

README

- Review of the 1st chapter of the DataCamp course
- Add your name and (`pledged`) in the `#+AUTHOR:` meta headline
- When you've completed the file, submit it in Canvas
- You'll get solutions after the deadline has passed

DONE Identify yourself

1. Add your name and (`pledged`) at the top
2. Run the `#+PROPERTY` and `#+STARTUP` lines with `C-c C-c`
3. You should open the R session in the same directory as this file
4. To check, run `getwd()` in the R console window

DONE Answer conceptual questions

- **What is the requirement for this course and where can you find this information?** - "Introduction to the Tidyverse" (Bottom of dashboard)

This is an introduction to the programming language R, focused on a powerful set of tools known as the Tidyverse. You'll learn the intertwined processes of data manipulation and visualization using the tools `dplyr` and `ggplot2`. You'll learn to manipulate data by filtering, sorting, and summarizing a real dataset of historical country data in order to answer exploratory questions. You'll then learn to turn this processed data into informative line plots, bar plots, histograms, and more with the `ggplot2` package. You'll get a taste of the value of exploratory data analysis and the power of Tidyverse tools. This is a suitable introduction for those who have no previous experience in R and are interested in performing data analysis.

- **What is the "Tidyverse"?** It's a bundle of R packages including `ggplot2` (which predates the "Tidyverse" by several years), `dplyr` for data frame manipulation, and many more. Its functions rely on data being "tidy", which corresponds to Codds 3rd normal form for relational or tabular data.



(Image source: hbctraining.github.io)

- **Base R is the foundation that every data scientist should know.**

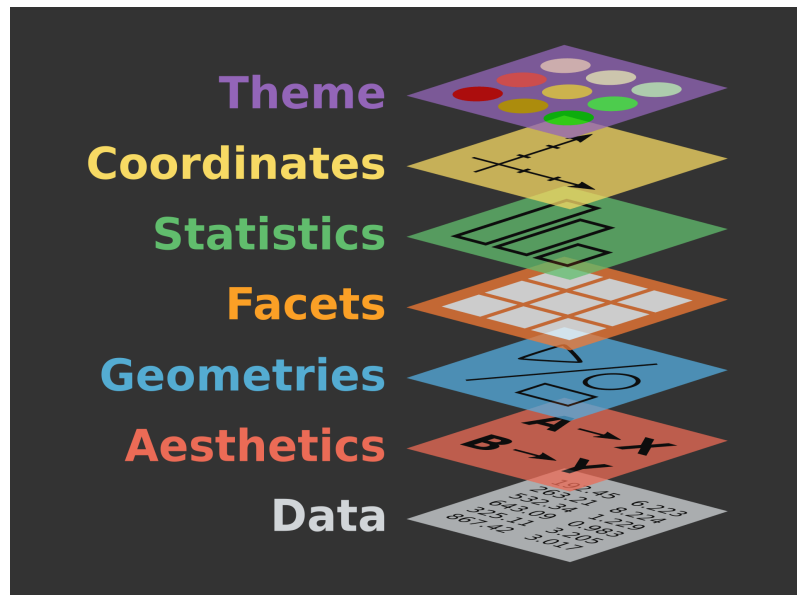
"If the user knows base-R (not difficult), she can handle any situation with just a few simple operations. The old adage applies: "Give a man a fish, and he can eat for a day. Teach him how to fish, and he can eat for a lifetime." From: "TidyverseSceptic" (Matloff, 2022)

- **What is the "Grammar of Graphics"?**

The "Grammar of Graphics" (gg) is a plotting framework by Leland Wilkinson (1999) implemented in R's `ggplot2` plotting package. Its core **ideas** are:

1. Graphics are distinct layers of 'grammatical' elements
2. Plots are given meaning through 'aesthetic' mappings

- **What are the layers of the 'Grammar of Graphics'?**



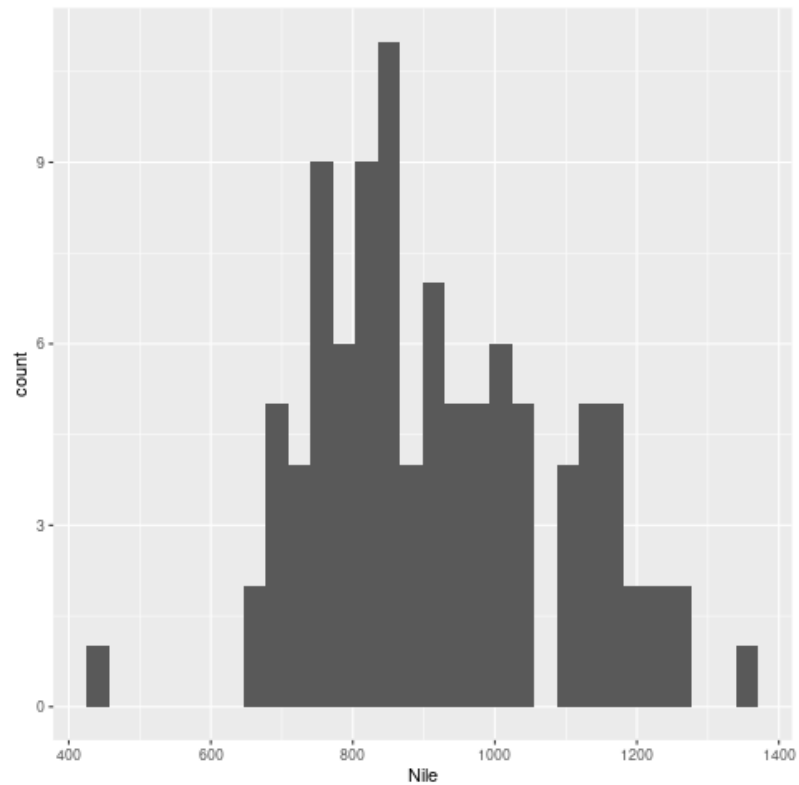
(Image source: r.qcbs.ca)

- What are some examples for these elements in ggplot2?

Data	{variables of interest}				
Aesthetics	<i>x-axis</i> <i>y-axis</i>	<i>colour</i> <i>fill</i>	<i>size</i> <i>labels</i>	<i>alpha</i> <i>shape</i>	<i>line width</i> <i>line type</i>
Geometries	<i>point</i>	<i>line</i>	<i>histogram</i>	<i>bar</i>	<i>boxplot</i>
Themes	<i>non-data ink</i>				
Statistics	<i>binning</i>	<i>smoothing</i>	<i>descriptive</i>	<i>inferential</i>	
Coordinates	<i>cartesian</i>	<i>fixed</i>	<i>polar</i>	<i>limits</i>	
Facets	<i>columns</i>	<i>rows</i>			

- Can you save a ggplot2 plot as an R object?

```
library(ggplot2)
g <- ggplot(
  data = data.frame(Nile),
  aes(Nile)) +
  geom_histogram()
g
```



- Show the structure and attributes of `g`:

```
attributes(g)
str(g)
```

```
$names
 [1] "data"      "layers"    "scales"    "guides"    "mapping"   "theme"
 [7] "coordinates" "facet"     "plot_env"  "layout"    "labels"

$class
 [1] "gg"      "ggplot"
List of 11
 $ data      :'data.frame': 100 obs. of  1 variable:
  ..$ Nile: Time-Series [1:100] from 1871 to 1970: 1120 1160 963 1210 1160 1160 83
 $ layers    :List of 1
  ..$ :Classes 'LayerInstance', 'Layer', 'ggproto', 'gg' <ggproto object: Class L
```

```

aes_params: list
compute_aesthetics: function
compute_geom_1: function
compute_geom_2: function
compute_position: function
compute_statistic: function
computed_geom_params: list
computed_mapping: uneval
computed_stat_params: list
constructor: call
data: waiver
draw_geom: function
finish_statistics: function
geom: <ggproto object: Class GeomBar, GeomRect, Geom, gg>
aesthetics: function
default_aes: uneval
draw_group: function
draw_key: function
draw_layer: function
draw_panel: function
extra_params: just na.rm orientation
handle_na: function
non_missing_aes: xmin xmax ymin ymax
optional_aes:
parameters: function
rename_size: TRUE
required_aes: x y
setup_data: function
setup_params: function
use_defaults: function
super: <ggproto object: Class GeomRect, Geom, gg>
  geom_params: list
  inherit_aes: TRUE
  layer_data: function
  map_statistic: function
  mapping: NULL
  position: <ggproto object: Class PositionStack, Position, gg>
compute_layer: function
compute_panel: function
fill: FALSE

```

```

required_aes:
reverse: FALSE
setup_data: function
setup_params: function
type: NULL
vjust: 1
super: <ggproto object: Class Position, gg>
  print: function
  setup_layer: function
  show.legend: NA
  stat: <ggproto object: Class StatBin, Stat, gg>
aesthetics: function
compute_group: function
compute_layer: function
compute_panel: function
default_aes: uneval
dropped_aes: weight
extra_params: na.rm orientation
finish_layer: function
non_missing_aes:
optional_aes:
parameters: function
required_aes: x|y
retransform: TRUE
setup_data: function
setup_params: function
super: <ggproto object: Class Stat, gg>
  stat_params: list
  super: <ggproto object: Class Layer, gg>
$ scales      :Classes 'ScalesList', 'ggproto', 'gg' <ggproto object: Class Scales
  add: function
  add_defaults: function
  add_missing: function
  backtransform_df: function
  clone: function
  find: function
  get_scales: function
  has_scale: function
  input: function
  map_df: function

```

```

n: function
non_position_scales: function
scales: list
train_df: function
transform_df: function
super: <ggproto object: Class ScalesList, gg>
$ guides      :Classes 'Guides', 'ggproto', 'gg' <ggproto object: Class Guides, gg>
  add: function
  assemble: function
  build: function
  draw: function
  get_custom: function
  get_guide: function
  get_params: function
  get_position: function
  guides: NULL
  merge: function
  missing: <ggproto object: Class GuideNone, Guide, gg>
add_title: function
arrange_layout: function
assemble_drawing: function
available_aes: any
build_decor: function
build_labels: function
build_ticks: function
build_title: function
draw: function
draw_early_exit: function
elements: list
extract_decor: function
extract_key: function
extract_params: function
get_layer_key: function
hashables: list
measure_grobs: function
merge: function
override_elements: function
params: list
process_layers: function
setup_elements: function

```

```

setup_params: function
train: function
transform: function
super: <ggproto object: Class GuideNone, Guide, gg>
  package_box: function
  print: function
  process_layers: function
  setup: function
  subset_guides: function
  train: function
  update_params: function
  super: <ggproto object: Class Guides, gg>
$ mapping      :List of 1
..$ x: language ~Nile
.. ..- attr(*, ".Environment")=<environment: R_GlobalEnv>
..- attr(*, "class")= chr "uneval"
$ theme        : list()
$ coordinates:Classes 'CoordCartesian', 'Coord', 'ggproto', 'gg' <ggproto object
  aspect: function
  backtransform_range: function
  clip: on
  default: TRUE
  distance: function
  expand: TRUE
  is_free: function
  is_linear: function
  labels: function
  limits: list
  modify_scales: function
  range: function
  render_axis_h: function
  render_axis_v: function
  render_bg: function
  render_fg: function
  setup_data: function
  setup_layout: function
  setup_panel_guides: function
  setup_panel_params: function
  setup_params: function
  train_panel_guides: function

```



```

transform: function
super: <ggproto object: Class CoordCartesian, Coord, gg>
$ facet      :Classes 'FacetNull', 'Facet', 'ggproto', 'gg' <ggproto object: Class
compute_layout: function
draw_back: function
draw_front: function
draw_labels: function
draw_panels: function
finish_data: function
init_scales: function
map_data: function
params: list
setup_data: function
setup_params: function
shrink: TRUE
train_scales: function
vars: function
super: <ggproto object: Class FacetNull, Facet, gg>
$ plot_env   :<environment: R_GlobalEnv>
$ layout     :Classes 'Layout', 'ggproto', 'gg' <ggproto object: Class Layout, gg>
coord: NULL
coord_params: list
facet: NULL
facet_params: list
finish_data: function
get_scales: function
layout: NULL
map_position: function
panel_params: NULL
panel_scales_x: NULL
panel_scales_y: NULL
render: function
render_labels: function
reset_scales: function
resolve_label: function
setup: function
setup_panel_guides: function
setup_panel_params: function
train_position: function
super: <ggproto object: Class Layout, gg>

```

```

$ labels      :List of 3
..$ x        : chr "Nile"
..$ y        : chr "count"
.. ..- attr(*, "fallback")= logi TRUE
..$ weight: chr "weight"
.. ..- attr(*, "fallback")= logi TRUE
- attr(*, "class")= chr [1:2] "gg" "ggplot"

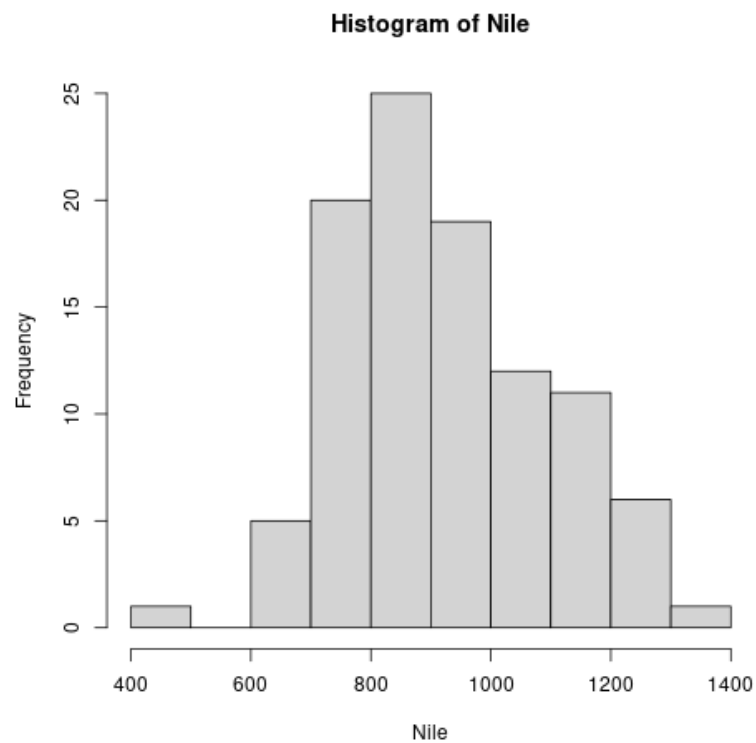
```

- Can you save a base R plot as an R object?

```

h <- hist(Nile)
h

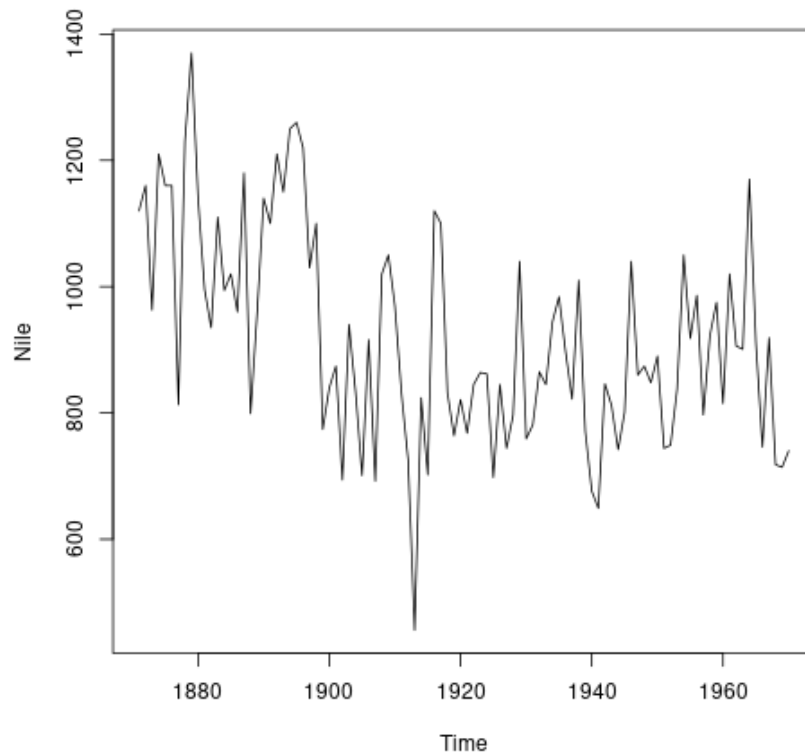
```



```

p <- plot(Nile)
p

```



```
attributes(h) # saving histogram to access its attributes and
# structure
str(h)
```

```
$names
[1] "breaks" "counts" "density" "mids" "xname" "equidist"
```

```
$class
[1] "histogram"
```

```
List of 6
```

```
$ breaks : int [1:11] 400 500 600 700 800 900 1000 1100 1200 1300 ...
```

```
$ counts : int [1:10] 1 0 5 20 25 19 12 11 6 1
```

```
$ density : num [1:10] 0.0001 0 0.0005 0.002 0.0025 0.0019 0.0012 0.0011 0.0006 0.000
```

```
$ mids : num [1:10] 450 550 650 750 850 950 1050 1150 1250 1350
```

```
$ xname      : chr "Nile"
$ equidist: logi TRUE
- attr(*, "class")= chr "histogram"
```

- **Can you combine ggplot2 and base R graphics in one plot array?** Answer: no. Base R graphics and ggplot2 graphics are completely different and cannot be mixed. In base R, plots are created by opening graphics devices, in ggplot2, plots are layered R objects.

DONE Create simple scatterplots

We're going to work with `MASS::mammals` using `ggplot2` and base R.

1. Load the relevant packages.

```
library(MASS)
library(ggplot2)
```

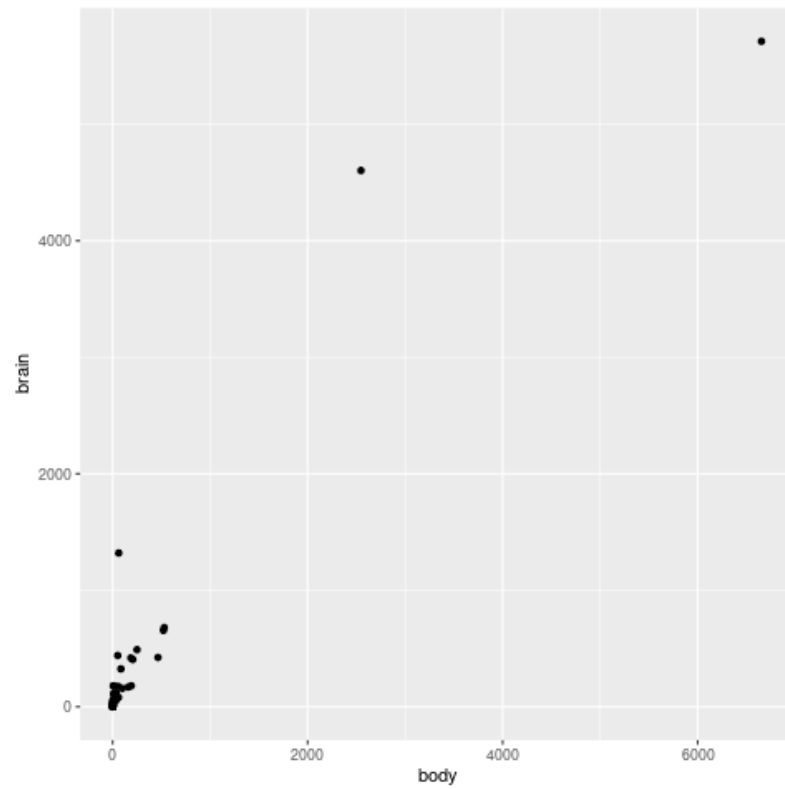
2. Show the data structure of `mammals`.

```
str(mammals)

'data.frame': 62 obs. of  2 variables:
 $ body : num  3.38 0.48 1.35 465 36.33 ...
 $ brain: num  44.5 15.5 8.1 423 119.5 ...
```

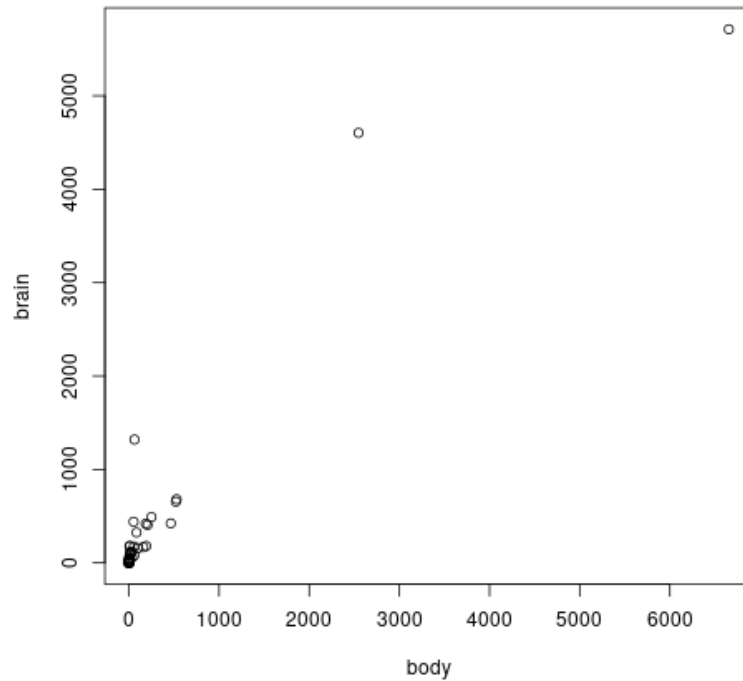
3. Create a scatterplot of `brain` vs `body` of the `mammals` data set in `ggplot2`.

```
ggplot(mammals, aes(x=body,y=brain)) +
  geom_point()
```



4. Create a scatterplot of `brain` vs `body` of the `mammals` data set in base R.

```
plot(mammals)
```



DONE Transform plots

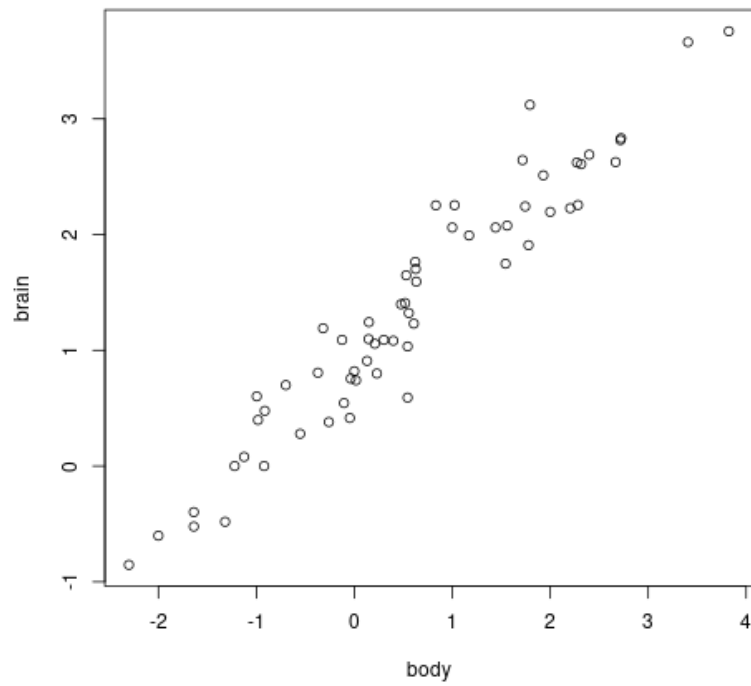
1. What's the problem with these plots and what could you do about it?

The problem: the data points are too bunched up because mammals have a wide spectrum of body and brain weights (there are very small and very large ones).

The solution: transform the x- and y-axis logarithmically.

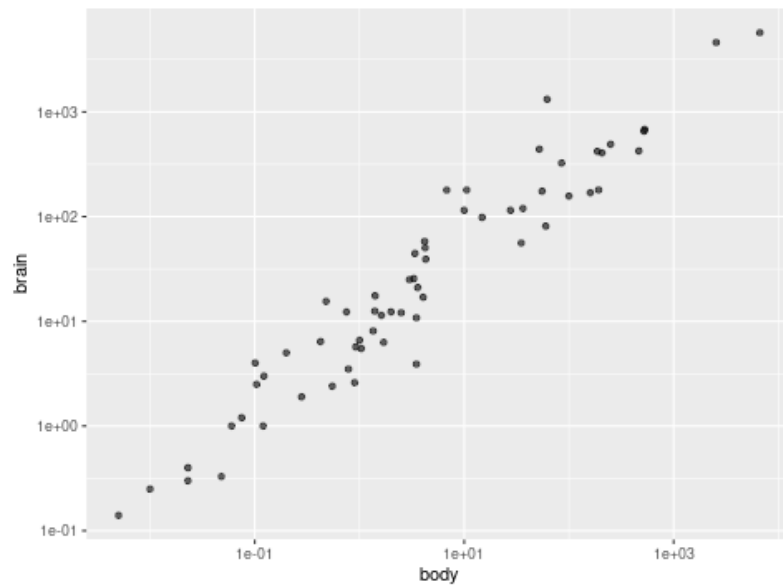
2. Implement the solution with `plot`.

```
plot(log10(mammals))
```



3. Implement the solution with `ggplot` - save the plot as `gg` for later, and print it.

```
gg <- ggplot(  
  data=mammals,  
  aes(x=body,y=brain)) +  
  geom_point(alpha=0.6) +  
  coord_fixed() +  
  scale_x_log10() +  
  scale_y_log10()  
gg
```



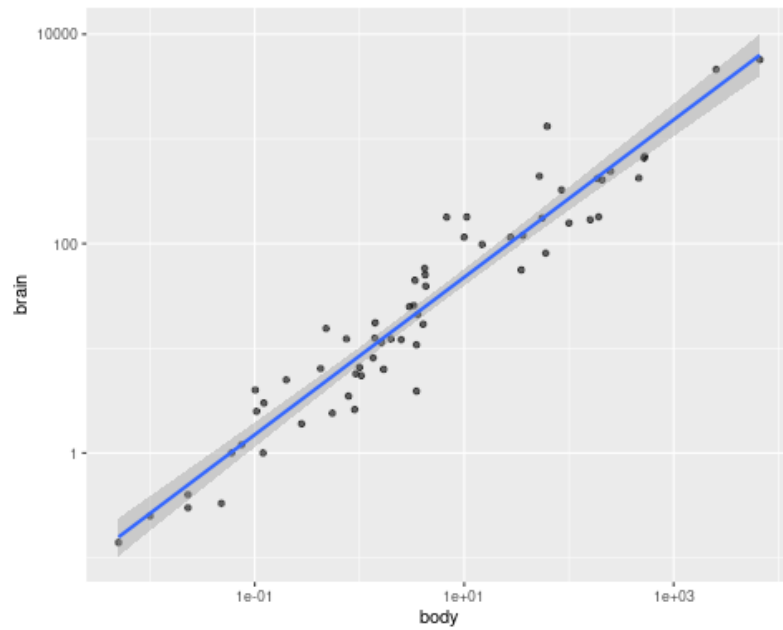
4. What does the `geom_point` argument `alpha` do?

Answer: it reduces the transparency of the points by 40%.

DONE Create trendlines with ggplot2 and base R

1. Create a linear trendline for the `ggplot2` plot `gg`. Inside the smoothing geometry, use `method="lm"` to fix the model.

```
gg +  
  geom_smooth(method="lm")
```

2. Create a linear model in **base R** using `lm` and `data=log10(mammals)`. Save it as `line` and print it.

```
line <- lm(brain ~ body, data=log10(mammals))
line
```

Call:

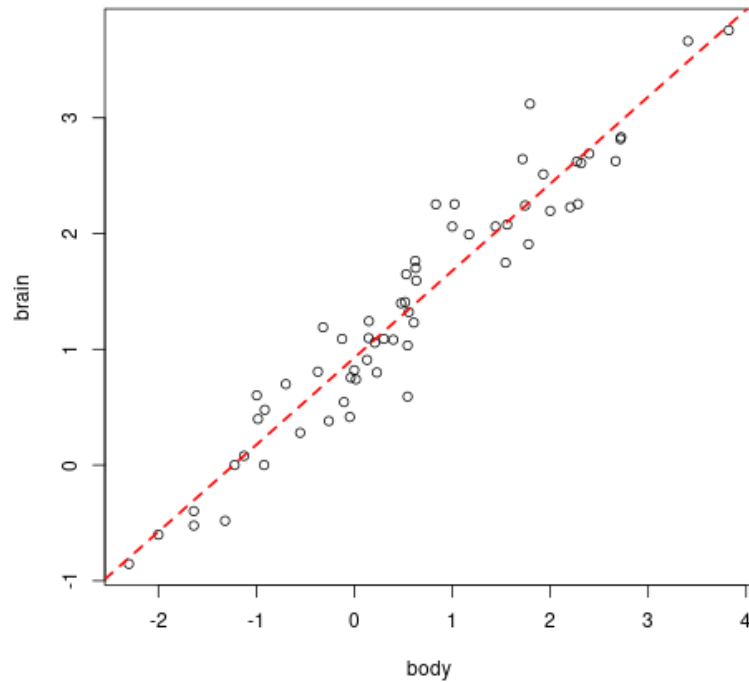
```
lm(formula = brain ~ body, data = log10(mammals))
```

Coefficients:

(Intercept)	body
0.9271	0.7517

3. Create a trendline plot in **base R** using the linear model. The line should be red, dashed and double wide.

```
plot(log10(mammals))
abline(line, col="red", lty=2, lwd=2)
```

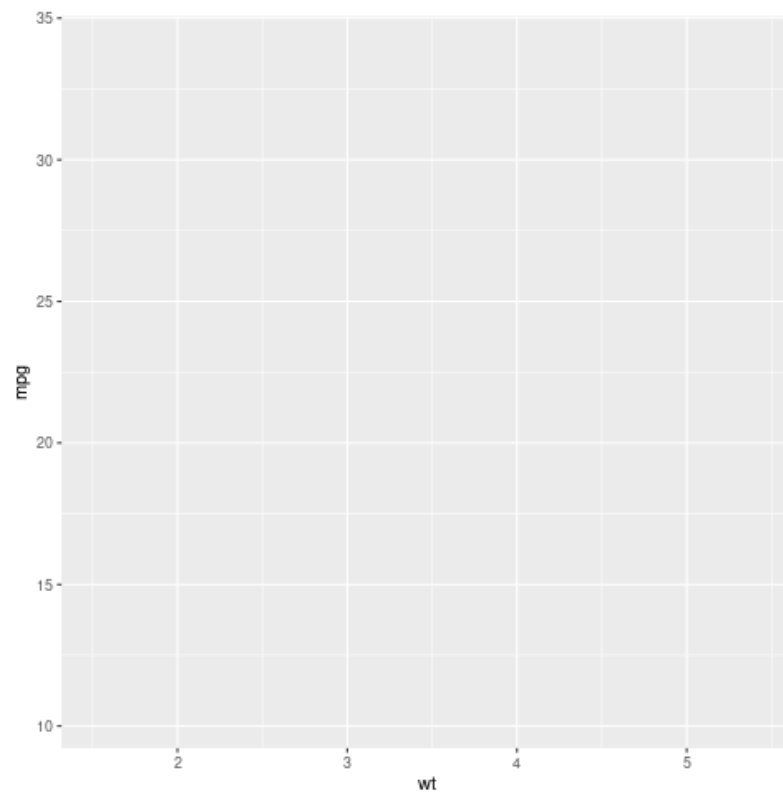


DONE Map 'aesthetics' to variables

Recall that the `mtcars` data frame lists the characteristics mileage (`mpg`), weight (`wt`) and number of cylinders (`cyl`) as **numeric** variables.

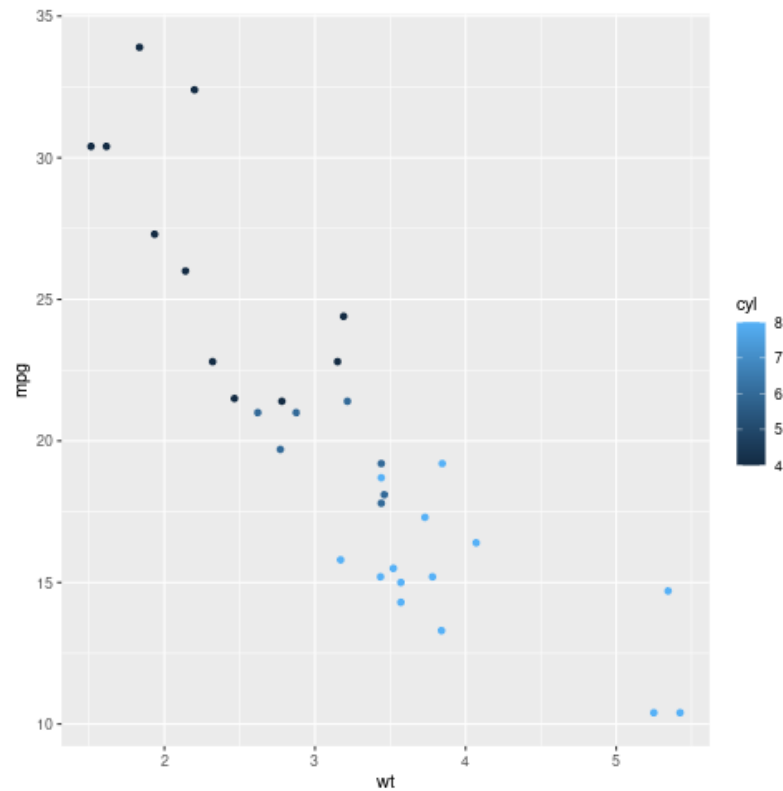
1. Create a `ggplot` of mileage vs. weight using `ggplot2`, save it as `gg` and print it.

```
gg <- ggplot(data=mtcars, aes(wt,mpg))
gg
```



2. Create a scatterplot where the color 'aesthetic' is mapped to the number of cylinders by adding a 'geometry' to gg.

```
ggplot(data=mtcars,  
  aes(wt,mpg)) +  
  geom_point(  
    aes(color=cyl))
```



3. What's the difference between mapping 'aesthetics' inside the 'geometry' or inside the `ggplot` function?

Answer: the `aes` function knows about the dataset from `data`. You can also pipe the data set into the function using `|>` or `%>%`. Without `aes`, you need to specify `geom_point(mtcars$cyl)`. You can subset data for a specific geometry by putting `aes` into the function.