashe     0.114  1091    1  
#> 2 Alleghany 0.061   487    0  
#> 3 Surry    0.143  3188    5  
#> 4 Currituck 0.07    508    1  
#> 5 Northampton 0.153  1421    9  
#> 6 Hertford  0.097  1452    7  
#> 7 Camden    0.062   286    0  
#> 8 Gates     0.091   420    0  
#> 9 Warren    0.118   968    4  
#> 10 Stokes   0.124  1612    1  
#> # ... with 90 more rows
```

Standard scatter plot



Which counties are these?



What changed?

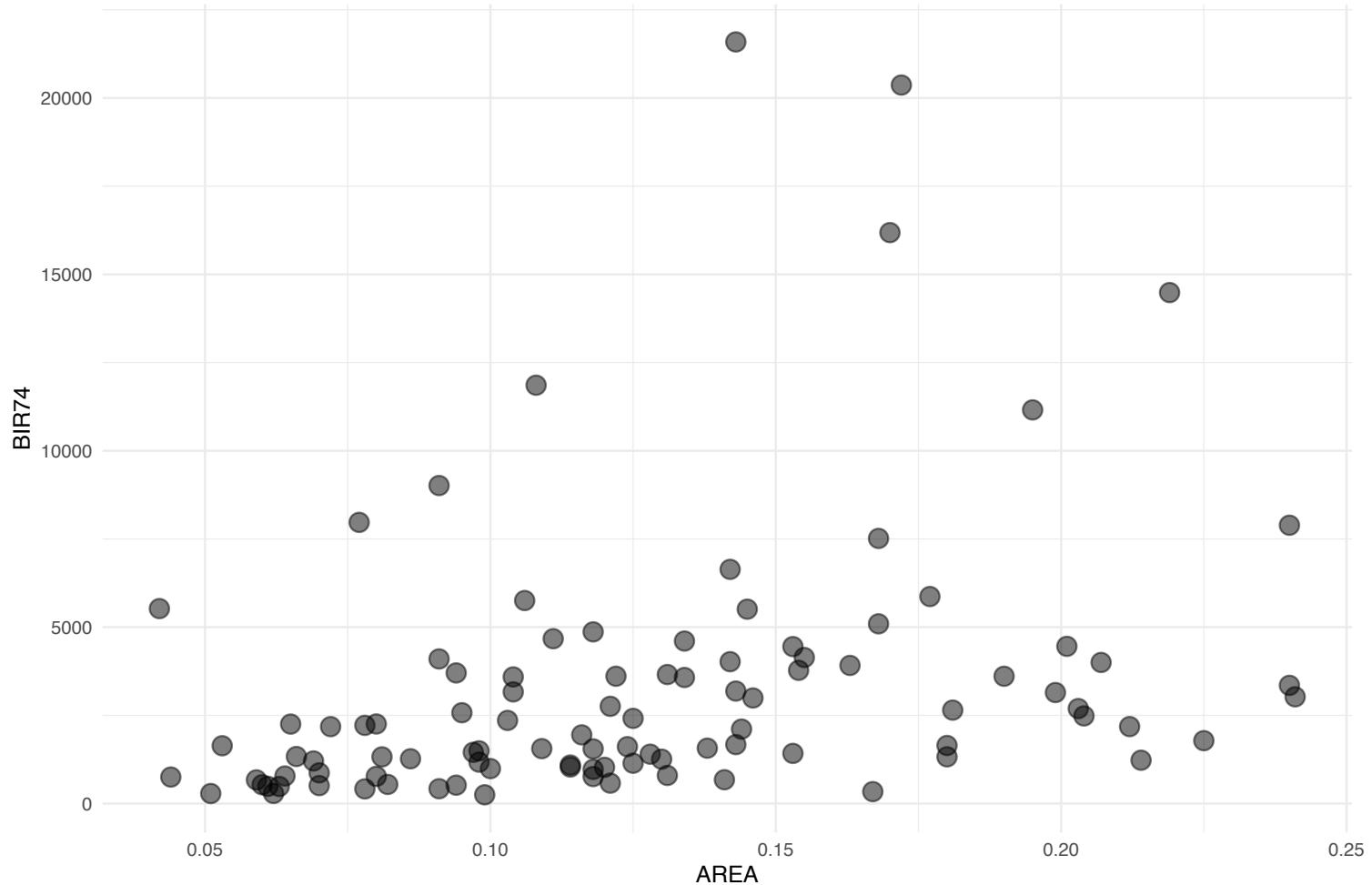
A scatter plot with `geom_point()`

```
ggplot(nc, mapping = aes(x = AREA, y = BIR74)) +  
  geom_point(size = 4, alpha = .5) +  
  theme_minimal()
```

A scatter plot with `geom_point_interactive()`

```
gg_name <- ggplot(nc, mapping = aes(x = AREA, y = BIR74)) +  
  geom_point_interactive(aes(tooltip = NAME), size = 4, alpha = .5) +  
  theme_minimal()  
  
girafe(code = {print(gg_name)}, height_svg = 6, width_svg = 9)
```

On hover functionality



What changed?

A scatter plot with `geom_point_interactive()`

```
gg_name <- ggplot(nc, mapping = aes(x = AREA, y = BIR74)) +  
  geom_point_interactive(aes(tooltip = NAME), size = 4, alpha = .5) +  
  theme_minimal()  
  
girafe(code = {print(gg_name)}, height_svg = 6, width_svg = 9)
```

On hover functionality with `data_id`

```
gg_hover <- ggplot(nc, mapping = aes(x = AREA, y = BIR74)) +  
  geom_point_interactive(aes(data_id = NAME, tooltip = NAME),  
                        size = 4, alpha = .5) +  
  theme_minimal()  
  
girafe(code = {print(gg_hover)}, height_svg = 6, width_svg = 9)
```

On click functionality

```
nc$wiki <- sprintf("window.open(\"%s%s\")",
                     "https://www.ncpedia.org/geography/", tolower(nc$NAME)
                     )

gg_name <- ggplot(nc, mapping = aes(x = AREA, y = BIR74)) +
  geom_point_interactive(aes(tooltip = NAME, onclick = wiki), size = 4,
                        theme_minimal())

girafe(code = {print(gg_name) })
```

Package ggiraph

- Add tooltips, animations, and JavaScript actions to ggplot graphics
- In general, instead of `geom_<plot_type>()` use `geom_<plot_type>_interactive()`
- Interactivity is added to ggplot geometries, legends and theme elements, via the following aesthetics:
 - `tooltip`: tooltips to be displayed when mouse is over elements.
 - `onclick`: JavaScript function to be executed when elements are clicked.
 - `data_id`: id to be associated with elements (used for hover and click actions)
- Function `girafe()` translates the graphic into an interactive web-based graphic
- See <https://github.com/davidgohel/ggiraph>

Exercise

Flint water data

```
flint <- read_csv("http://www2.stat.duke.edu/~sms185/data/health/flint.csv")
```

```
#> # A tibble: 271 x 6
#>   id    zip  ward draw1  draw2  draw3
#>   <dbl> <dbl> <dbl>  <dbl>  <dbl>  <dbl>
#> 1 1     48504     6  0.344  0.226  0.145
#> 2 2     48507     9  8.13   10.8   2.76
#> 3 4     48504     1  1.11   0.11   0.123
#> 4 5     48507     8  8.01   7.45   3.38
#> 5 6     48505     3  1.95   0.048  0.035
#> 6 7     48507     9  7.2    1.4    0.2
#> 7 8     48507     9  40.6   9.73   6.13
#> 8 9     48503     5  1.1    2.5    0.1
#> 9 12    48507     9  10.6   1.04   1.29
#> 10 13    48505     3  6.2    4.2    2.3
#> # ... with 261 more rows
```

Create a visualization of the data from object `flint`. Incorporate topics from today's lecture.

More visualization resources

- <http://r-statistics.co/Top50-Ggplot2-Visualizations-MasterList-R-Code.html>
- <https://plot.ly/ggplot2/>

References

- <https://yihui.name/en/2018/08/gifski-knitr/>
- <http://r-statistics.co/Top50-Ggplot2-Visualizations-MasterList-R-Code.html>
- <https://ggplot2.tidyverse.org/articles/extending-ggplot2.html>
- <https://github.com/davidgohel/ggiraph>
- <https://ganimate.com>
- <https://github.com/thomasp85/patchwork>
- <https://github.com/erocoar/ggpol>