

# Before the workshop

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1. Download R :

<https://cloud.r-project.org/>

2. Download Rstudio Desktop (Open Source License):

<https://www.rstudio.com/products/rstudio/download/>

3. Access the workshop materials:

- PowerPoint Presentation
- Sample Code

# Introduction to R

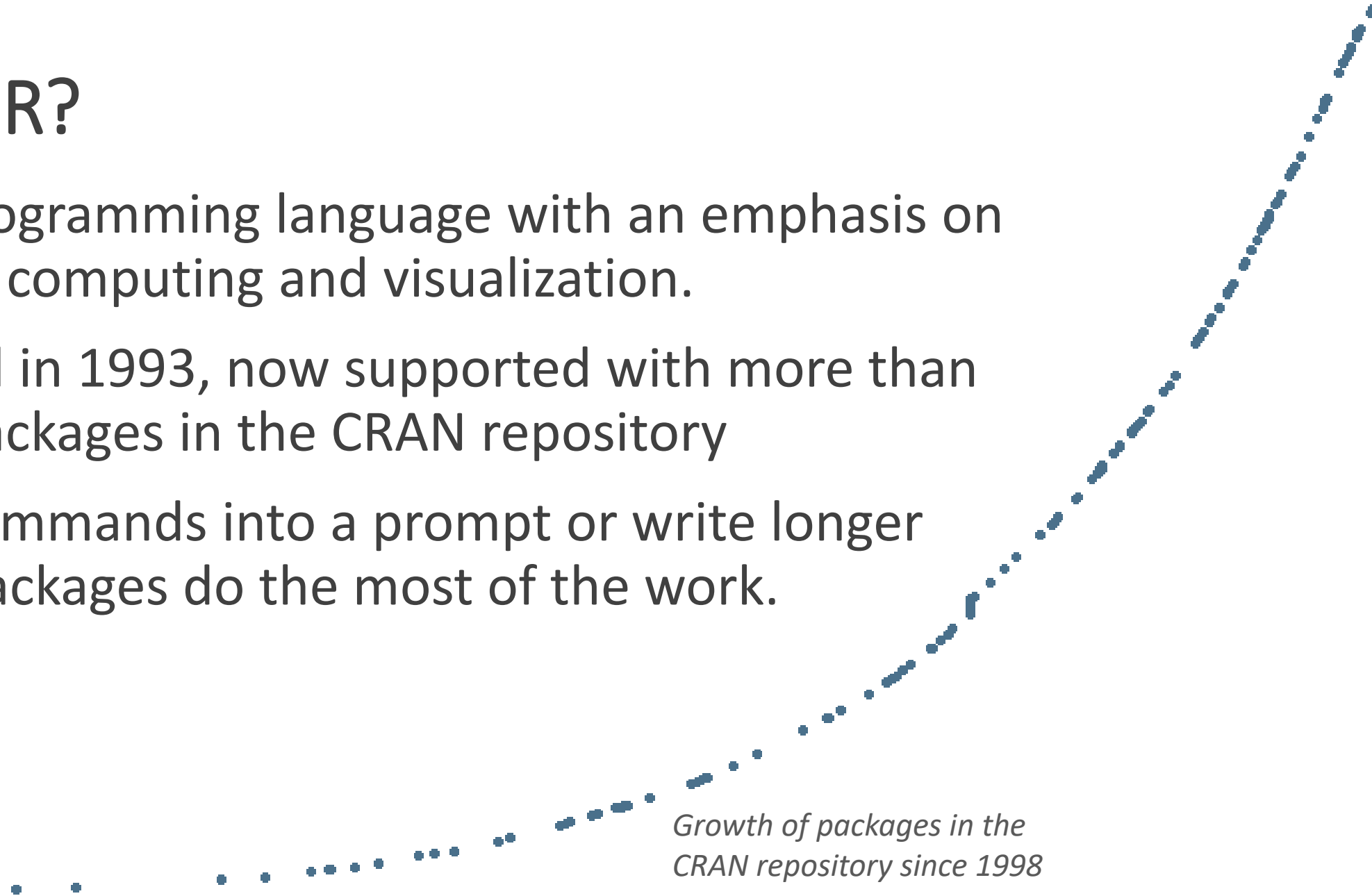
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SCIR XXX

September YY, 2022

# What Is R?

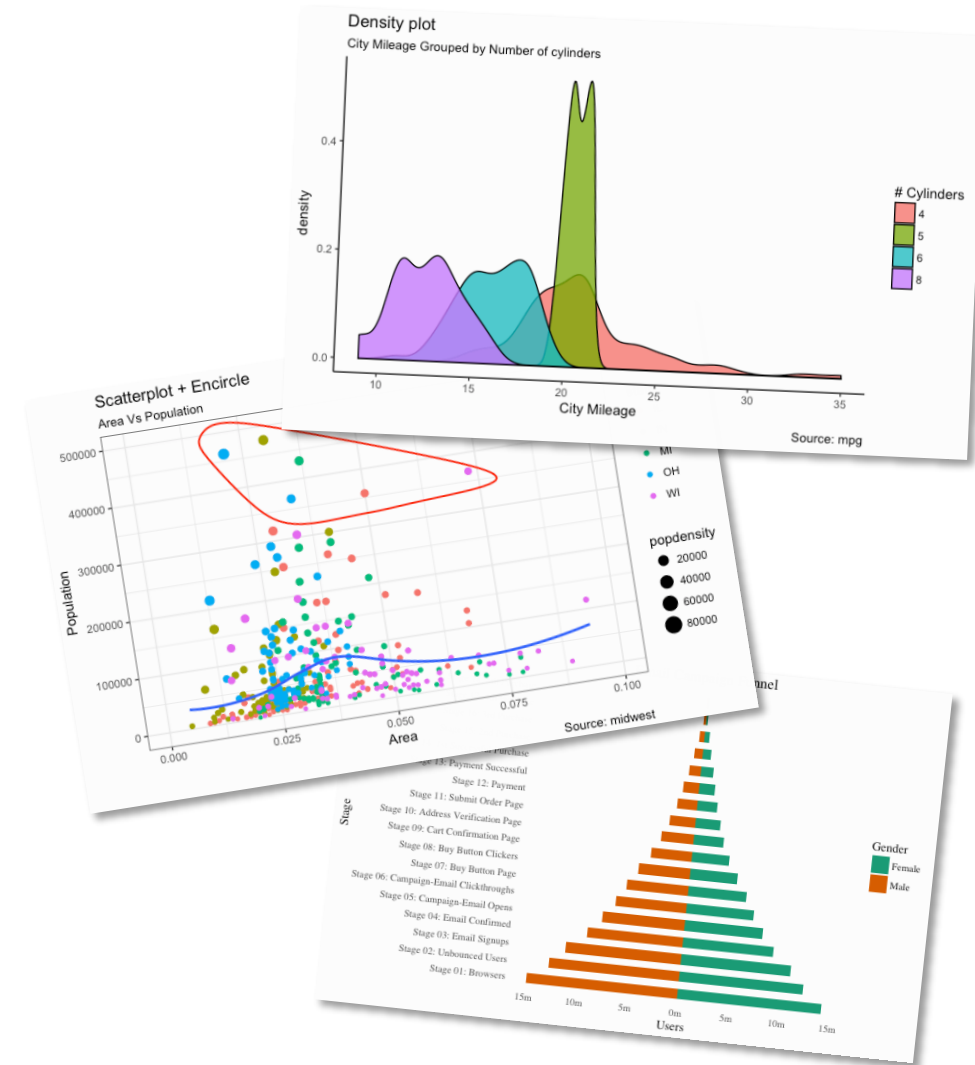
- Free programming language with an emphasis on statistical computing and visualization.
- Created in 1993, now supported with more than 14,000 packages in the CRAN repository
- Type commands into a prompt or write longer scripts. Packages do the most of the work.



*Growth of packages in the  
CRAN repository since 1998*

# Why learn R?

- Open source
- Works with bigger and more diverse data sets than Excel
- Repeat the same analysis on different data using scripts
- Perform sophisticated statistical functions, text mining, data visualization
- Specialized packages have been created for specific disciplines (e.g. economics, biology)



Sample visualizations with ggplot2.  
Source: <http://r-statistics.co/Top50-Ggplot2-Visualizations-MasterList-R-Code.html>

# Disadvantages of R

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- Syntax is not intuitive
- Documentation can be hard to interpret (but it's all in one place and formatted the same way)
- Many packages use a lot of memory and processing power

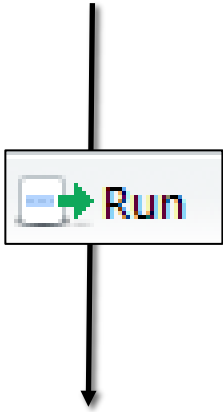
# Today we will learn:

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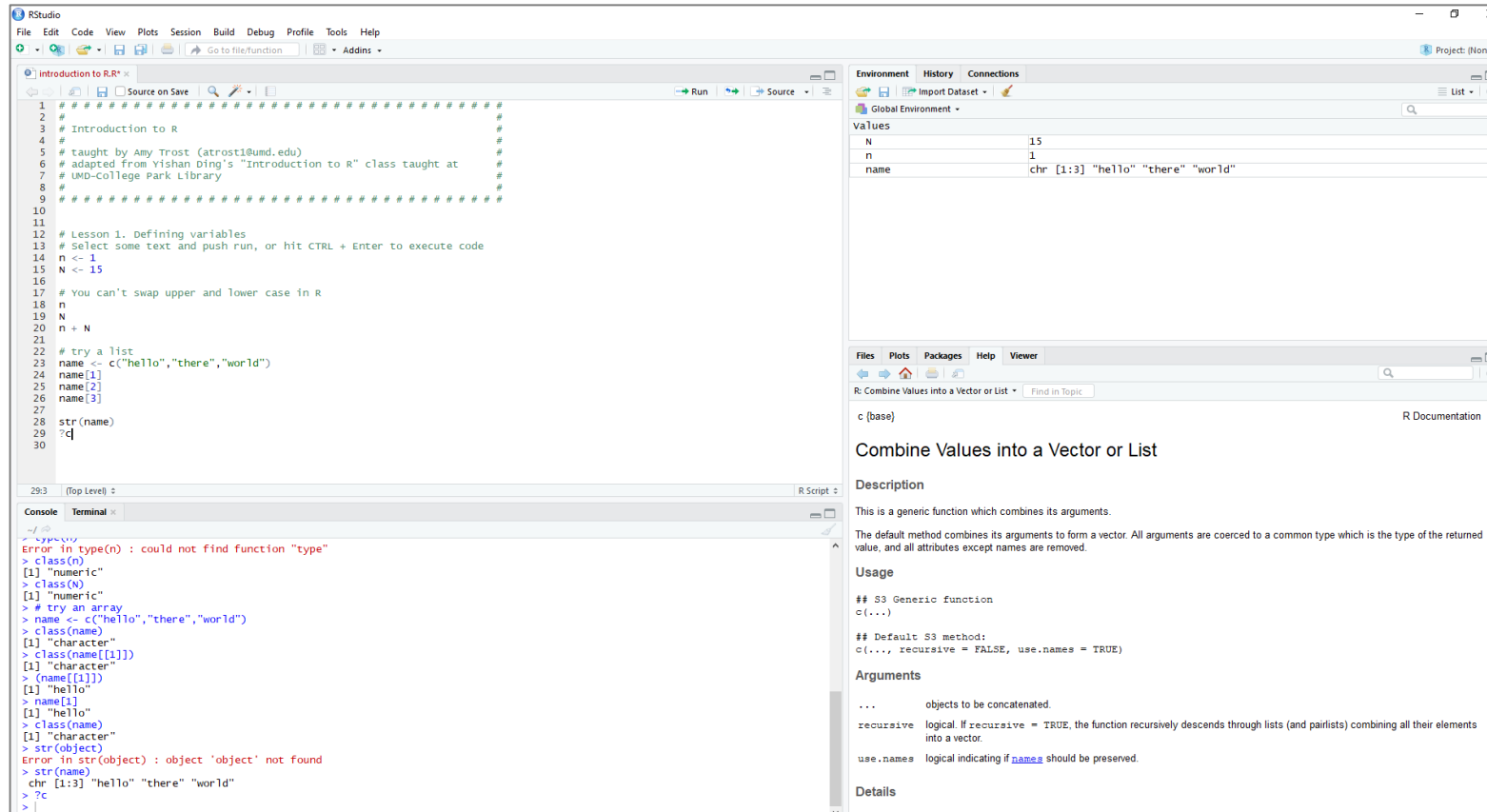
- ☐ **Basic Commands**
- ☐ **Importing Data**
- ☐ **Data Types**
- ☐ **Manipulating Data**
- ☐ **Visualizing Data**

# RStudio: an integrated development environment (IDE) for R

Source editor: Text editor for scripts.



Console: Execute code and examine the log.  
-- Can also type commands here too!

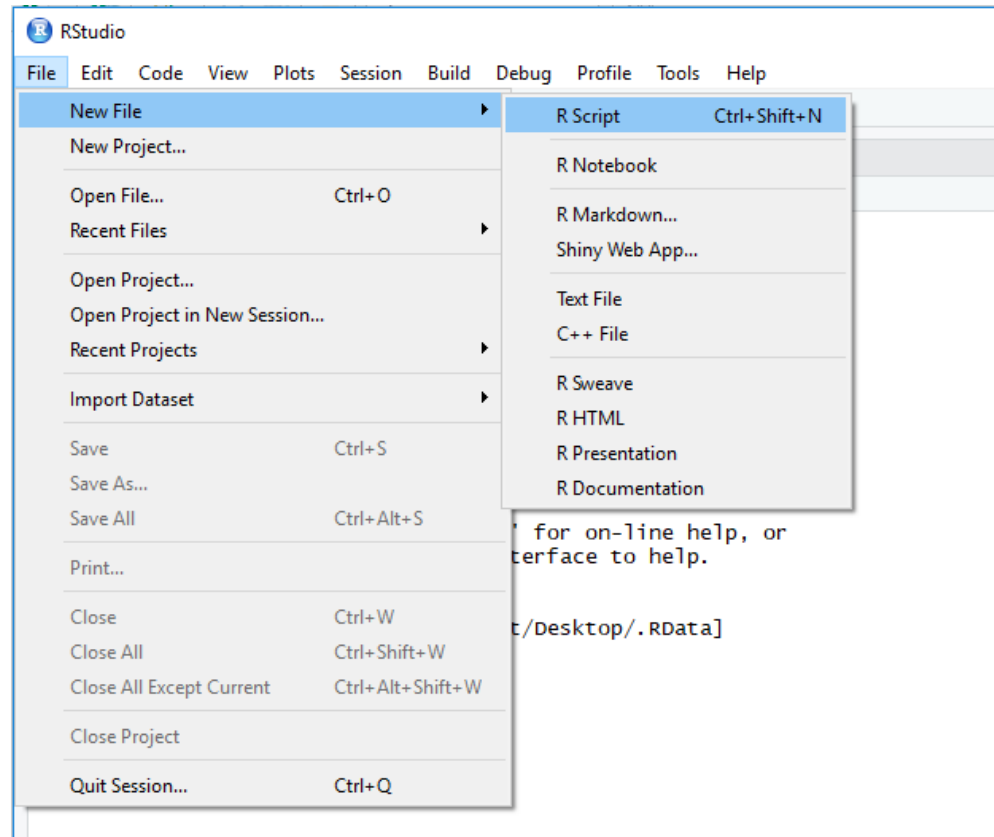


Objects: names, properties, and variables

Library: View plots, install packages, access help



# Getting Started—Launching a script



*Scripts allow you to edit commands, save your work, and execute many commands at once*



# Getting Started--Basic Commands

# marks a  
comment; R  
ignores these



```
# Assigning values to variables
n <- 1
N <- 15

n
N      # You can't swap upper and lower case in R
n + N

# try a list
name <- c("hello", "there", "world")
name[1]
name[2]
name[3]

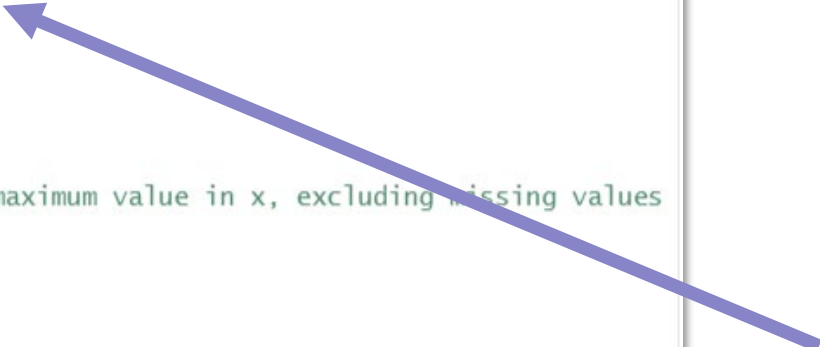
# try a simple function
test_scores <- c(90, 72, 85, 93, 84)
test_average <- mean(test_scores)

## -----
## Statistical Functions
## -----

max(test_scores)      # Find the maximum value in x, excluding missing values
min(test_scores)      # minimum
mean(test_scores)     # mean
median(test_scores)   # median
sum(test_scores)      # sum
var(test_scores)      # variance
sd(test_scores)       # standard deviation

# get help for a command
?mean
```

Try typing some  
of this in to your  
console (or just  
run the sample  
code)



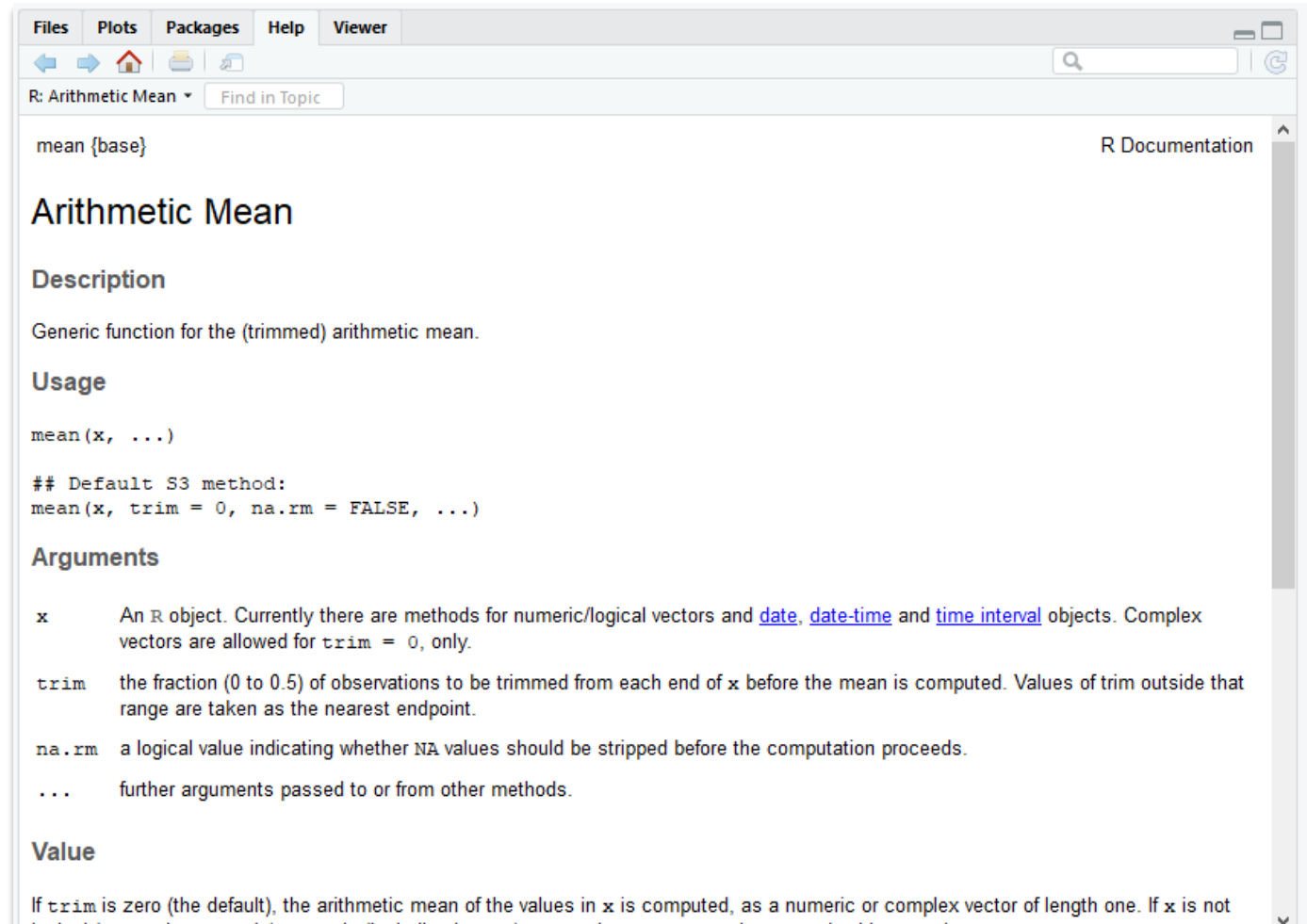
# Statistical Functions

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- ❑ `max(x, na.rm = TRUE)`
- ❑ `min(x, na.rm = TRUE)`
- ❑ `mean(x, na.rm = TRUE)`
- ❑ `median(x, na.rm = TRUE)`
- ❑ `sum(x, na.rm = TRUE)`
- ❑ `sd(x, na.rm = TRUE)`

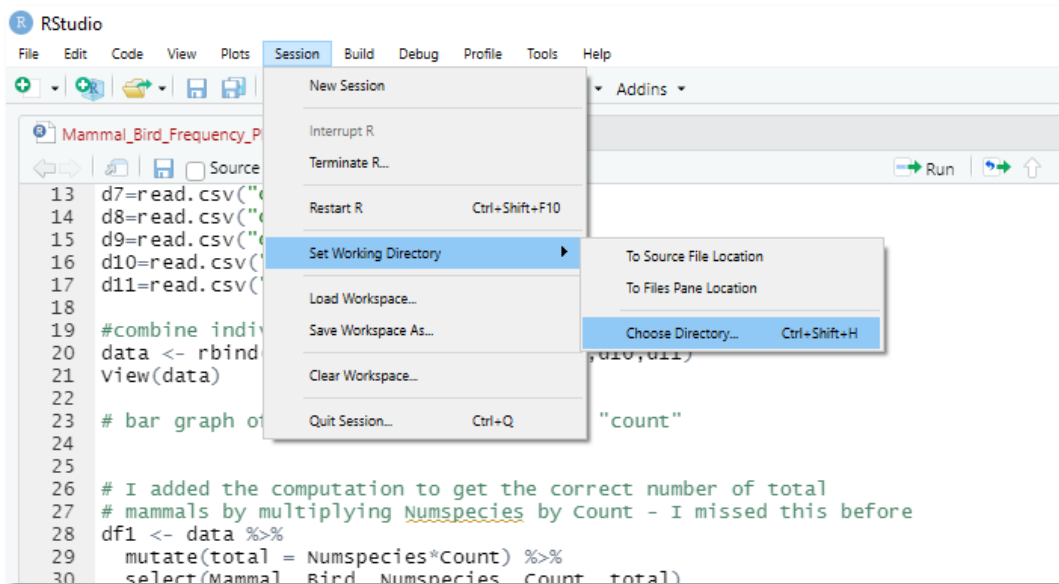
# Getting Help

A question mark before a command will launch a help page in Rstudio, for example, typing “?mean” launches



# Importing Data (2 files for this session)

1. **Set Working Directory** (under Session) to choose the folder where your data is.
2. **Use the read.csv command:** `species <- read.csv("wikipedia species list.csv")`  
`countries <- read.csv("worldbank google country data.csv")`



*You can also use read.csv on the full file path, e.g.*

```
countries <- read.csv("C:/Users/atrost/Desktop/worldbank  
google country data.csv")
```

*This can get messy, though, and on windows you need to make sure the slashes are forward / and not backward \*

# Explore the dataset

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- ❑ `view()` open the dataset in a new window
- ❑ `class()` type of data set
- ❑ `head()` preview the dataset in console
- ❑ `dim()` number of rows & columns (dimension) of data set
- ❑ `nrow()` number of rows
- ❑ `ncol()` number of columns
- ❑ `colnames()` column names
- ❑ `rownames()` row names
- ❑ `summary()` shows type and distribution of all of the variables in a data set

# Selecting elements of a data set

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## 1. Subset by row/column index

**DATA [ROW, COLUMN]**

- ❑ `countries[6,]`
- ❑ `countries[6:10,]`
- ❑ `countries[,1]`
- ❑ `countries[10:20, 1:3]`

## 2. Subset by variable name

**DATA\$VARIABLE\_NAME**

- ❑ `species$common_name`
- ❑ `species$common_name[30:55]`

# Data Types

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	Class	Example
1	Integer	<ul style="list-style-type: none"><li>• 3, 4, 17484, -65</li></ul>
2	Numeric	Can be a decimal, irrational: <ul style="list-style-type: none"><li>• 3</li><li>• 3.0</li><li>• 3.14159</li></ul>
3	Character	"a", "b", "Montgomery College"
4	Logical	<ul style="list-style-type: none"><li>• TRUE</li><li>• FALSE</li></ul>
5	Complex	<ul style="list-style-type: none"><li>• <math>1 + 4i</math></li></ul>

## How to know the class?

- `class()` *what kind of object is it (high-level)?*
- `typeof()` *what is the object's data type (low-level)?*
- `length()` *how long is it? What about two dimensional objects?*
- `attributes()` *does it have any metadata?*

## How to change the class?

- `as.numeric()`, `as.logical()`, `as.character()`, etc.

# Some data structures

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	Structure type	What is this?
1	Vector	A simple group of numbers, characters, or logical values
2	List	Flexible, sometimes complicated, stores many types of variables nested within each other.
3	Matrix	Like a vector with more dimensions
4	Data Frame	Very useful!! Represents a data table, with rows and columns of information.
5	Factors	Represent categories of information
6	Tibble	A specialized data frame-like object that is actually a list; used in tidyr and some other packages.



View data types  
in the  
“environment”  
pane of RStudio

Environment	History	Connections	Tutorial
Import Dataset 185 MiB			
R Global Environment			
Data			
countries	212 obs. of 9 variables		
\$ Country	: chr "Solomon Islands" "Suriname" "Samoa" "Vanuatu" ...		
\$ Region	: chr "EAST ASIA AND PACIFIC\		
\$ Income.Classification	: chr "LOWER-MIDDLE INCOME" "UPPER-MIDDLE-INCOME" "LOWER-MIDDLE ...		
\$ Avg.elevation.in.m	: int 216 162 435 250 82 600 273 283 218 168 ...		
\$ X..covered.by.cropland	: num 0 1.1 0 0 0.3 1.2 0 6.7 1.4 5.3 ...		
\$ X..covered.by.trees	: num 71.9 68.4 67.8 67.5 66.9 64.4 61.2 60.1 58.9 57.3 ...		
\$ mean.rainfall.mm	: int 3179 2252 3939 2763 3437 3058 2735 2952 2029 2061 ...		
\$ income.per.capita.in.USD	: int 2067 3106 3495 2874 22332 2102 6112 7465 3836 3219 ...		
\$ Life.expectancy.in.years	: num 73.1 71.8 73.5 70.6 76 64.7 NA 76.3 70 74.8 ...		
species	412 obs. of 8 variables		
\$ common_name	: chr "Christmas Island Pipistrelle" "Jamaican greater funnel-eared bat" ...		
\$ species_group	: chr "bats" "bats" "bats" "bats" ...		
\$ latin_name	: chr "Pipistrellus murrayi" "Natalus jamaicensis" "Natalus primus" "Cole...		
\$ est_pop	: num 0 50 100 100 160 250 250 300 300 500 ...		
\$ IUCN_abbr	: chr "EX" "CR" "CR" "CR" ...		
\$ IUCN_category	: chr "Extinct" "Critically Endangered" "Critically Endangered" "Critical...		
\$ Trend	: chr "Steady" "Decrease" "Decrease" "Decrease" ...		
\$ domesticated_YN	: logi FALSE FALSE FALSE FALSE FALSE FALSE ...		
values			
n	1		
N	15		
name	chr [1:3] "hello" "there" "world"		
test_average	84.8		
test_scores	num [1:5] 90 72 85 93 84		

# Changing Data Types

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```
# changing from integer to numeric
```

```
countries$avg_elev_m <- as.numeric(countries$avg_elev_m)
```

```
# turning a bunch of variables into factors (categories)
```

```
countries$Region <- as.factor(countries$Region)
```

```
countries$Income.Classification <- as.factor(countries$Income.Classification)
```

```
# view newly organized data set
```

```
summary(countries)
```

```
levels(countries$Region)
```

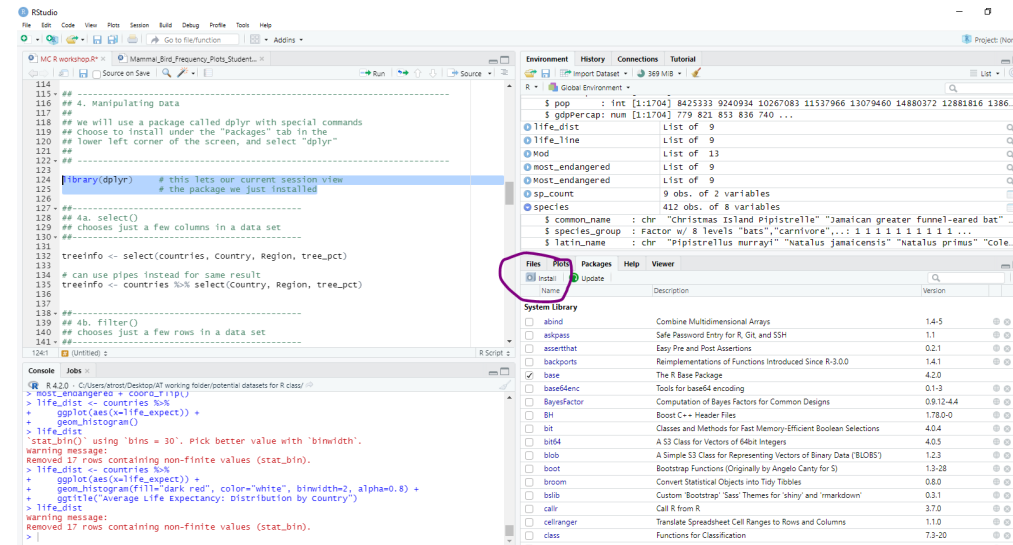
# Exercise 1

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1. How many columns are in the endangered species data set? How many rows?
2. How could you select just the first 3 columns of the countries data set?
3. How could you select just the latin names for the species data set?
4. What kind of variable is the countries data set?
5. What kind of variable is cropland\_pct in the countries data set?
6. Use `as.factor()` to change the species data set. With 3 separate lines of code, turn `species_group`, `IUCN_abbr`, and `IUCN_category` into factors. Summarize the species data set when you are through.

# Manipulating Data: Install dplyr package

1. Select “Install” under Packages in R studio. You only have to do this once for a specific computer.



2. Run the library command so that R knows to look for this package in a particular session.

```
library(dplyr)
```

```
# this lets our current session view
```

```
# the package we just installed
```

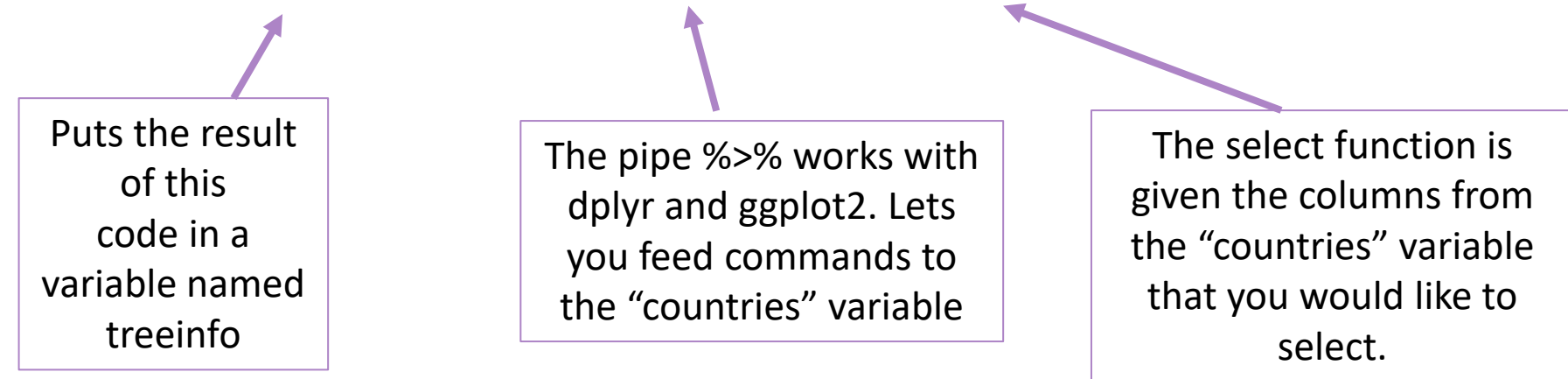
# Manipulating Data: Select

---

```
treeinfo <- select(countries, Country, Region, tree_pct)
```

# can use pipes instead for same result

```
treeinfo <- countries %>% select(Country, Region, tree_pct)
```



Puts the result of this code in a variable named treeinfo

The pipe %>% works with dplyr and ggplot2. Lets you feed commands to the "countries" variable

The select function is given the columns from the "countries" variable that you would like to select.

# Manipulating Data: Filter

# can filter on multiple categories

# just continue long commands on a new line

```
whales_dolphins_endangered <- species %>%  
  filter(species_group == "cetacean", IUCN_abbr == c("CR", "EN"))
```

# can also combine with select to get certain columns

```
whales_dolphins_endangered <- species %>%  
  filter(species_group == "cetacean", IUCN_abbr == c("CR", "EN")) %>%  
  select(common_name, est_pop, IUCN_category, Trend)
```

Using the “c” function inside the filter function to filter for more than one value. Two equals signs lets R know that you’re not trying to change the values of species\_group and IUCN\_abbr, you just want to find places where the values already match

A second pipe lets you perform a second operation (in sequence) on the “species” data

# Manipulating Data: group\_by, summarize, arrange

---

```
species %>% group_by(IUCN_category) %>%  
  summarize(n())
```

Summarize with “n” means you are counting the number of species in each category. The pipe %>% lets you perform two functions, group\_by and summarize

```
countries %>% group_by(Income_class) %>%  
  summarize(median(income_percap, na.rm=TRUE))
```

In this example summarize by median income (remove NAs)

```
countries %>% group_by(Income_class) %>%  
  summarize(med_income = median(income_percap, na.rm=TRUE)) %>%  
  arrange(med_income)
```

Give median income a variable name so you can reference it in the next line and arrange your results

# Exercise 2

---

1. Filter the species data to only include species with an estimated population of less than 10,000. Select just the species\_name, the species\_group, and the est\_pop .
2. Group the country data by region, then summarize using the median rainfall in mm. Arrange from largest to smallest, using:

```
arrange(desc(med_rainfall))
```



# Manipulating Data: Mutate

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The backwards arrow feeds the new (mutated) column back in to the original countries data set, making an additional column (variable)

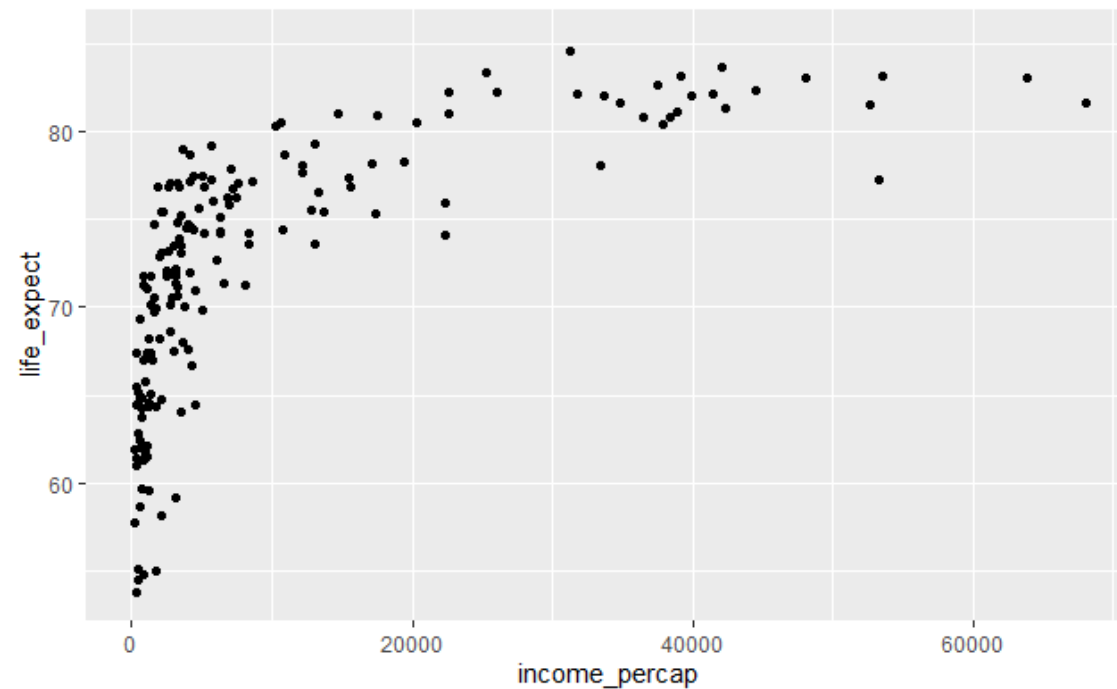
```
countries <- countries %>%  
  mutate(rainfall_in = rainfall_mm * .0393701)
```

Mutate creates a new data column (variable) based on existing information. Here we are creating a new column that converts rainfall in mm to inches.

# Visualizing Data: Scatterplot

```
ggplot(countries, aes(x=income_percap, y=life_expect)) +  
  geom_point()
```

Geom\_point tells R you want a scatter plot. You can ask for multiple geoms at a time



# Visualizing Data: Fancier Scatterplot

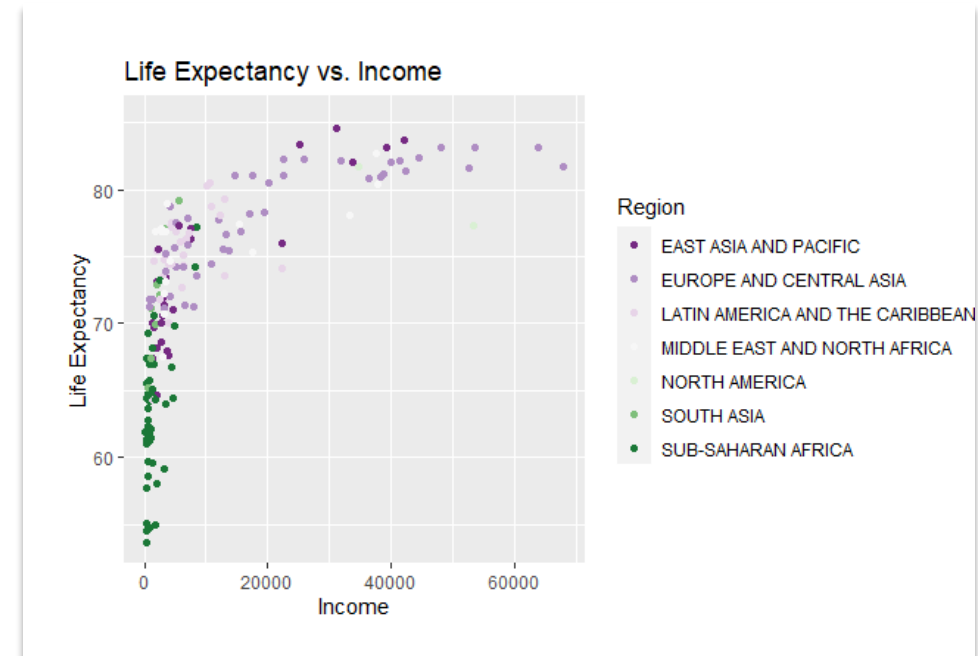
Gives the graph a name so you can call it up later (and add to it etc). Repeating “country\_scatter” at the end calls up the actual graphic.

Can list variable name then pipe it instead of including “countries” in the ggplot command

Assigning axis labels and a title here

```
country_scatter <- countries %>%  
  ggplot(aes(x=income_percap,y=life_expect)) +  
  labs(title = "Life Expectancy vs. Income",y="Life Expectancy", x="Income") +  
  geom_point(aes(col=Region)) +  
  scale_color_brewer(palette="PRGn")  
country_scatter
```

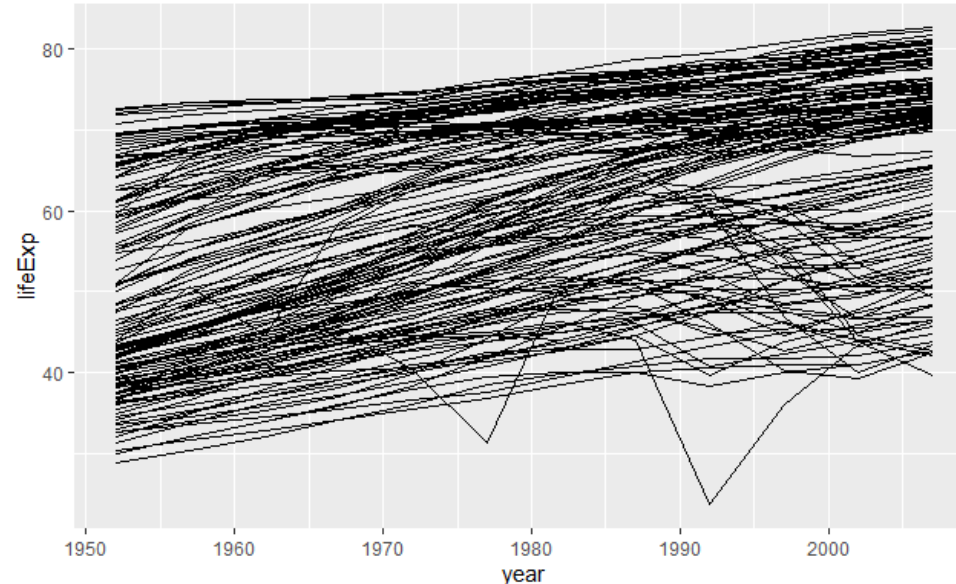
Col=Region tells R to assign color to points based on region. The scale\_color\_brewer is a special way to add built in color palettes using R.



# Visualizing Data: Line Graph

```
life_line <- gapminder %>%  
  ggplot(aes(x=year, y=lifeExp, by=country)) +  
  geom_line()  
life_line
```

geom\_line for a line graph



Tells ggplot to create a line for each country

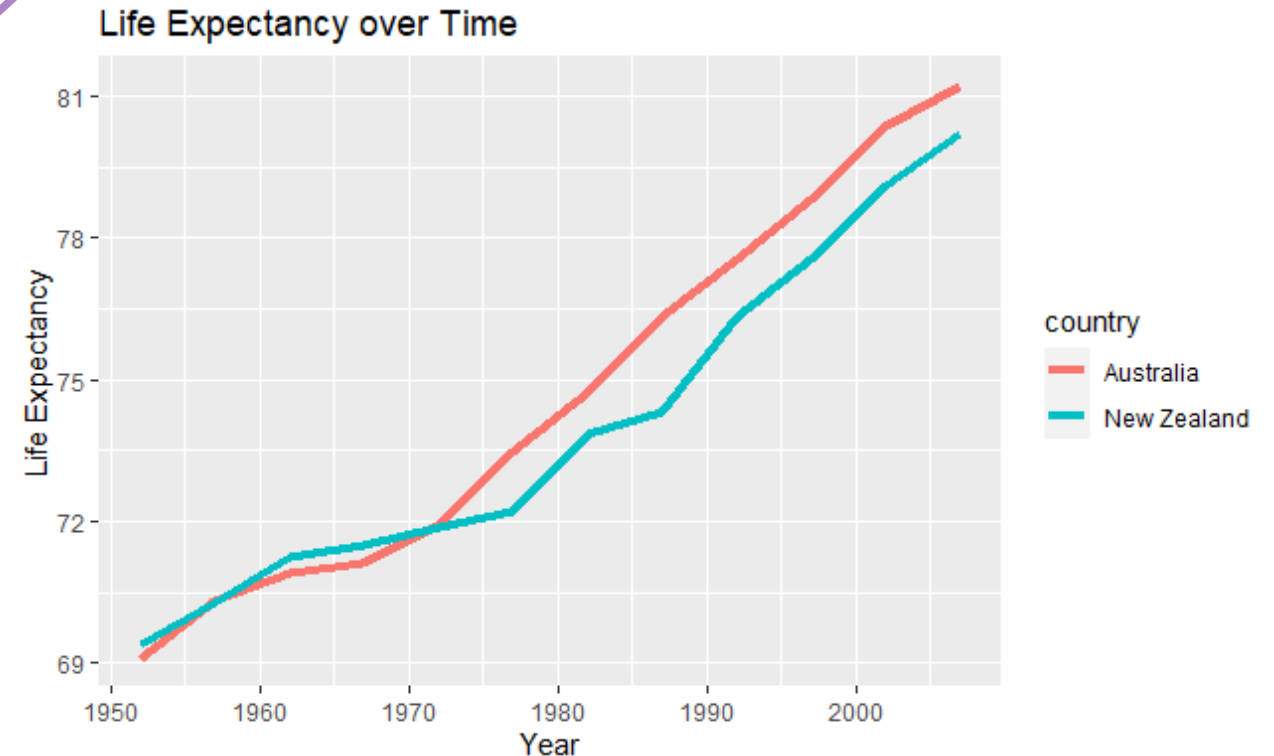
# Visualizing Data: Filtered Line Graph

```
life_line <- gapminder %>%  
  filter(continent == "Oceania") %>%  
  ggplot(aes(x=year, y=lifeExp, by=country)) +  
  labs(title = "Life Expectancy over Time", y="Life Expectancy", x="Year") +  
  geom_line(aes(col=country), size=1.5)  
life_line
```

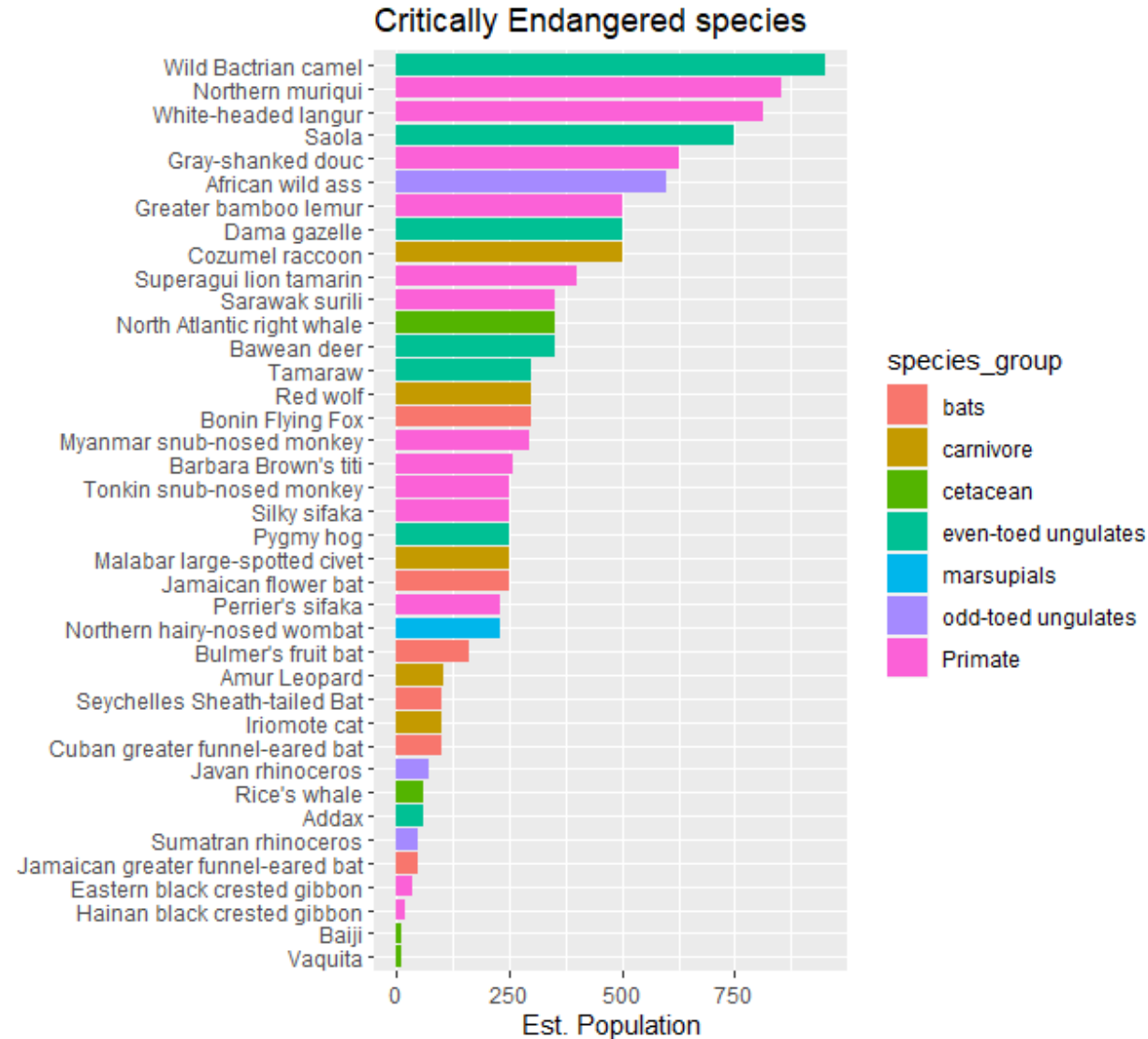
Telling R to only include countries in Oceania

Assigning axis labels and a title here

Col= country tells R to assign color to points based on country. The size=1.5 makes the lines a bit thicker



# Visualizing Data: Bar Graph



```
most_endangered <- species %>%  
  arrange(est_pop) %>%  
  filter(IUCN_abbr == "CR", est_pop < 1000) %>%  
  ggplot(aes(x=reorder(common_name, est_pop), y=est_pop)) +  
  labs(title = "Critically Endangered species", y="Est. Population", x=NULL) +  
  geom_col(aes(fill=species_group))  
most_endangered + coord_flip()
```

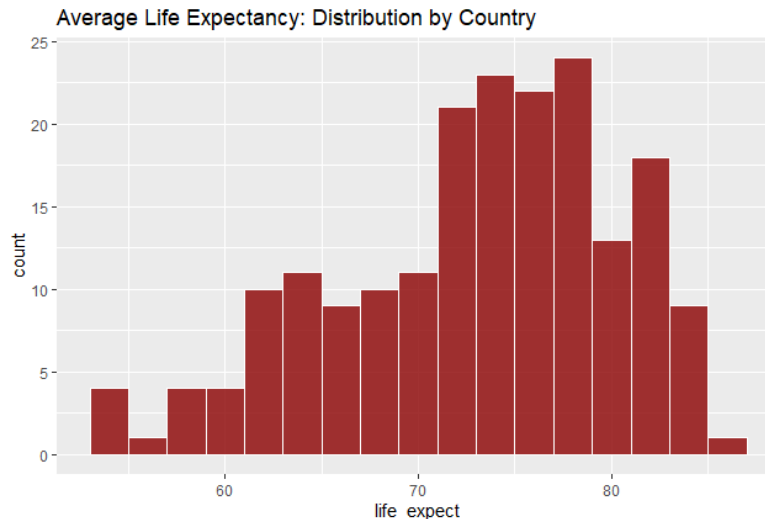
# Visualizing Data: Histogram

```
life_dist <- countries %>%  
  ggplot(aes(x=life_expect)) +  
  geom_histogram(fill="dark red", color="white", binwidth=2, alpha=0.8) +  
  ggtitle("Average Life Expectancy: Distribution by Country")  
life_dist
```

Creates a new bar for each  
group of 2 years

Makes bars 80% opaque

Makes the borders of the  
bars white



# Exercise 3

---

1. Filter the countries data set to include only data from South Asia. With this subset, create a bar chart that reports average rainfall in mm for each country.
2. Create a scatter plot with the countries data set that shows the relationship between tree cover and rainfall. Color the dots by region, label the x & y axes, and give the chart a title.



# Thank You

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QUESTIONS? PLEASE CONTACT ME AT  
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