

How to Create Data Visualizations Using Best Practices

R Breakout Session

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Data for Public Good Symposium

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Why do we visualize data?

Questions for You

- What do you use data visualizations for?
 - Data exploration/understanding
 - Figures for organizational reports
 - Figures for academic papers/grants
- Who is your typical audience when producing data visualizations?
 - Yourself
 - Members of your organization
 - Members of another organization
 - Members of the scientific community
- What is most important to you when thinking about data visualization?
 - Clarity
 - Reproducibility
 - Ease of creation

Why do we visualize data?

(and why exploratory data visualization is important)

The DatasauRus Dozen

Consider a dataset with (x,y) data coming from 12 groups:

Show

8

 entries

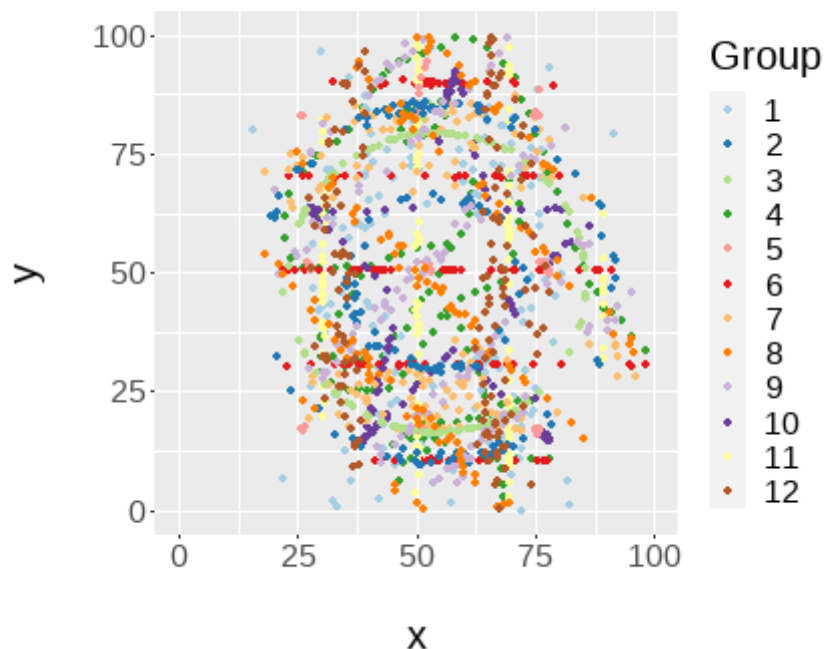
Search:

	Group	x	y
1	4	55.3846	97.1795
2	4	51.5385	96.0256
3	4	46.1538	94.4872
4	4	42.8205	91.4103
5	4	40.7692	88.3333
6	4	38.7179	84.8718
7	4	35.641	79.8718
8	4	33.0769	77.5641

The DatasauRus Dozen

In aggregate, the data show no discernible pattern:

Visualizing the Data



Summary Statistics

Show entries Search:

	Statistic	Value
1	Mean(x)	54.27
2	Mean(y)	47.83
3	SD(x)	16.71
4	SD(y)	26.85
5	Corr(x,y)	-0.07

Showing 1 to 5 of 5 entries

Previous

1

Next

The DatasauRus Dozen

Each group has summary statistics that are nearly identical:

Show

8

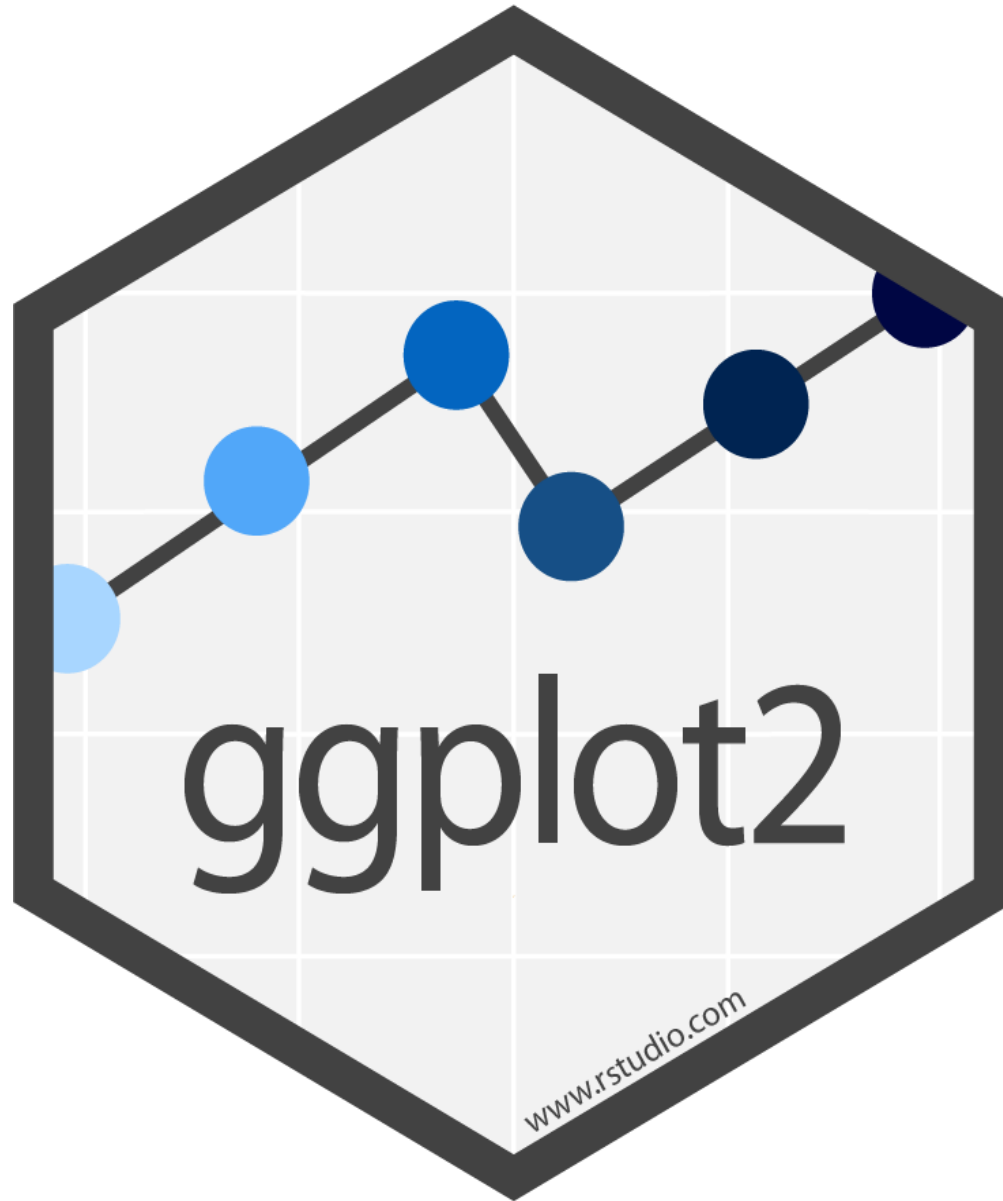
 entries

Search:

Group		Mean(x)	Mean(y)	SD(x)	SD(y)	Corr(x,y)
1	1	54.27	47.83	16.77	26.94	-0.06
2	2	54.27	47.83	16.77	26.94	-0.07
3	3	54.27	47.84	16.76	26.93	-0.07
4	4	54.26	47.83	16.77	26.94	-0.06
5	5	54.26	47.84	16.77	26.93	-0.06
6	6	54.26	47.83	16.77	26.94	-0.06
7	7	54.27	47.84	16.77	26.94	-0.07
8	8	54.27	47.84	16.77	26.94	-0.07

The DatasauRus Dozen

Yet, plotting reveals that each group is quite distinct:



ggplot2

- Though R has built-in functions for visualizing data, today's focus will be the `ggplot2` package
- `ggplot2` is based on the Grammar of Graphics by Leland Wilkinson (see: ggplot2.tidyverse.org)
- Grammar states that all data visualizations consist of three parts:
 1. Data
 2. Aesthetics
 3. Geometry
- `ggplot2` allows you to uniquely describe any plot as a combination of these three parts, via seven parameters:

```
ggplot(data = <DATA>) +  
  <GEOMETRY_FUNCTION>(  
    mapping = aes(<MAPPINGS>),  
    stat = <STAT>,  
    position = <POSITION>  
  ) +  
  <COORDINATE_FUNCTION>() +  
  <FACET_FUNCTION>()
```

1. Data

- `ggplot2` works best with **tidy data**:
 1. Each variable in the data set has its own column
 2. Each observation has its own row
 3. Each value has its own cell
- It is (usually) best practice to have a single data source (see: cfss.uchicago.edu/notes/tidy-data/)

country	year	cases	population
Afghanistan	1999	745	19987071
Afghanistan	2000	666	20095360
Brazil	1999	37737	172006362
Brazil	2000	80488	174004898
China	1999	212258	1272915272
China	2000	216766	1280425583

variables

country	year	cases	population
Afghanistan	1999	745	19987071
Afghanistan	2000	666	20095360
Brazil	1999	37737	172006362
Brazil	2000	80488	174004898
China	1999	212258	1272915272
China	2000	216766	1280425583

observations

country	year	cases	population
Afghanistan	99	745	19987071
Afghanistan	00	666	20095360
Brazil	99	37737	172006362
Brazil	00	80488	174004898
China	99	212258	1272915272
China	00	216766	1280425583

values

Untidy vs. Tidy Data

Untidy Data

Show entries Search:

	name	quiz1	quiz2	test1
1	Billy		D	C
2	Suzy	F		
3	Lionel	B	C	B
4	Jenny	A	A	B

Showing 1 to 4 of 4 entries

Tidy Data

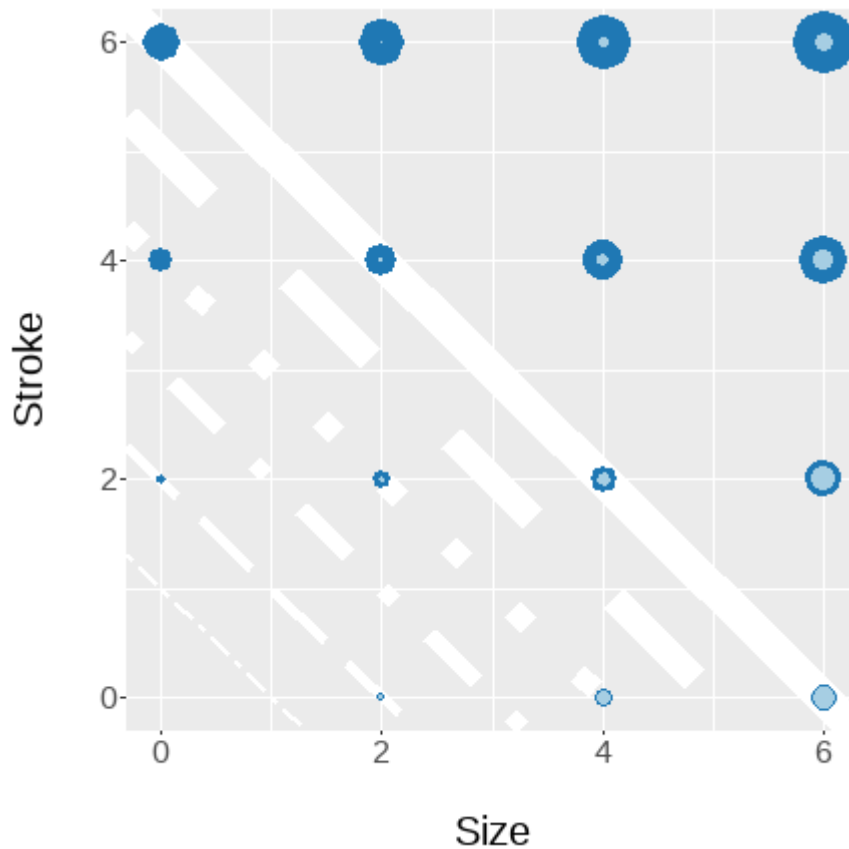
Show entries Search:

	Name	Assessment	Grade
1	Billy	Quiz 1	
2	Billy	Quiz 2	D
3	Billy	Test 1	C
4	Jenny	Quiz 1	A

Showing 1 to 4 of 12 entries

2. Aesthetics

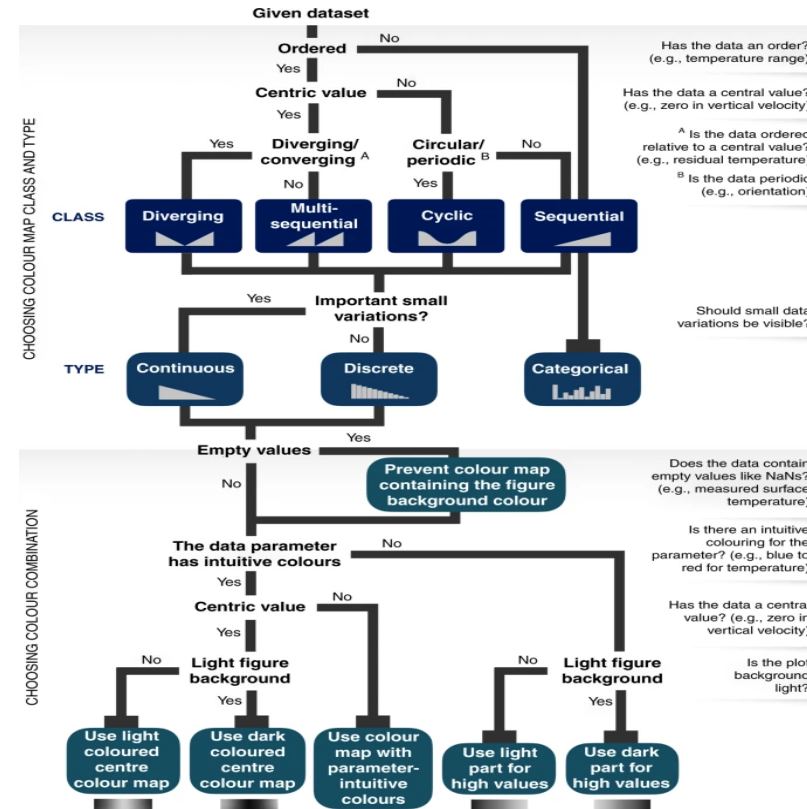
- Aesthetics map the columns in our data to standard elements of a plot
- These define the 'visual cues' we use to distinguish aspects of our data:
 1. Coordinate Positions (x- and y- axes)
 2. Color (i.e. 'outside' color)
 3. Fill (i.e. 'inside' color)
 4. Point Shapes
 5. Line Types
 6. Element Sizes



A Note on Color

Use color **intentionally** and **thoughtfully**:

- Meet basic accessibility guidelines for color
 - Color blindness
 - Intended medium (color vs. black and white)
- Do not reinforce gender or racial stereotypes
 - Blue for boys and pink for girls
 - Colors associated with skin tones
- Consider data class, type, and display
 - Continuous, Ordinal, Nominal
 - Increasing/Decreasing, Centric Valued



[nature.com/articles/s41467-020-19160-7](https://www.nature.com/articles/s41467-020-19160-7)

Color Palettes

- R accommodates RGB, CMYK, or Hex color and has preset palettes to choose from (e.g. `ggsci` package):
 - Academic Journals (e.g. JAMA), Institutional (e.g. U Chicago), Just for Fun (e.g. Star Trek)
- Default colors for most data visualization software are difficult to see if color blind (`ggplot2` included)
- Thankfully, there are color blind friendly options (what I have been using thus far)

Viridis

```
library(scales)
# viridis also prints well in greyscale!
show_col(viridis_pal()(5), ncol = 5)
```

RColorBrewer

```
library(RColorBrewer)
# display.brewer.all(colorblindFriendly = T)
show_col(brewer.pal(5, "Set2"), ncol = 5)
```

3. Layers and Geometries

- A layer combines data, aesthetic mapping, a geom (geometric object), a stat (statistical transformation), and a position adjustment
- You will create layers using a `geom_XXXX()` function, overriding the default position and stat if needed
- Choice of geometry is specific to your data:
 - **Reference Lines:** `geom_abline()`, `geom_hline()`, `geom_vline()`
 - **Bar Charts:** `geom_bar()`, `geom_col()`, `stat_count()`
 - **Densities:** `geom_density()`, `stat_density()`
 - **Histograms:** `geom_freqpoly()`, `geom_histogram()`, `stat_bin()`
 - **... and many, many more!**
- See: ggplot2.tidyverse.org/reference/

Putting it All Together

Installing ggplot2

- Even though the package is sometimes just referred to as "ggplot," the package name is `ggplot2`
- `ggplot2` is a package included in the **tidyverse**: tidyverse.org
- To install the tidyverse packages (if you have not already), you will need to run:

```
install.packages("tidyverse") ## Run this only once
```

- To load the tidyverse packages, run:

```
library(tidyverse) ## Run this each session
```

- Be sure to load the tidyverse packages at the start of each session

What does ggplot() do?

- The `ggplot()` function returns a "gg/ggplot" object
- In the **Grammar of Graphics** workflow, ggplot objects are 'built up':
 1. Define a new plot (ggplot object) associated with your data
 2. Use (+) to add additional instructions (layers) to the object to build your plot
 3. Within layers, add aesthetics, geometries, statistical transformations, and/or position adjustments

```
ggplot(data = <DATA>) +  
  <GEOM_FUNCTION>(      # 1. Define a new plot  
    mapping = aes(<MAPPINGS>), # 2. Add layers  
    stat = <STAT>,           # 3. Within layers:  
    position = <POSITION>    #   Add aesthetics  
                             #   Add statistical transformations  
                             #   Add position adjustments  
  )
```

- `ggplot()` can also be passed any default aesthetic mappings you would like all layers to use

```
ggplot(data = <DATA>, mapping = aes(<MAPPINGS>)) +  
  <GEOM_FUNCTION>()
```

Mapping Values with aes()

- `aes()` is the function that defines a mapping from data to geometry aesthetics
- Signature: `aes(x, y, ...)`
 - The first two parameters default to x and y values
 - All other parameters should be named: `aes(x, y, <aesthetic> = column_name)`
- `aes()` expects parameters to be "symbols" or "expressions"
 - Do not use quotes or variables (see `aes_q` or `aes_string`)
 - May contain functions (e.g. `aes(x=a+b, y=sqrt(z))`)
 - Never use "\$" inside an `aes()`

Mapping Values with aes()

- By default, layer `aes()` values are inherited from `ggplot()`
- Disable inheritance with `<GEOM_FUNCTION>(..., inherit.aes = FALSE)`
- You may also add, override, or remove `aes()` values:

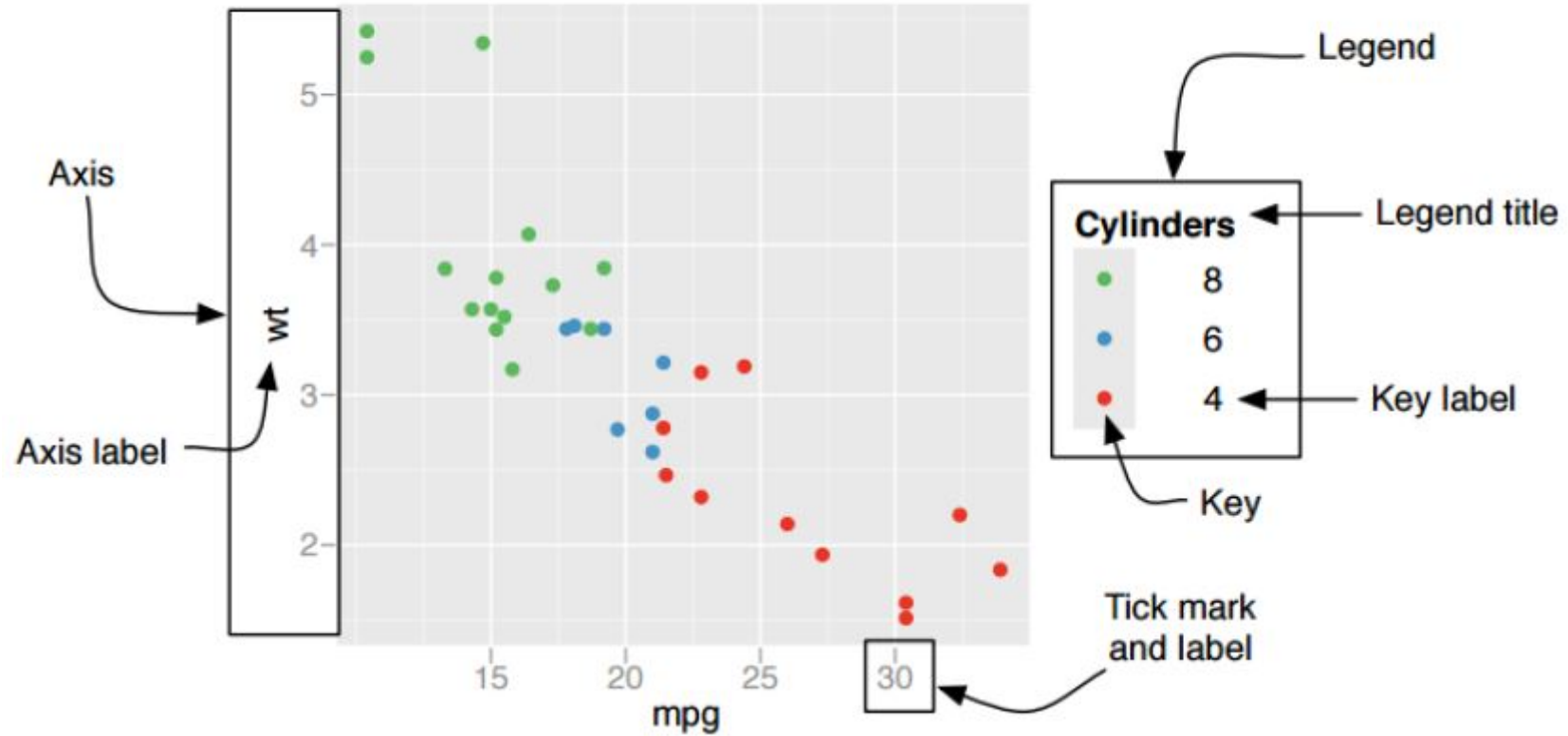
Show entries Search:

	ggplot()	geom_XXXX()	Resulting Aesthetics	Operation
1	aes(x, y)	aes(color = Group)	aes(x, y, color = Group)	Add
2	aes(x, y)	aes(y = y2)	aes(x, y2)	Override
3	aes(x, y)	aes(y = NULL)	aes(x)	Remove

Showing 1 to 3 of 3 entries Previous 1 Next

There are many ways to get the same plot!

Plot parts



Labeling

- You can label your axes, titles, and more with `labs()`:

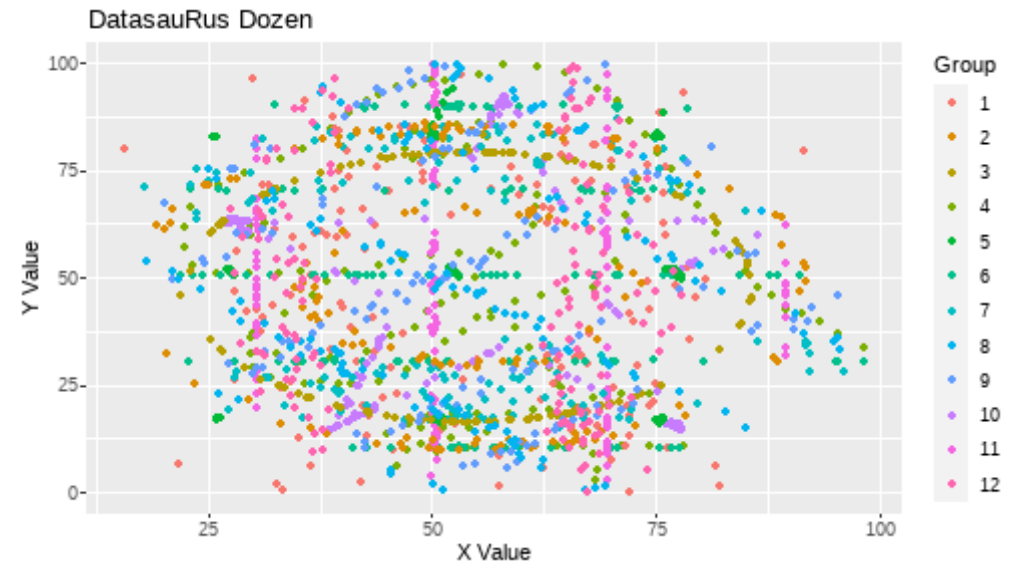
```
+ labs(x="X Name",  
      y="Y Name",  
      title="Plot Title"  
      subtitle="Plot Subtitle")
```

- You can also include mathematical expressions (see `?plotmath`)

```
+ labs(title=expression(y==alpha+beta*x))
```

- Setting values to `""` shows no label, setting values to `NULL` removes space for label as well

```
ggplot(datasaurus_dozen2,  
       aes(x, y, color=Group)) +  
  geom_point() +  
  labs(x="X Value",  
       y="Y Value",  
       title="DatasauRus Dozen")
```



Formatting Tick Labels

- Format 0-1 as percents: + `scale_y_continuous(labels=scales::percent_format())`
- Format as dollar amounts: + `scale_y_continuous(labels=scales::dollar_format("$"))`
- Format in thousands: + `scale_fill_continuous(labels=scales::unit_format("k", 1e-3))`
- You can pass any function as the `labels=` parameter

Formatting Tick Marks

- You can set where your tick marks fall with `scale_x_continuous()`
 - `breaks`= vector of values where to draw major tick marks
 - `labels`= vector of values with what to draw at those tick marks
 - `minor_breaks`= vector of values where to draw minor (unlabeled) tick marks
 - `trans`= optional transformation to apply to axis
 - `expand`= how far to extend axis past observed data
 - `limits`= lower and upper bound for tick marks
- For date-time axes: `date_minor_breaks`= with units like "1 month" or "2 years"

Themes

- The overall "look" of a plot is set by the theme
- Just call one of the theme functions to see all the values you can customize
- The `ggthemes` package has many themes to try out
- You can also create your own theme objects

```
p <- ggplot(mpg, aes(cty, hwy, color=factor(cyl))) +  
  geom_jitter() +  
  geom_abline(color="grey50", size=2) +  
  ggtitle("My Plot!")
```

```
p + theme_grey() # Default Theme  
p + theme_bw()  
p + theme_linedraw()  
p + theme_light()  
p + theme_dark()  
p + theme_minimal()
```

Customizing a Theme

- You can customize parts of themes

```
p + theme(plot.title = element_text(color="red", margin=margin(t=20, b=20)))  
p + theme(panel.background = element_rect(fill="linen"))
```

- Increase font size for presentation slides

```
p + theme_grey(base_size=18)
```

- Set default theme for session

```
theme_set(theme(...))
```

- The ggplot2 book or the "theme" help page on the ggplot2 website has more info

Examples!



SurvivoR Data

- **SurvivoR** is a collection of data sets in R detailing events across all 40 seasons and 596 episodes of the U.S. television show, Survivor
- Data include castaway information, vote history, immunity and reward challenge winners, and jury votes
- gradientdescending.com/survivor-data-from-the-tv-series-in-r/
- github.com/doehm/survivoR

```
devtools::install_github("doehm/survivoR") # Data are already tidy!
```

Season Summaries

```
library(survivoR)
season_summary
```

Show

3

 entries

Search:

	season_name	season	location	country	tribe_setup	full_name	winner
1	Survivor: Borneo	1	Pulau Tiga, Sabah, Malaysia	Malaysia	Two tribes of eight new players	Richard Hatch	Richard
2	Survivor: The Australian Outback	2	Herbert River at Goshen Station, Queensland, Australia	Australia	Two tribes of eight new players	Tina Wesson	Tina
3	Survivor: Africa	3	Shaba National Reserve, Kenya	Kenya	Two tribes of eight new players	Ethan Zohn	Ethan

Viewership Over Time

```
p1_dat <- season_summary %>%
  select(season, viewers_premier, viewers_finale, viewers_reunion, viewers_mean) %>%
  pivot_longer(cols = -season, names_to = "episode", values_to = "viewers") %>%
  mutate( episode = str_replace(episode, "viewers_", ""))

DT::datatable(p1_dat, fillContainer = F, options = list(pageLength = 3))
```

Show

3

 entries

Search:

	season	episode	viewers
1	1	premier	15.51
2	1	finale	51.69
3	1	reunion	36.7

Showing 1 to 3 of 160 entries

Previous

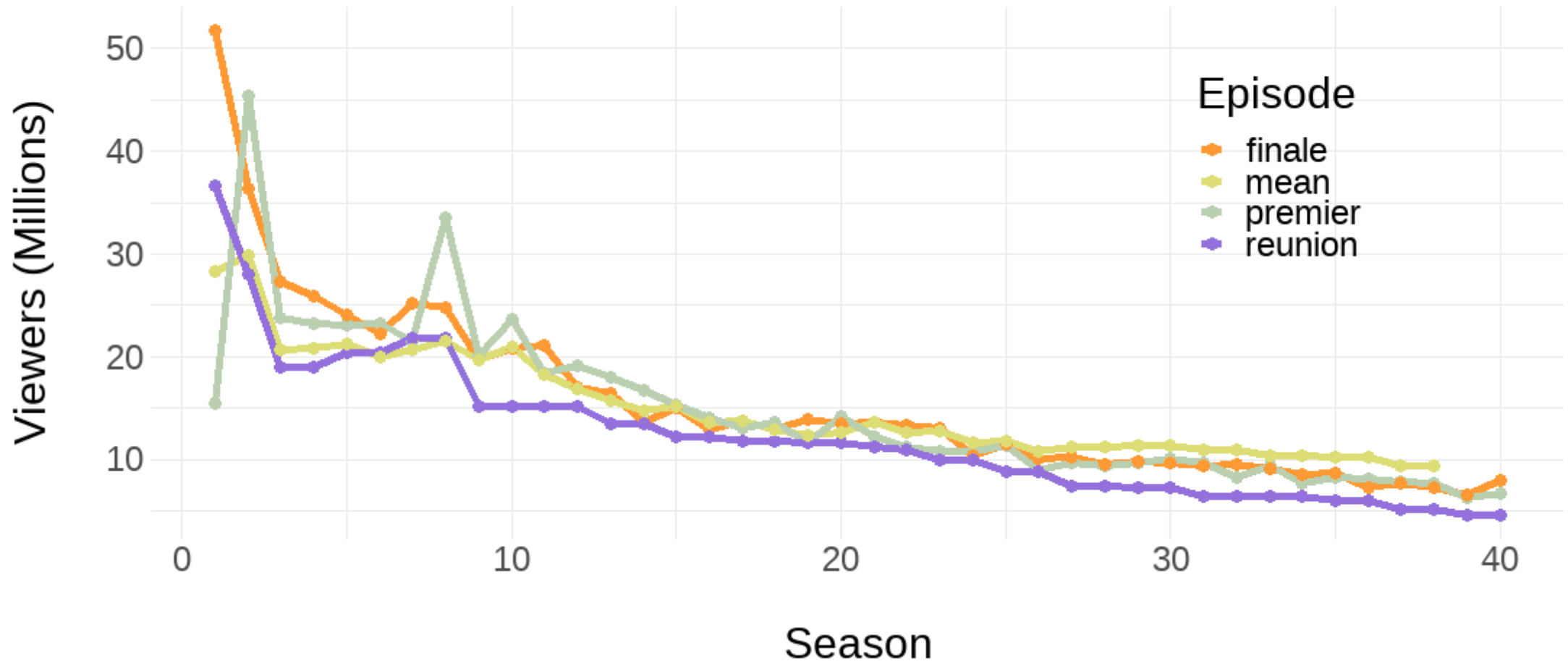
1

2
3
4
5
...
54
Next

Viewership Over Time

```
ggplot(p1_dat, aes(x = season, y = viewers, colour = episode)) +  
  geom_line(size = 2) + # Note 'size' is not in aes()  
  geom_point(size = 4) + # geom_point after geom_line overlays points on lines  
  theme_minimal() + # Many theme options, this keeps minor axes obly  
  scale_colour_survivor(16) + # Custom palette with colors from the show  
  labs(  
    title = "Survivor viewers over the 40 seasons",  
    x = "\nSeason", # '\n' is shorthand for 'newline'  
    y = "Viewers (Millions)\n", # '\n' adds space between axis title and ticks  
    colour = "Episode"  
  ) +  
  theme(  
    text = element_text(size = 30), # Changes the size for all text elements  
    legend.position = c(0.8, 0.7) # Puts the legend in the top right corner  
  )
```

Survivor viewers over the 40 seasons



Votes Received by Each Finalist

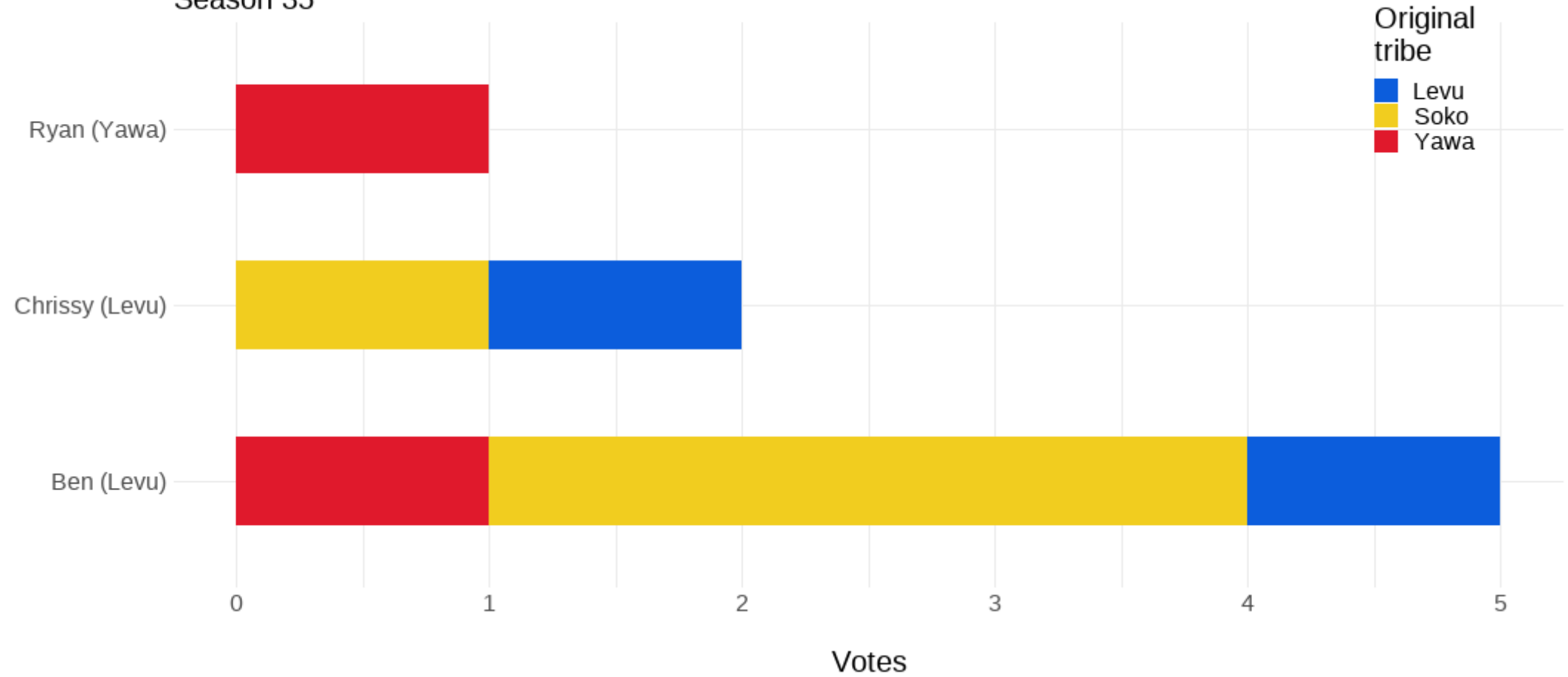
```
library(glue)

labels <- castaways %>%
  filter(season == 35, str_detect(result, "Sole|unner")) %>%
  mutate(label = glue("{castaway} ({original_tribe}")) %>%
  select(label, castaway)

jury_votes %>%
  filter(season == 35) %>%
  left_join(castaways %>% filter(season == 35) %>% select(castaway, original_tribe), by = "castaway")
  group_by(finalist, original_tribe) %>%
  summarise(votes = sum(vote)) %>%
  left_join(labels, by = c("finalist" = "castaway")) %>%
  {ggplot(., aes(x = label, y = votes, fill = original_tribe)) +
    geom_bar(stat = "identity", width = 0.5) +
    scale_fill_survivor(35, tribe = .$original_tribe) +
    coord_flip() +
    theme_minimal() +
    labs(x = "", y = "\nVotes", fill = "Original\ntribe",
         title = "Votes received by each finalist", subtitle = "Season 35") +
    theme(legend.position = c(0.9, 0.9),
          text = element_text(size = 20))}
```

Votes received by each finalist

Season 35

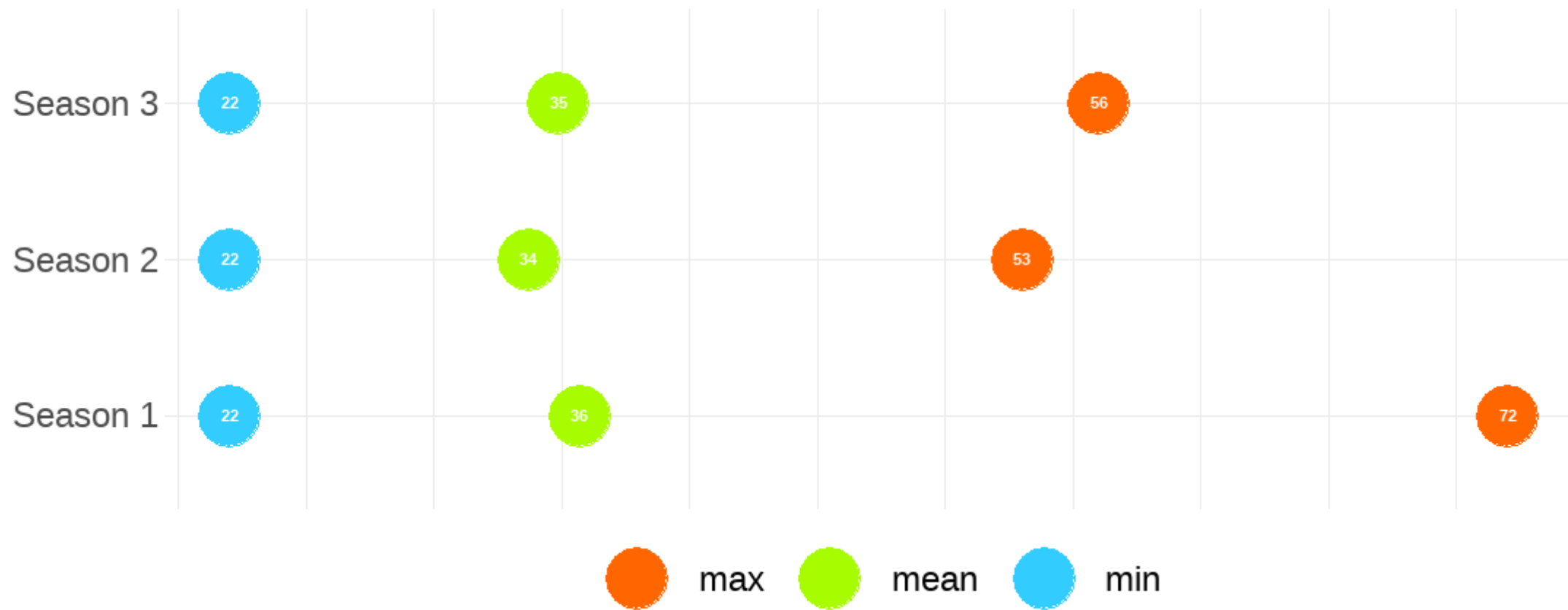


Age Distribution by Season

```
cols <- RColorBrewer::brewer.pal(3, "Blues")

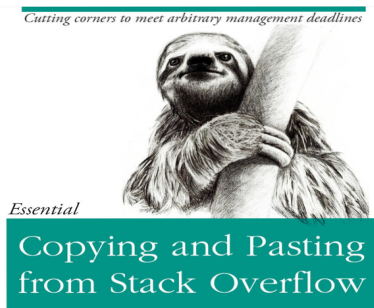
castaways %>%
  filter(season %in% 1:3) %>%
  group_by(season) %>%
  summarise(min_age = min( age, na.rm = T),
            mean_age = mean(age, na.rm = T),
            max_age = max( age, na.rm = T)) %>%
  pivot_longer(min_age:max_age) %>%
  mutate(name = name %>% str_replace("_age", "")) %>%
  ggplot(aes(x = value, y = paste("Season", season), color = name)) +
    theme_minimal() +
    geom_point(size = 20) +
    geom_text(aes(label = round(value)), colour = "white", fontface = "bold") +
    scale_colour_survivor(2) +
    labs(title = "Castaway Age Distribution",
         subtitle = "First Three Seasons",
         y = NULL,
         colour = NULL) +
    theme(text = element_text(size = 30),
          axis.text.x = element_blank(),
          axis.title.x = element_blank(),
```

Castaway Age Distribution First Three Seasons

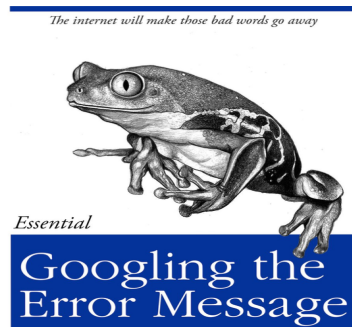


Getting Help

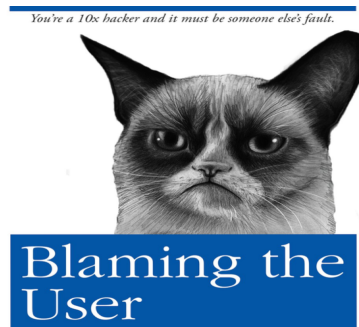
- Use R's built-in help documentation (e.g. `> ?ggplot`)
- Go to the `ggplot2` website: ggplot2.tidyverse.org/reference/
- Read the creator's (Hadley Wickham) book: *ggplot2: Elegant Graphics for Data Analysis*, second edition
- Google (my preferred method)



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arstechnica.com/gaming/2017/10/tim-oreilly-on-why-the-future-probably-wont-be-all-that-be-terrible/

References

These Slides

- Adapted from "ggplot2," a presentation by Matthew Flickinger, Ph.D. on Sept 12, 2018
- Slide were produced in R using the `xaringan` package: github.com/yihui/xaringan

The DatasauRus Dozen

- <https://cran.r-project.org/web/packages/datasauRus/vignettes/Datasaurus.html>
- <https://www.autodesk.com/research/publications/same-stats-different-graphs>

SurvivoR Data

- gradientdescending.com/survivor-data-from-the-tv-series-in-r/
- github.com/doehm/survivoR

Data Visualizations

- stephanieevergreen.com

Questions?

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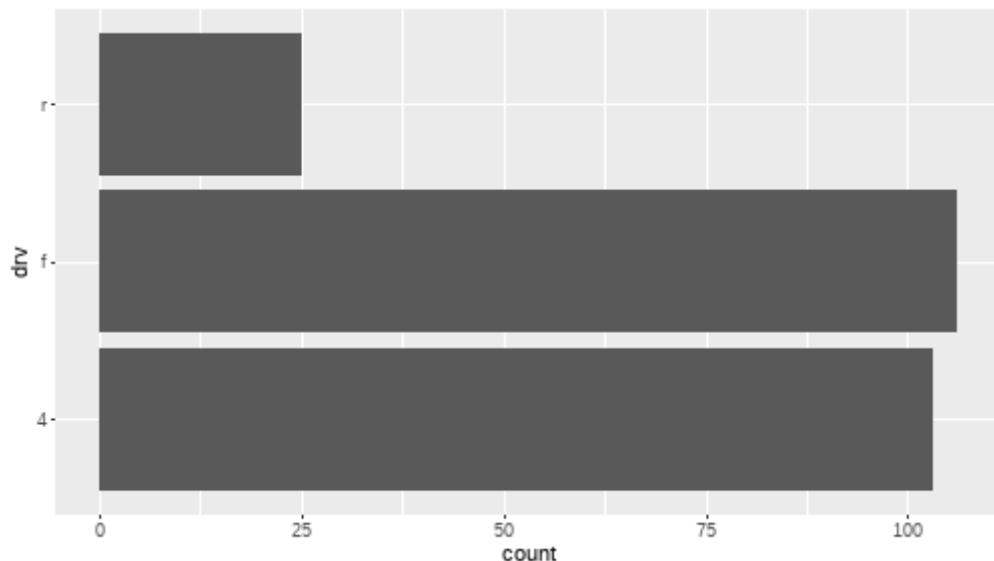
Potpourri

(time permitting)

Coordinate Transformations

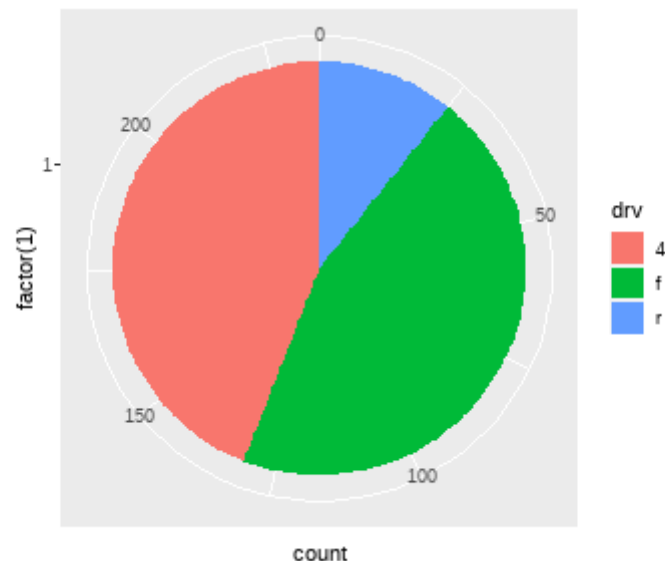
`coord_flip()`: Swap x & y axes

```
ggplot(mpg, aes(x=drv)) +  
  geom_bar() +  
  coord_flip()
```



`coord_polar()`: Make "pie" charts

```
ggplot(mpg) +  
  geom_bar(aes(factor(1), fill=drv), width=1) +  
  coord_polar(theta="y")
```

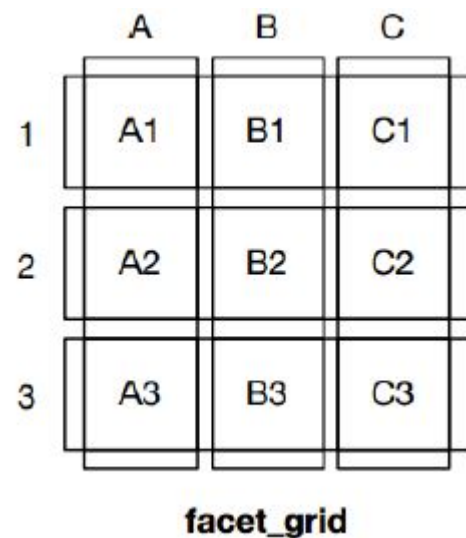


Coordinate Transformations

- `coord_cartesian()`: limit the plotting window
 - `xlim` = range of x values `c(lower, upper)`
 - `ylim` = range of y values `c(lower, upper)`
 - Differs from changing limits on scales which will subset data
- `coord_fixed()`: fix the distance ratio for x and y

Faceting

- Divide plot into subgroups and draw layers for each subset of data based on faceting variables
- Two primary options:
 - `facet_grid()`: Variables for rows/columns
 - `facet_wrap()`: No panel structure
- Each facet gets all layers
- Each layer's data split on same variables



facet_wrap()

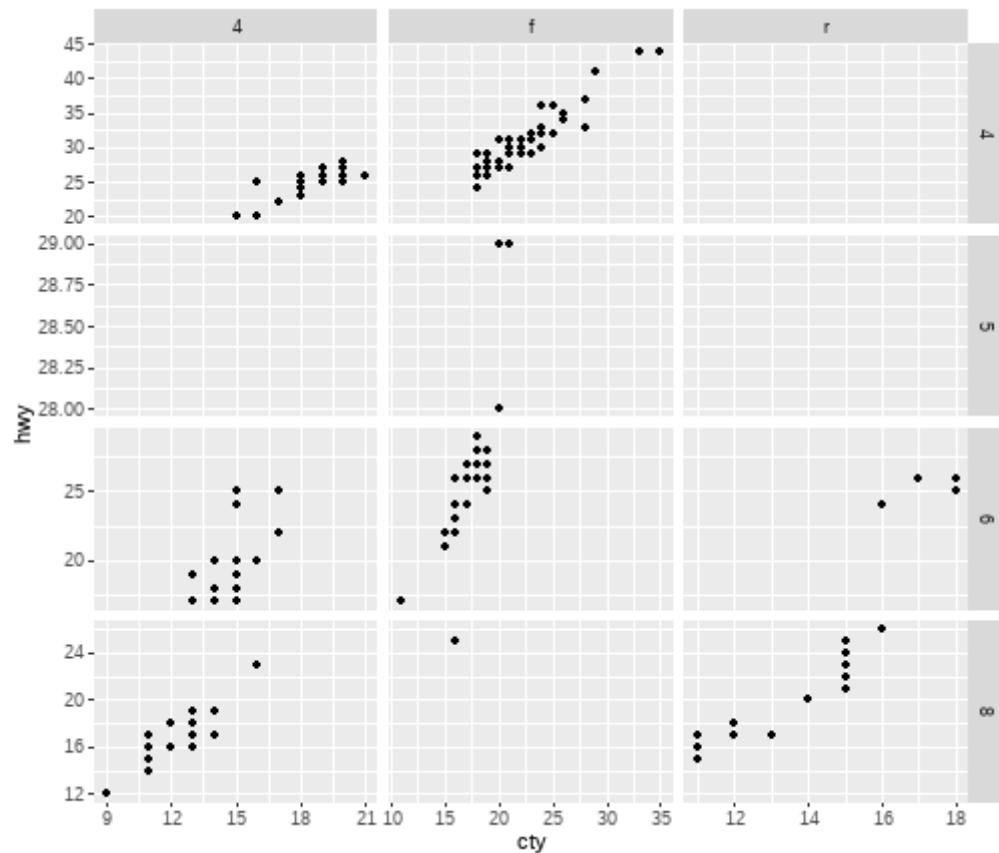
- Requires a right hand formula: `~ var1 + var2`
- Each combination of columns gets its own panel
- `scales=` options:
 - "fixed": scales same in all
 - "free_x": x range can vary
 - "free_y": y range can vary
 - "free" domain and range can change for each panel
- `ncol` = number of columns

```
library(gapminder)
ggplot(gapminder, aes(x=lifeExp)) +
  geom_density(fill="grey40") +
  facet_wrap(~continent)
```

facet_grid()

- Requires two arguments:
 - rows = vars(var1)
 - cols = vars(var2)
- Shared axis across panels
- Results in "rectangular" output

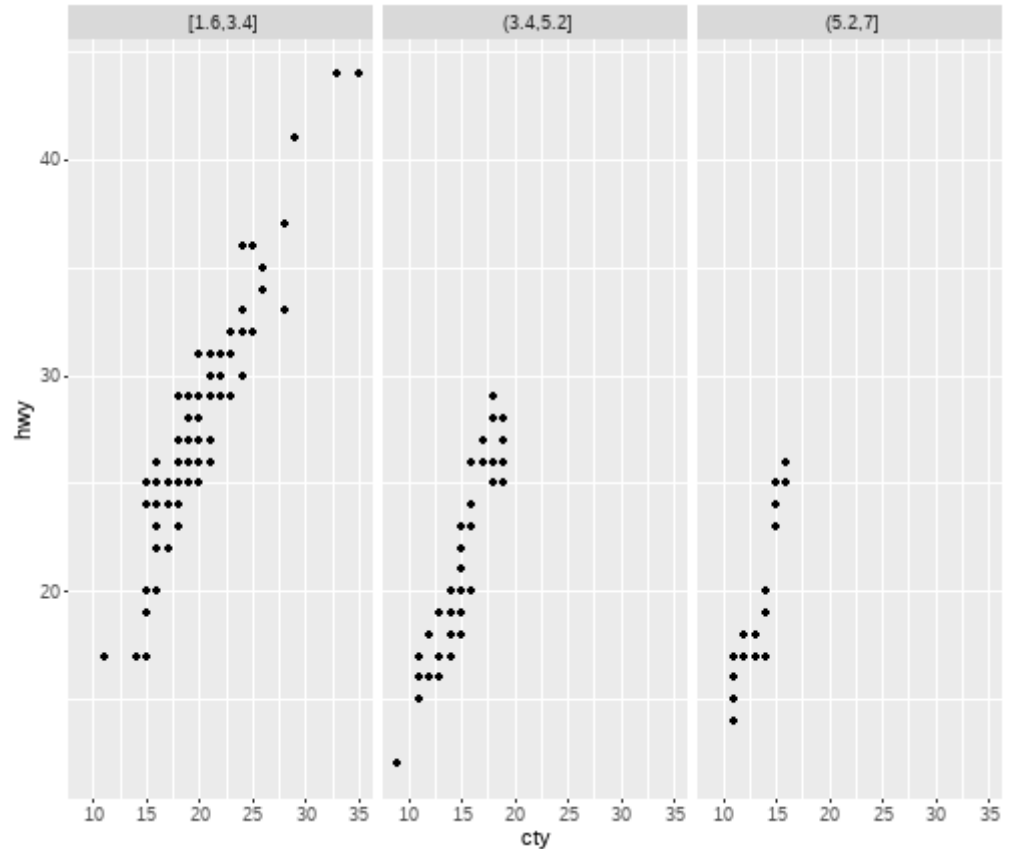
```
ggplot(mpg, aes(cty, hwy)) +  
  geom_point() +  
  facet_grid(rows=vars(cyl),  
             cols=vars(drv),  
             scales="free"  
  )
```



Faceting with Continuous Variables

- Faceting works best with categorical variables
- To use with categorical variables, we cut the values into groups
- Useful helper functions:
 - **Bins of same length:** `cut_interval()`
 - **Bins of a certain width:** `cut_width()`
 - **Bins with roughly equal counts:** `cut_number()`
- For example:

```
mpg %>%  
  mutate(d = cut_interval(displ, 3)) %>%  
  ggplot(aes(cty, hwy)) +  
  geom_point() +  
  facet_wrap (~d)
```



Legend

- Only mapped `aes()` values appear in the legend
- You can disable the legend for a layer with `show.legend=FALSE`
- You can also control where the legend goes:
 - "Outside"
 - `+ theme(legend.position="right")`
 - `+ theme(legend.position="bottom")`
 - "Inside"
 - `+ theme(legend.position=c(0, 1))`
 - (0,0) is lower left, (1,1) is top right, can be any number in between
 - No Legend: `+ theme(legend.position="none")`

Legend Guides

- The `guides()` function can control how keys are drawn in the legend
- Layout key values:

```
# Create more than 1 column
+ guides(fill = guide_legend(ncol=2))

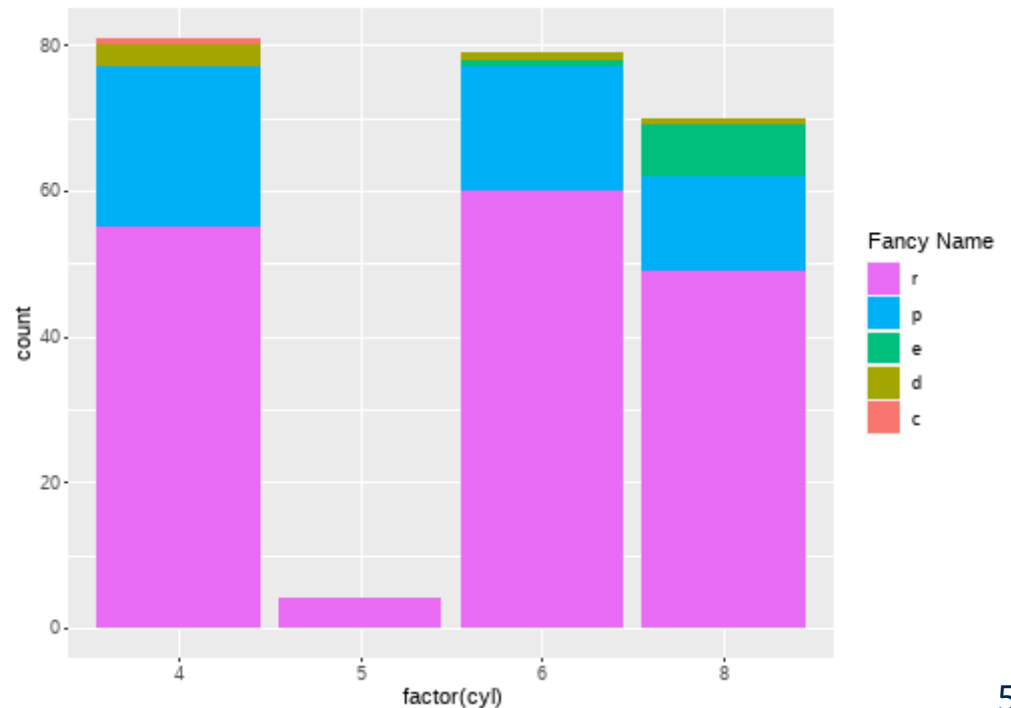
# Change row direction
+ guides(fill = guide_legend(ncol=2, byrow=T))

# Reverse order:
+ guides(fill = guide_legend(reverse=T))
```

- Remove guide: `+ guides(color="none")`
- Set title:

```
+ guides(color=guide_legend("Name"))
+ labs(color="Name")
```

```
ggplot(mpg, aes(factor(cyl), fill=fl)) +
  geom_bar() +
  guides(fill=guide_legend(reverse=T)) +
  labs(fill="Fancy Name")
```



Scales

- Scales describe how raw data values should be converted to aesthetic values
- Default scales are determined by the class of the variables in your data
- Each aesthetic (e.g. color, fill, size, shape) can have at most one scale
- Scales can have guides:
 - Axes for positions
 - Legends for everything else
- Color is mapped differently depending if year is a numeric or character vector
- Default color scales
 - **Numeric:** `scale_color_continuous()`
 - **Factor:** `scale_color_discrete()`

Scaling Based on Data Type

```
p <- gapminder %>%  
  filter(country=="United States") %>%  
  ggplot(aes(gdpPercap, lifeExp))
```

- Compare the following output based on whether `year` is continuous or a factor

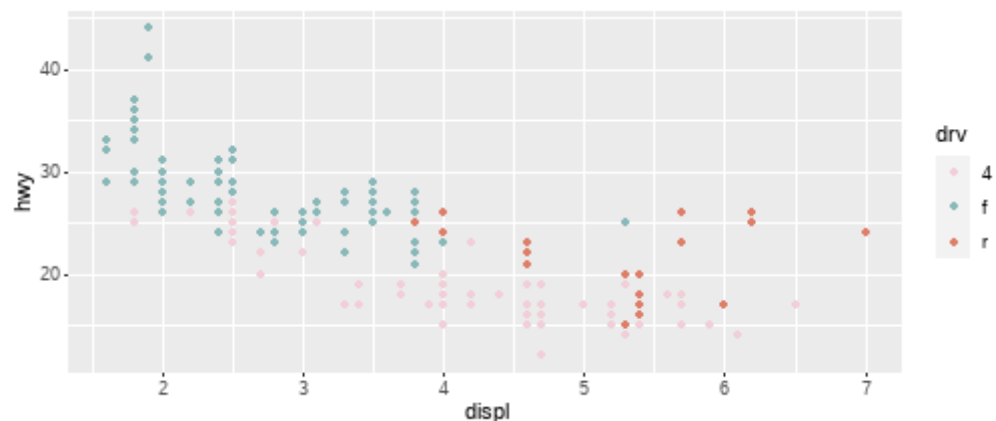
```
p + geom_point(aes(color=year))
```

```
p + geom_point(aes(color=factor(year)))
```

Manually Setting Discrete Colors

- You can customize the default color scales
- Or you can create your own manual scale
 - <http://colorbrewer2.org/>
 - `RColorBrewer::display.brewer.all()`
- Get RGB/HEX values from anywhere
 - <http://colormind.io/>
 - <http://color.adobe.com>
- `scale_color_manual()` expects:
 - Vector of colors
 - Named vector of colors
 - Vector of colors for levels named in breaks

```
p <- ggplot(mpg, aes(displ, hwy)) +  
  geom_point(aes(color=drv))  
  
cust_cols <- c("4"="#F2CED8",  
               "f"="#88B8B8",  
               "r"="#DE7E68")  
  
p + scale_color_manual(values=cust_cols)
```



Manually Setting Continuous Colors

- Continuous values are plotted with gradients:

```
scale_color_gradient() # 2 Color  
scale_color_gradient2() # 3 Color  
scale_color_gradientn() # N Color
```

- Examples:

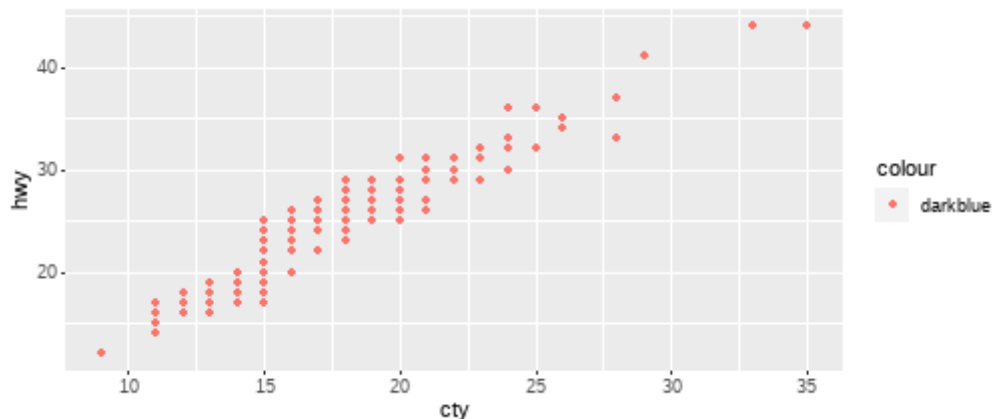
```
p <- ggplot (mpg, aes(cty, hwy)) +  
  geom_point(aes(color=displ))  
  
p + scale_color_gradient(low="white",  
                        high="orchid")  
  
p + scale_color_gradient2(low="white",  
                        high="orchid",  
                        mid="tomato")  
  
p + scale_color_gradientn(colors=c("blue",  
                                  "wheat",  
                                  "green"))
```

Setting v.s. Mapping

- Notice the difference that `color=` makes inside v.s. outside the `aes()` function
- Only things inside an `aes()` get a legend (only the mappings)
- If you have a column that has color values, use `scale_color_identity()` to prevent remapping

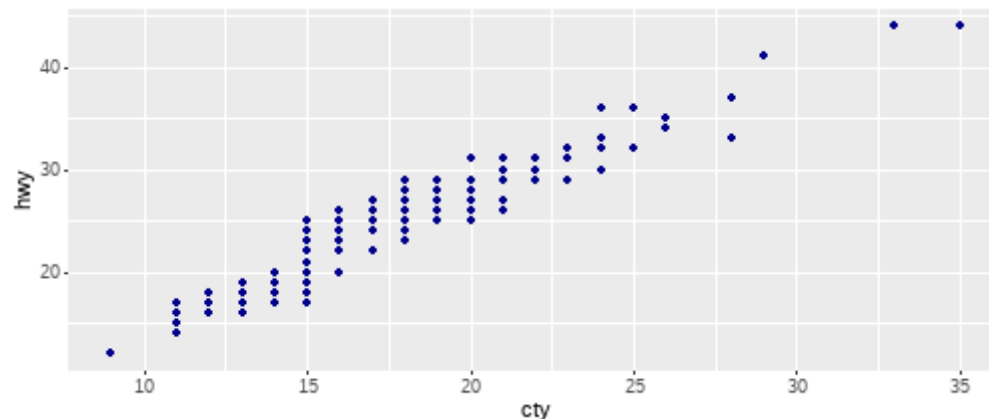
Odd Behavior

```
ggplot(mpg, aes(cty, hwy)) +  
  geom_point(aes(color = "darkblue"))
```



Correct Behavior

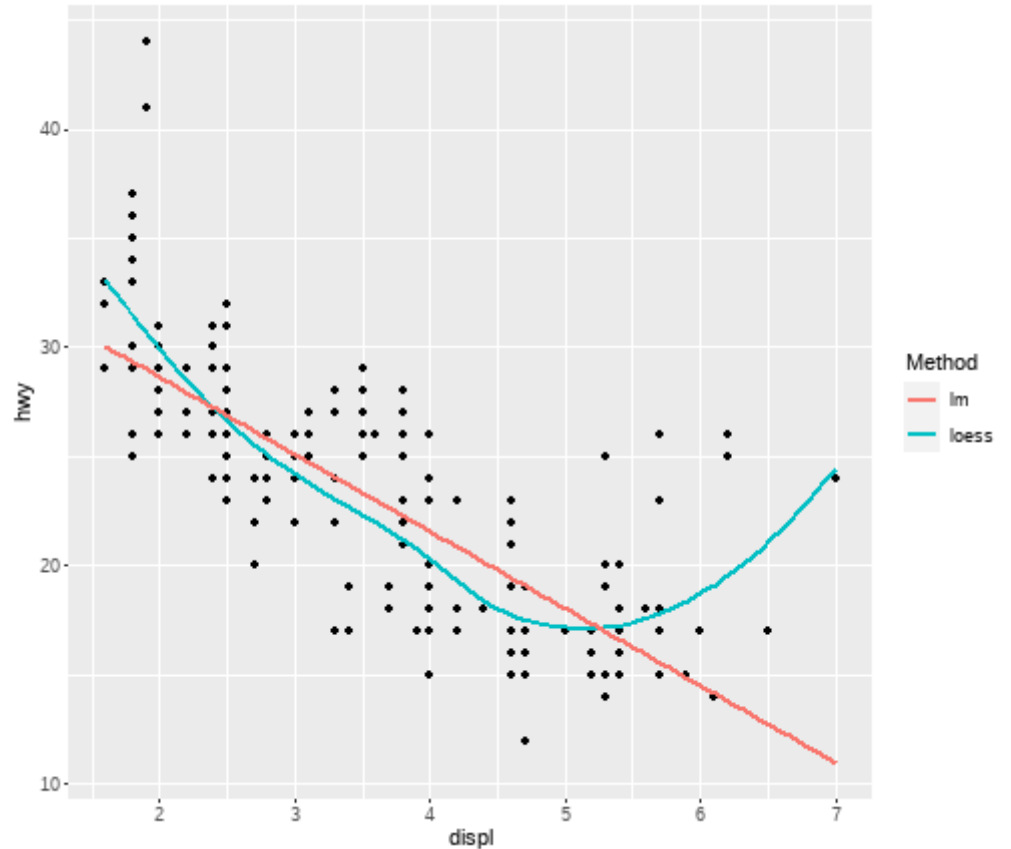
```
ggplot(mpg, aes(cty, hwy)) +  
  geom_point(color="darkblue")
```



Literal String Mappings

- Specifying a literal mapping can be useful if using multiple layers
- Here we add two layers with different smoothers
- Specify `color=` in the `aes()` for a nice legend

```
ggplot(mpg, aes(displ, hwy)) +  
  geom_point() +  
  geom_smooth(aes(color = "loess"),  
    method = "loess", se = FALSE) +  
  geom_smooth(aes(color = "lm"),  
    method = "lm", se = FALSE) +  
  labs(color = "Method")
```



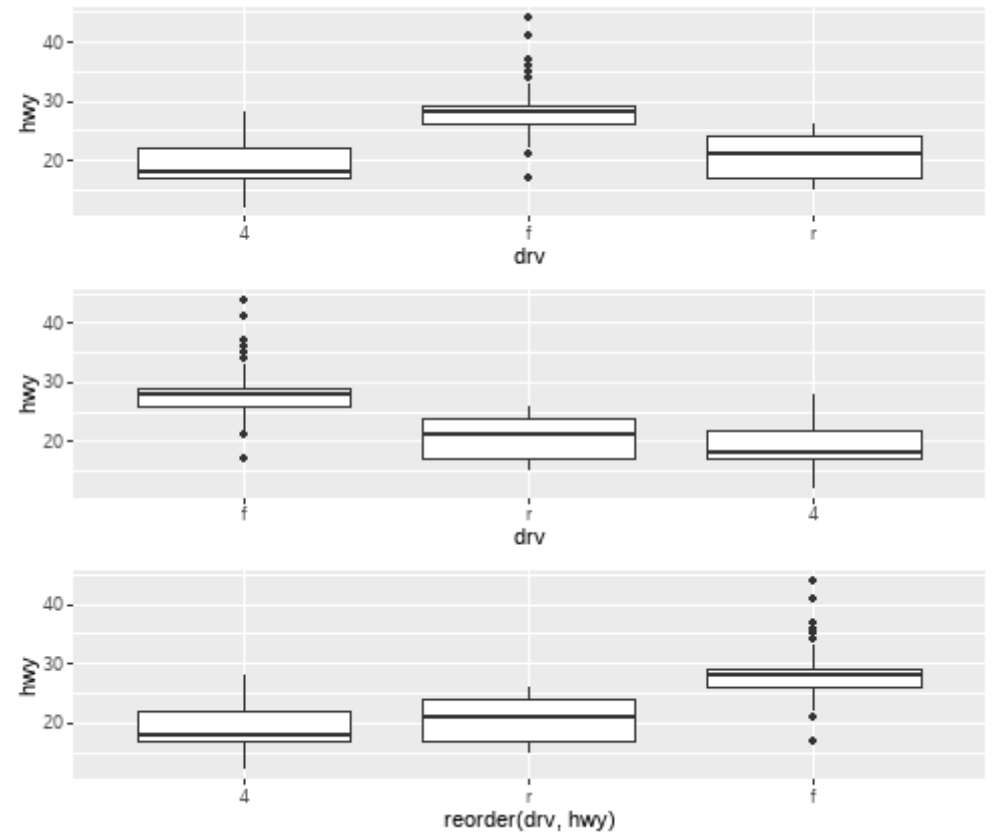
Axes are Scales!

- You can change transformations of x/y axes via scales
 - `scale_x_log10()`
 - `scale_x_sqrt()`
 - `scale_x_reverse()`
- Can also take finer control over tick marks
 - Numeric Values: `scale_x_continuous()`
 - Date/Time Values: `scale_x_datetime()`
- Control display of factor levels
 - `scale_x_discrete()`
 - Choose new labels for factor levels

Discrete Axis Plotting Order

- Discrete axes are drawn in the order of the `levels()` of the corresponding factor
- You can change that order by changing the axis scale
- Or you can re order the factor itself (see ? `reorder`)

```
p <- ggplot(mpg, aes(y=hwy))  
  
# Default  
p + geom_boxplot(aes(drv))  
  
# Use Scale  
p + geom_boxplot(aes(drv)) +  
  scale_x_discrete(limits=c("f", "r", "4"))  
  
# Use Data  
p + geom_boxplot(aes(reorder(drv, hwy)))
```



Maps

Arranging Multiple Plots

- The `patchwork` package makes it incredibly simple to combine separate ggplots into the same graphic
- Uses arithmetic operators (e.g. `+`, `|`, `/`) and order of operations to combine ggplot objects

```
library(patchwork)

p1 <- ggplot(mtcars) +
  geom_point(aes(mpg, disp))

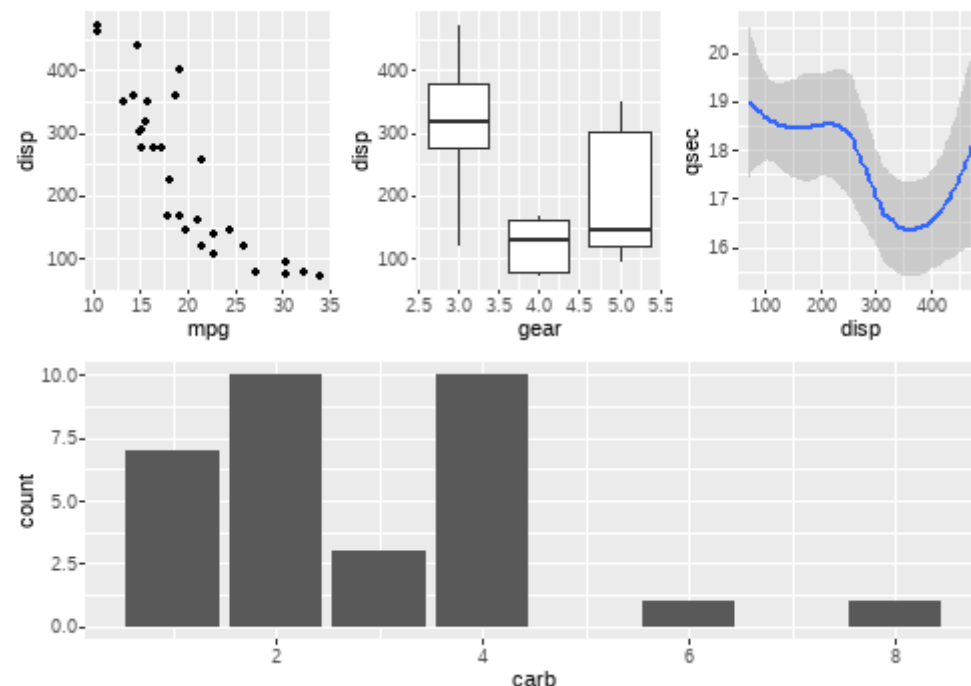
p2 <- ggplot(mtcars) +
  geom_boxplot(aes(gear, disp, group = gear))

p3 <- ggplot(mtcars) +
  geom_smooth(aes(displ, qsec))

p4 <- ggplot(mtcars) +
  geom_bar(aes(carb))

# Put them all together!

(p1 | p2 | p3) / p4
```



Vector v.s. Raster

- Two main image format categories
- Vector images:
 - Remembers the shapes drawn
 - Infinitely zoom-able
 - pdf, svg eps,
- Raster/bitmap images:
 - Remembers just the pixels of the image
 - Number of pixels depends on the dots per inch (DPI) of your image
- Typically vector is better, but if you have lots of points, raster may be easier to work with

ggsave()

- Once you've made your masterpiece, use `ggsave()` to save it
- It will create a file in your current working directory (`getwd()/setwd()`)
- Saves last plot printed
- Looks at file name to determine type:

```
ggsave("plot.pdf"); ggsave("plot.eps");  
ggsave("plot.png"); ggsave("plot.jpg");  
ggsave("plot.tiff"); ggsave("plot.svg");
```

- Can also pass plot object to save any plot:

```
ggsave("plot.png", ggplot(mpg, aes(cty, hwy)) + geom_point())
```

Thank You!