



filter()

1
4
5
8

# Data Handling Simplified: Discover the Power of R



Instructor: Justin Barker

Maps, Data and Government Information Center

madgichelp@trentu.ca





**Collections Support** 



**Data Visualization Lab** 



Advanced computers, spatial, statistical and visualization software

Bata Library 402

GIS, Statistical, and Data Visualization Support



ESRI / ArcGIS





Research Data Management and Archiving



Research Collaborations



Regional
Environmental
History Atlas
Project



Workshops and teaching support



Bata Library 415

8:30 am to 4:30 pm, Monday to Friday



https://www.trentu.ca/library/madgic



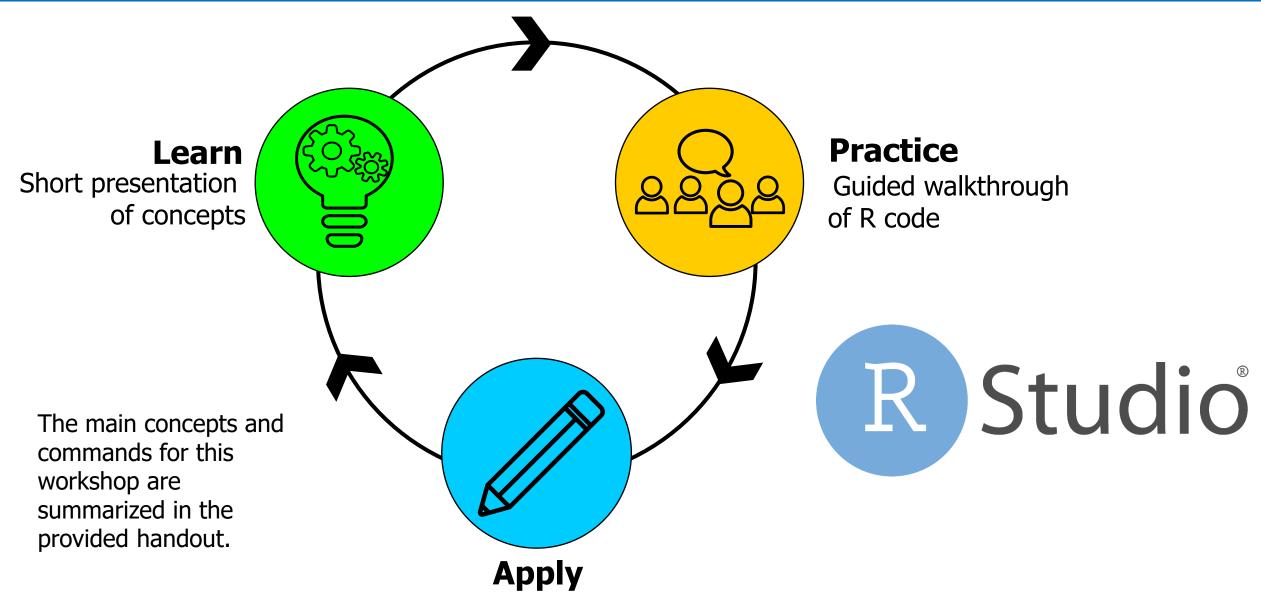
# Workshop content

All content for this workshop was e-mailed to you this morning: Subject: **R Data Handling Simplified files** 

- Slides
- Cheat sheet
- Script
- Data for workshop



# Workshop structure



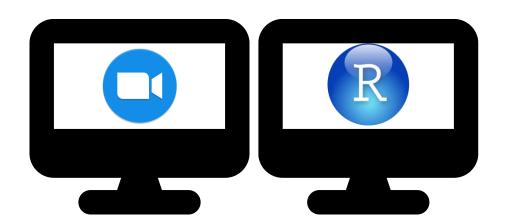
A multiple-choice poll to test your knowledge

# Following along

We'll go through the presentation and R script step by step.

Feel free to follow along using your own RStudio session, which works best if you have two screens.

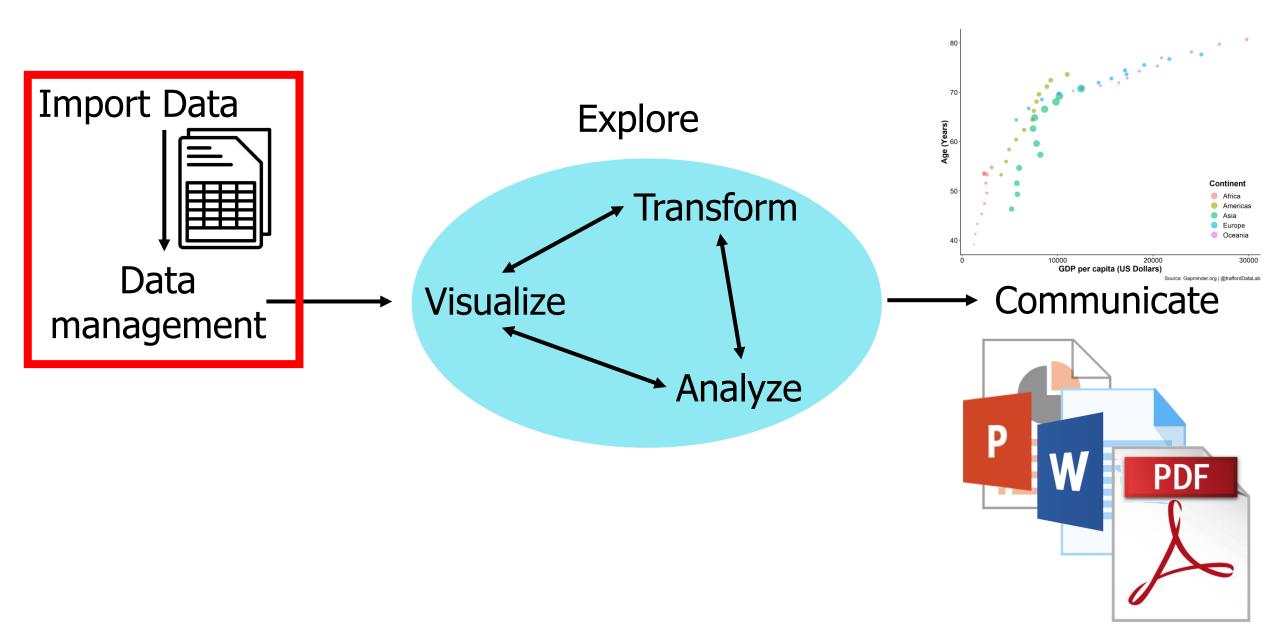
If not, you can easily replicate the script on any computer for future reference.





# R workflow

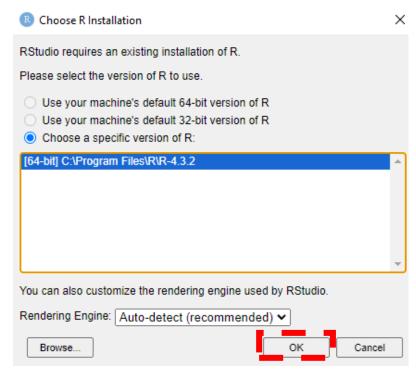






# Please open RStudio with the appropriate script for your computer: **DataHandlingR\_Windows.r** or **DataHandlingR\_Mac.r**

If this is your first time opening RStudio, you may receive the two pop-ups:



Enable Automated Crash Reporting

May we upload crash reports to RStudio automatically?

Crash reports don't include any personal information, except for IP addresses which are used to determine how many users are affected by each crash.

Crash reporting can be disabled at any time under the Global Options.

Accept the default R Installation

Allow R to upload crash reports



# Review of R Fundamentals

- Run code in the source pane (text editor) with Ctrl + Enter or the button
- Any text to the left of at least one # is a comment. Comments are non-executable text. Use them to annotate the code
  - Four or five # provide headers or sub-headers in the script, respectively.
  - Use Ctrl + Shift + C to quickly (un)comment lines
- Create objects in R with the assignment operator, <-</li>

```
> Spp_Ptbo_2021 <- 81
> Spp_Ptbo_2021
[1] 81
```

- Text data should always be input between ' or ''.
   Make sure text values are green!
  - This includes file paths

```
> read.csv("file.csv")
```

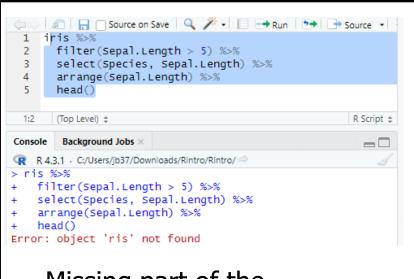


# Caution: Highlighting to run

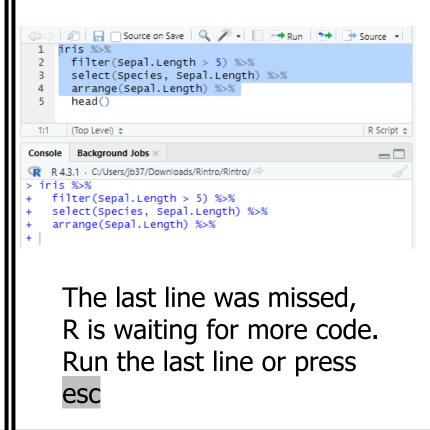


You can highlight part of a line, one line, or multiple lines.

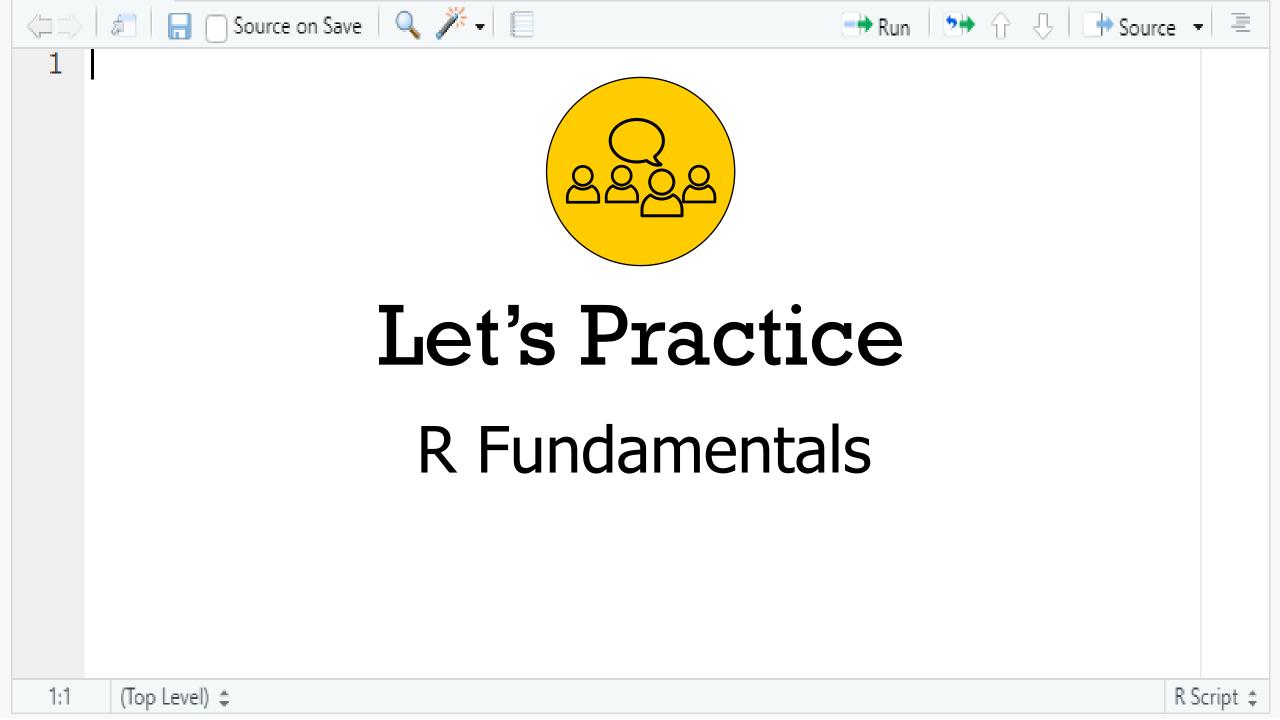
Be careful about what you are highlighting. Highlighting incomplete code may return an error or cause further issues.



Missing part of the beginning returns an error saying the data frame is not found



```
iris %>%
     filter(Sepal.Length > 5) %>%
     select(Species, Sepal.Length) %>%
     arrange(Sepal.Length) %>%
                                              R Script :
     (Top Level) $
Console Background Jobs
                                               =\Box
R 4.3.1 . C:/Users/jb37/Downloads/Rintro/Rintro/ 
  filter(Sepal.Length > 5) %>%
   select(Species, Sepal.Length) %>%
   arrange(Sepal.Length) %>%
  head()
 Species Sepal.Length
 setosa
 setosa
                5.1
 setosa
                5.1
 setosa
 setosa
                5.1
 setosa
                5.1
   All lines highlighted were
   run in their entirety
```







The Christmas Bird Count is a yearly census of birds performed by volunteers, indicating species counts per region.

Location

Year

Common Name

Count

Flags

Region where the count was conducted

Year when the count was conducted

Common name of the species being reported

Total number of observed individuals for the region and species.

Regional Editor flags (High count, low count, unusual count)



# **Importing Data**

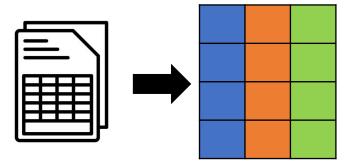


Most R work starts with bringing your data in.

R can read data from all common file types:

- Text (.txt)
- Comma separated values (.csv)

Base R



- Excel worksheet (.xlsx)
- SPSS dataset (.por)
- SAS data (.xpt)
- Stata (.dta)
- Raster (.tiff, .grd, .bil, .asc, .sdat, .rst, .nc, .envi, .img)
- Shapefiles (.shp)
- JSON (.json)
- SQL database queries

Requires additional packages



# **Importing Data**



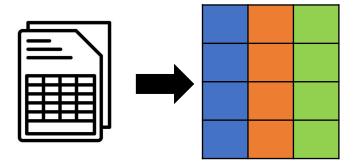
Most R work starts with bringing your data in.

R can read data from all common file types:

- Text (.txt)
- Comma separated values (.csv)



- SPSS dataset (.por)
- SAS data (.xpt)
- Stata (.dta)
- Raster (.tiff, .grd, .bil, .asc, .sdat, .rst, .nc, .envi, .img)
- Shapefiles (.shp)
- JSON (.json)
- SQL database queries



See readr and forgein packages functions to import other file types



# Formatting files before importing



# Raw Spreadsheet:

4	А	В	С	D	Е	F	G	Н
1	Name	100m	Long jump	Long jump	High.jump			
2	SEBRLE	NA	Disqualified	14.83	2.07		Mean	8.672938
3	CLAY	10.76	7.4	14.26	1.86		Median	9.2
4	KARPOV	11.02	7.3	14.77	2.04		Min	1.85

R interpretation:

	Name	X100m	Long.jump Long	.jump.1	High.jump	Х	X.1	X.2
1	SEBRLE	NA	"Disqualified"	14.83				8.672937
2	CLAY	10.76	"7.4"	14.26				9.200000
3	KARPOV	11.02	"7.3"	14.77	2.04	NA	Min	1.850000
4	BERNARD	11.02	"7.23"	14.25	1.92	NA	Мах	16.360000

Start column names with a letter

Remove blank columns and rows

Do not duplicate names or use spaces

Remove notes or other unnecessary data

Use NA or leave cells blank for missing values



# Importing functions



```
Text files (.txt):
```

file: file to be imported

header: file to be imported contains column names as the first line (TRUE) or not (FALSE)

\*Check your cheat sheet for different deliminator codes

# Comma Separated Values (.csv):

## **Excel Worksheets (.xlsx)**:

```
> library(openxlsx)
```

- > read.xlsx(xlsxFile = "file.xlsx", sheet = 1)
- > read.xlsx(xlsxFile = "file.xlsx", sheet = "Sheet1")

sheet: Sheet number (first sheet is 1) or name

# R environment data (.rda):

>load("file.rda")

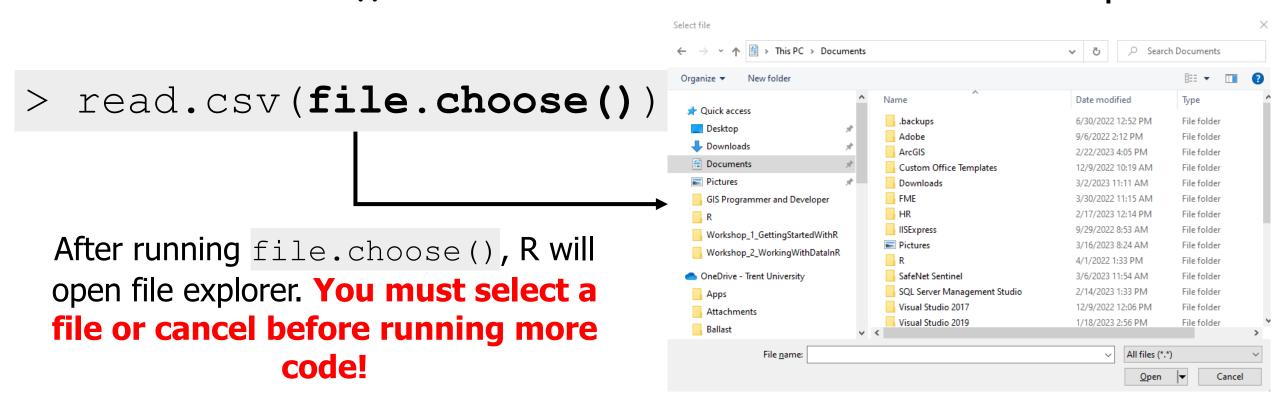
R environment data loads all objects saved in a previous R session



# Importing data: file.choose()



• file.choose(): Allows user to select file from file explorer





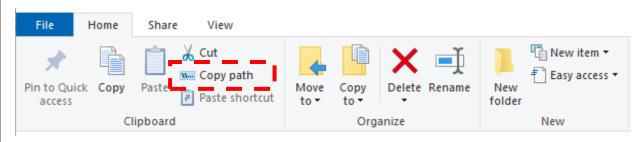
# Importing data: File path



- File path: Open a file by providing the full file path with forward slashes (/)
- > read.csv("E:/Data/Data.csv")

### Windows:

- 1. Find and select the file in File Explorer.
- 2. Click the Copy path button



- 3. Paste into the appropriate line in R
- 4. Use Find and Replace to change all "\" with "/"

### Mac:

- 1. Find and select the file in File Explorer.
- 2. Press CMD + Option + C
- 3. Paste into the appropriate line in R



# Importing data: Working directory



 Working directory: Open files within a folder and subfolder by setting the working directory

```
> setwd("E:/Data/")
```

> read.csv("Data.csv")

A working directory is the default location of any files that are imported from or exported to



# Inspect the data



Data should be inspected after importing to ensure it was imported correctly. This can be done through various functions:

Function	Returns
head()	Displays the first 6 rows or values
tail()	Displays the last 6 rows or values
str()	Displays the data frame structure (column names, data types, and first few values)
View()	Look at the whole data frame in a new window or tab
names()	Column names
rownames()	Row names
ncol()	Number of columns
nrow()	Number of rows
summary()	Basic statistics of each column

Note: These functions can be used with any data structure (vector, matrix, list, array)



# Inspect the data: str()



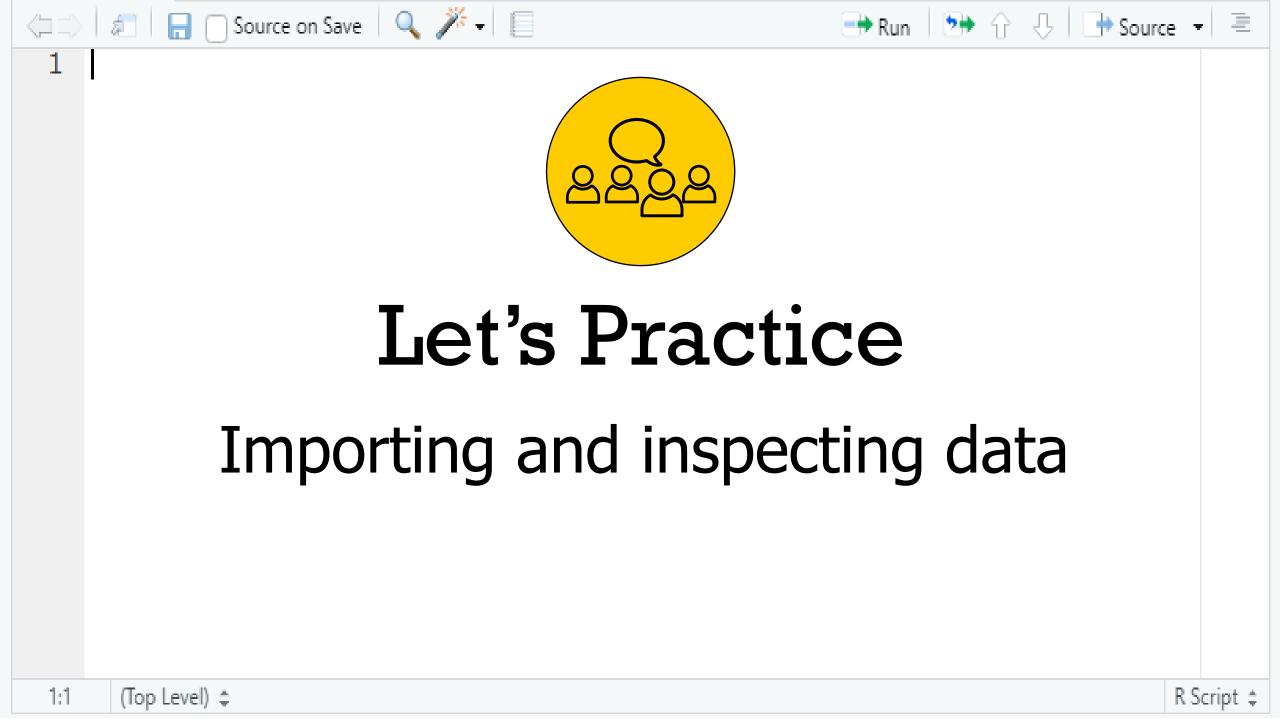
For a quick inspection, str() is recommended.

```
> str(Ptho 2020)
data.frame':
               104 obs. of 5 variables:
                      "Snow Goose" "Cackling Goose" "Canada Goose" "Wood Duck"
  Common_name
                 chr |
                      NA NA 1505 NA 12 1251 NA NA NA NA ...
                 int
  Count
  Flag
                 chr
                       NA NA NA NA ...
                      "Peterborough" "Peterborough" "Peterborough" "Peterborough"
                 chr
  Location_name:
                      2020 2020 2020 2020 2020 2020 2020 2020 2020 ...
  Year
```

- Column names
- Column data types
- Data structure
- Number of rows (observations)
- Number of columns (variables)
- Shows first few values per column

head() or tail() are good simple alternatives to show the first or last six rows.

glimpse() is synonymous with str()





# Challenge 1

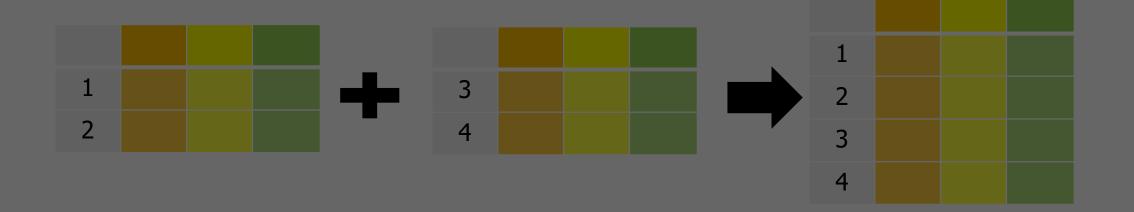


# Open and examine *ChristmasBirdCount\_2022\_SelectCities.txt,* Which of the following will successfully import it into R?

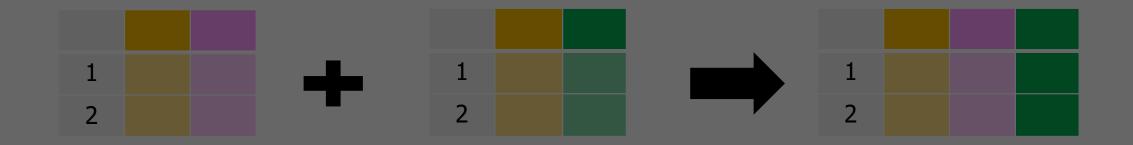
- a) read.table()
- b) read.csv()
- c) read.xlsx()
- d) None of the above

```
"Location" | "Year" | "Common Name" | "Count" | "Flags"
"Guelph" 2022 | "American Black Duck" 66 NA
"Guelph" 2022 | "American Crow" 424 NA
"Guelph" | 2022 | "American Kestrel" | 1 | NA
"Guelph" 2022 | "American Robin" 366 NA
"Guelph" 2022 | "American Wigeon" 2 NA
"Guelph" 2022 Bald Eagle 13 NA
"Guelph" | 2022 | "Belted Kingfisher" | 15 | "HC"
"Guelph" | 2022 | "Blue Jay" | 142 | NA
"Guelph" | 2022 | "Bohemian Waxwing" | 2 | NA
"Guelph" 2022 | "Brown Creeper" | 10 | NA
"Guelph" 2022 | "Bufflehead" 1 NA
"Guelph" 2022 | "Canada Goose" 4906 | NA
"Guelph"|2022|"Carolina Wren"|1|NA
"Guelph" 2022 | "Cedar Waxwing" | 175 | NA
"Guelph" 2022 | "Common Goldeneye" | 8 | NA
"Guelph" 2022 | "Common Merganser" | 57 | NA
```

# Data frame manipulation using dplyr





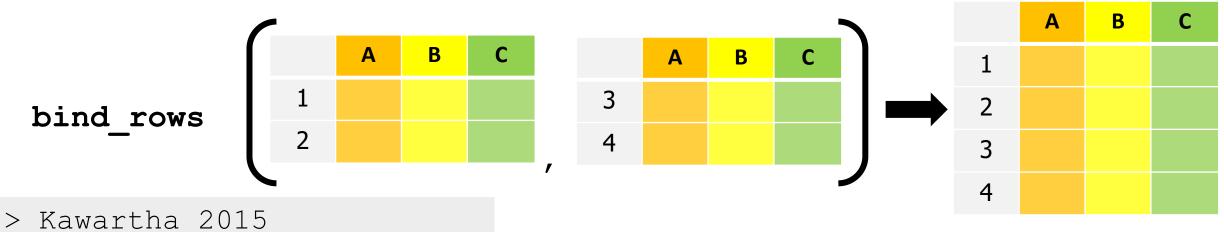




# Combine data frames by rows



## Combine rows with the same number of columns and names bind rows ():



```
Common Name Count Year
       Canada Goose 293 2015
2 American Black Duck 1 2015
      Mallard 382 2015
  Greater Scaup 7 2015
> Kawartha 2016
        Common Name Count Year
       Canada Goose 40 2016
     Trumpeter Swan 19 2016
3 American Black Duck 2 2016
           Mallard 26 2016
```

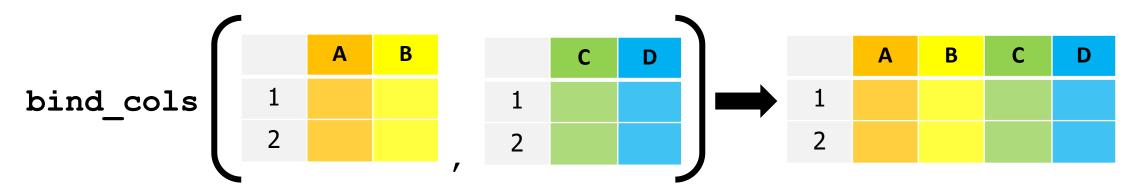
```
> Kawartha 2015 16 <- bind rows (Kawartha
+ 2015, Kawartha 2016)
> head(Kawartha 2015 16)
        Common Name Count Year
       Canada Goose 293 2015
2 American Black Duck 1 2015
           Mallard 382 2015
 Greater Scaup 7 2015
     Canada Goose 40 2016
     Trumpeter Swan 19 2016
```



# Combine data frames by column



# Combine columns with the same number of rows with bind\_cols():

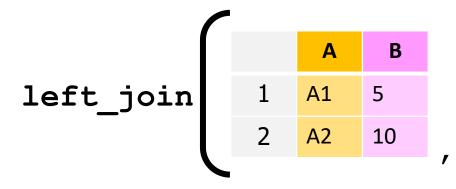




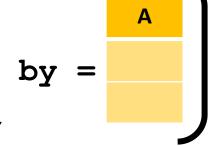
# Combine data frames by value



## Combine columns with the same values with left join():



	Α	С
1	A2	8
2	A1	7



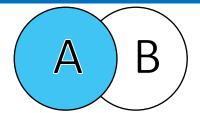
		Α	В	С
<b></b>	1	A1	5	7
	2	A2	10	8

```
> head(Kawartha 2015)
              Name Y2015
       Canada Goose 293
2 American Black Duck 1
      Mallard 382
      Greater Scaup
> head (Kawartha 2016)
              Name Y2016
       Canada Goose
            Mallard 26
      Greater Scaup
4 American Black Duck
```

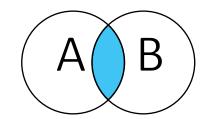
```
> Kawartha 15 16 <- left join(</pre>
+ Kawartha 2015, Kawartha 2016,
+ by = "Name")
> head(Kawartha 2015 16)
             Name Y2015 Y2016
   Canada Goose 293 40
2 American Black Duck 1 2
      Mallard 382 26
  Greater Scaup
```

# **Joins**

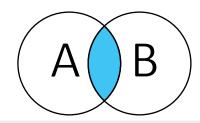




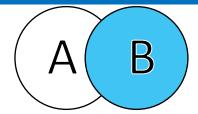
Join matching rows from B to A



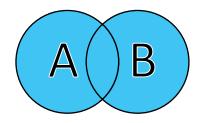
Join all matching rows in A and B, including duplicates



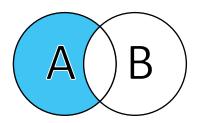
All rows in A that have a match in B, without duplicates



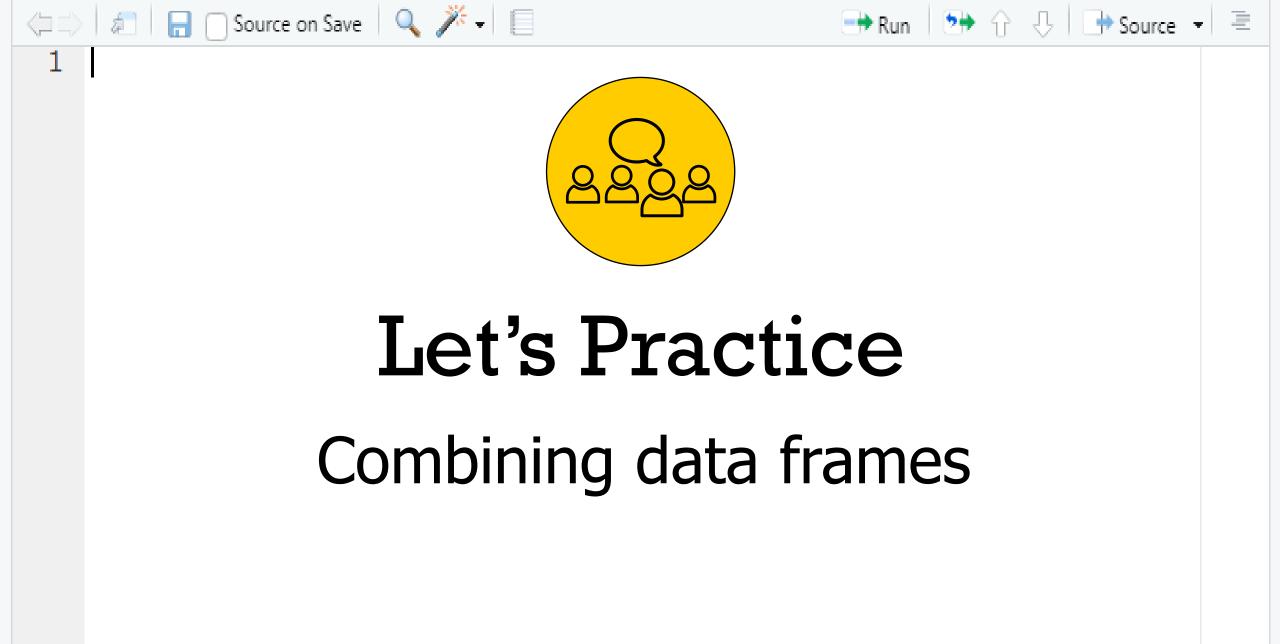
Join matching rows from A to B



Join data, retain all values, all rows



All rows in A that do not have a match in B





# Challenge 2



# Given the AVONET data set, how would one add AVONET to

CBC data joined?

```
> str(CBC_data_joined)
                                             > str(AVONET)
'data.frame': 3096 obs. of 7 variables:
                                             'data.frame':
                                                          9993 obs. of 5 variables:
                                              $ Scientific_Name: chr "Accipiter albogularis" "Accip
$ Location : chr "Guelph" "Guelph"
                                                            : num 10.6 8.8 9.2 8.9 8.7 6.6 8.3 8
                                              $ Beak.Width
                           2020 2020 2020 202
$ Year
       : num
                                              $ Beak.Depth : num 14.7 11.6 13.5 11.9 11.1 12 10.
$ Common_Name : chr "American Black Du
                                              $ Habitat : chr "Forest" "Shrubland" "Woodland"
ald Eagle" ...
                                              $ Trophic.Level : chr "Carnivore" "Carnivore" "Carniv
          : num 48 499 239 5 10 ..
$ Count
$ Flags : chr NA NA NA NA ...
$ Scientific_Name : chr "Anas rubripes" "C
us" "Haliaeetus leucocephalus" ...
 $ Num_Species_Reported: int 71 71 71 71 71
```

- a) bind rows (CBC Ptbo wScinames, AVONET)
- b) bind\_cols(CBC\_Ptbo\_wScinames, AVONET)
- c) left\_join(CBC\_data\_joined, AVONET, by = "Scientific\_name")

# Break time

5-10 minutes

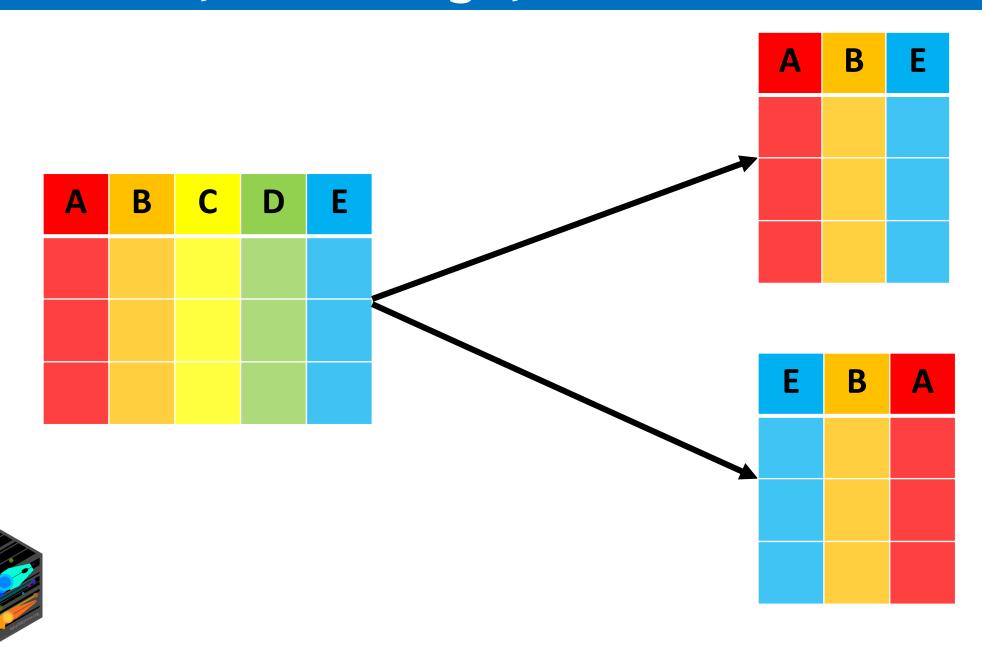


# Data frame manipulation using dplyr



# Subset, re-arrange, and remove columns





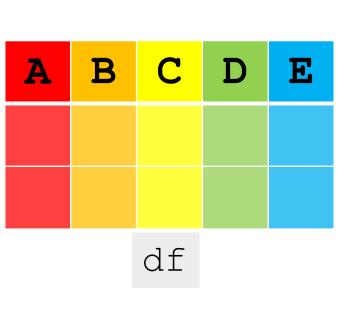




# Subset columns



Columns can be subset by name or position using select():

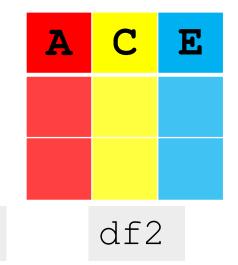


# By name:

> df2 <- select(df, A, C, E)

# By position:

> df2 <- select(df, 1, 3, 5)

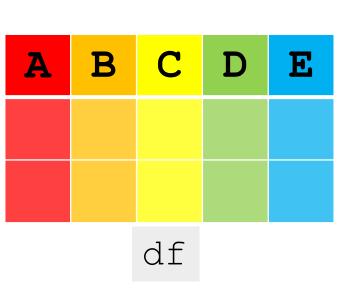




# Re-arrange columns



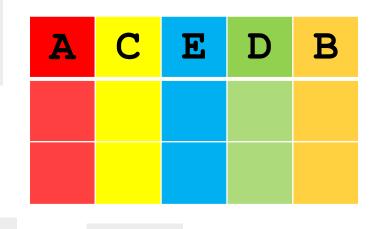
Changing column order in select() will return the order provided:



# By name:

```
> df2 <- select(df, A,
+ C, E, D, B)</pre>
```

# By position:



df2



#### Remove columns



Columns can be removed by placing a minus sign, –, before the column name or position in select():

## By name:

A B C D E

df

> df2 <- select(df, -D, -E)</pre>

By position:

> df2 <- select(df, -4, -5)</pre>

A B C

df2



#### **Selection helpers**



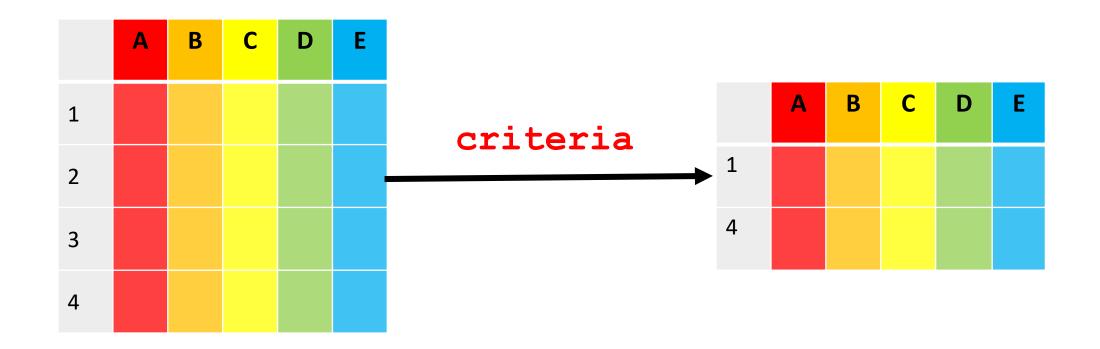
Columns can be selected based on a common pattern or criteria:

- everything () all columns, unless otherwise selected
- last col() select the last column, possibly with an offset
- starts\_with("A") columns that start with a specified pattern
- ends\_with("s") columns that end with a specified pattern
- contains ("a") columns that contain a specified pattern
- •: select sequential columns (e.g. Col\_A:Col\_J)



## **Subset rows**











## Select rows by value: filter()



Rows may be isolated based on a criteria

Different data types require specific criteria.

All common data types can be filtered based on a few common operators:

Description	Operator
Equals	==
Not equal to	! =
Missing values	is.na()
Non-missing values	!is.na()
Any of	X %in% c()
Not any of	!X %in% c()
	•

X represents a single column name



#### Select rows by numeric value



#### filter(data frame, col <criteria>)

#### Numeric filters include:

Description	<b>Operator</b>
Greater than	>
Less than	<
Greater than or equal to	>=
Less than or equal to	<=
Between two numbers	between(X, left, right)

```
X represents a single column name
```

```
>head(TO 20, 3)
     Common Name Count Flag
   Canada Goose 1229 <NA>
 Mute Swan 136 <NA>
3 Trumpeter Swan 102 <NA>
> filter(TO 20, Count <= 10)
      Common Name Count Flag
        Wood Duck 7 <NA>
 Northern Pintail 1 <NA>
    Canvasback 3 <NA>
4 Ring-necked Duck 1 <NA>
     Merganser sp. 8 <NA>
         Duck sp. 3 < NA >
      Wild Turkey 3 <NA>
                    3 <NA>
      Common Loon
```



#### Select rows by character value



Filtering character values is made easier through the stringr package:

Description	Operator
Contains	str_detect(X,"s")
Begins with	str_detect(X, "^s")
Ends with	str_detect(X, "s\$")
Contains digits	str_detect(X, "\\d")

X represents a single column name "s" represents a string



#### Filter by multiple criteria: AND



Multiple criteria can be applied using the AND (&) or OR (|) operators:

- **AND**: All criteria must be met
- OR: One of the criteria must be met

```
filter(data frame, col <criteria>)
```

Common names that end in "d" AND have counts less than 1000



## Filter by multiple criteria: OR



Multiple criteria can be applied using the AND (&) or OR (I) operators

- AND: All criteria must be met
- **OR**: One of the criteria must be met

```
filter(data frame, col <criteria>)
```

```
> filter(TO 20, str detect(Common Name, "d$")
              Count < 1000
             Common Name Count Flag
               Mute Swan 136 <NA>
                                           Add a | with
          Trumpeter Swan 102 <NA>
                Mallard 2713 <NA>
                                           Shift + \
         Northern Pintail 1 <NA>
              Canvasback 3 <NA>
                Redhead 825 <NA>
```

Common names that end in "d" OR have counts less than 1000



## Sort rows: arrange ()



Sort rows by one or more columns in ascending order (A - Z, 0 - 100):

#### Sort by a single column

```
> arrange(TO 20, Count)
               Common Name Count Flag
          Northern Pintail
                             1 < NA >
          Ring-necked Duck 1 <NA>
3
          Red-necked Grebe 1 <NA>
                Bald Eagle 1 <NA>
       Red-shouldered Hawk
                             1 < NA >
             American Coot
                             1 <NA>
          Purple Sandpiper
                              1 <NA>
         American Woodcock
                               US
```

#### Sort by multiple columns

```
> arrange(TO 20, Count,
            Common Name)
               Common Name Count Flag
             American Coot
                             1 <NA>
         American Woodcock
                             1 US
                Bald Eagle
                             1 <NA>
     Downy/Hairy Woodpecker
                             1 < NA >
                             1 <NA>
               Fox Sparrow
             Glaucous Gull
                             1 <NA>
          Northern Pintail
                             1 <NA>
      Northern Saw-whet Owl
                             1 <NA>
```

Sorts by Count THEN Common\_Name



#### Sort rows in descending order



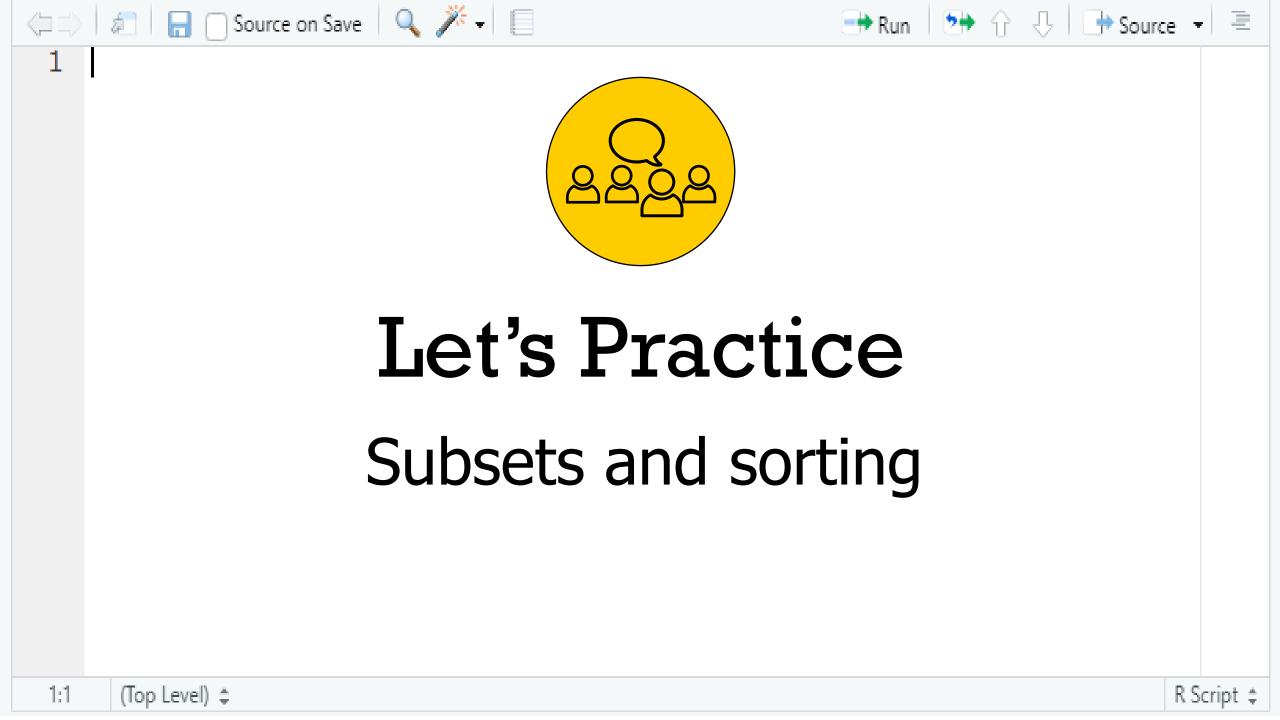
Sort rows by one or more columns in descending order (Z - A, 100 - 0) by using desc() around each desired column:

```
arrange(data frame, desc(<by>))
```

#### Sort by a single column

```
> arrange(TO 20, desc(Count))
                Common Name Count
                    Mallard
                             2713
2
          European Starling 2703
3
4
                             2370
           Long-tailed Duck
              House Sparrow 2012
5
                             1475
     Black-capped Chickadee
              Greater Scaup 1365
6
           Ring-billed Gull
                             1353
8
                             1229
               Canada Goose
```

#### Sort by multiple columns





## Challenge 3



Match the functions with their proper application

select()

Sort rows

filter()

Subset, re-arrange, and remove columns

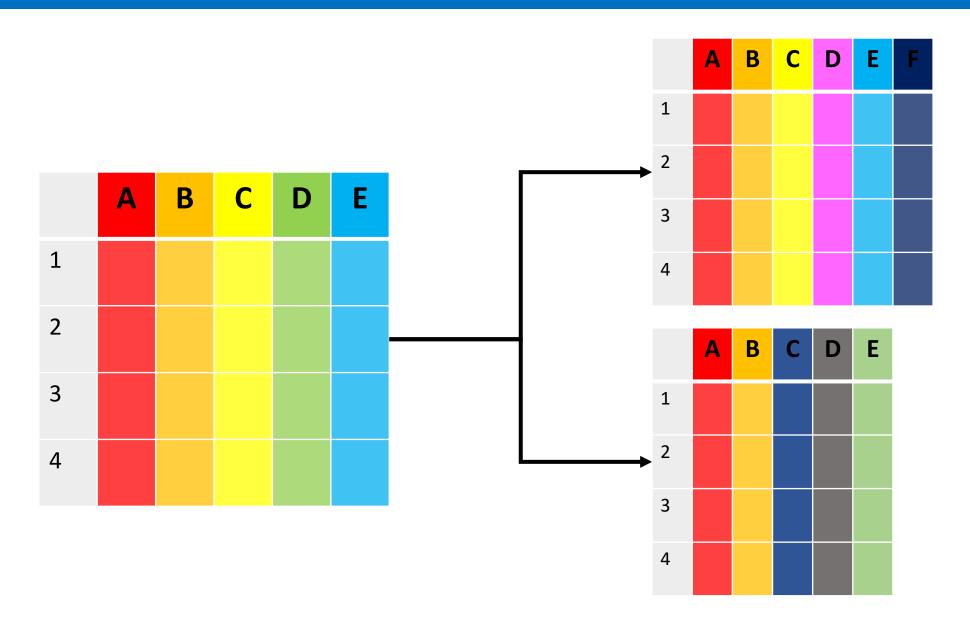
arrange()

Subset rows



#### Add or alter columns











## Creating new columns: mutate()



#### Create new columns from existing data:

```
mutate(data_frame, new_col = function(existing_col))
```

```
> head(CBC Barrie, 2)
      Common Name Y2019 Y2020
1 Accipiter sp. 1 1
2 American Black Duck 56 75
> mutate (CBC Barrie, Count Change = Y2020 - Y2019)
           Common Name Y2019 Y2020 Count Change
        Accipiter sp. 1 1
2 American Black Duck 56 75
```



#### Alter column values: mutate()



If a name of an existing column is given in mutate(), that column is altered:

```
mutate(data_frame, existing_col = function(existing_col))
```



#### Replace values



Use mutate() with ifelse() to change specified values:

```
mutate(data_frame, existing_col = function(existing_col))
```

```
> head(CBC Barrie, 5)
         Common Name Y2019 Y2020
       Accipiter sp.
  American Black Duck
                         75
   American Coot
                          NA
  American Crow
                          99
                    109
  American Robin
                     NA
                          NA
  mutate(CBC Barrie, Y2019 = ifelse(is.na(Y2019), 0, Y2019))
         Common Name Y2019 Y2020
       Accipiter sp.
   American Black Duck
                   56
                          75
    American Coot
                          NA
  American Crow
                    109
                          99
 American Robin
                          NA
```

If Y2019 is NA, then replace with 0, else return original Y2019 value

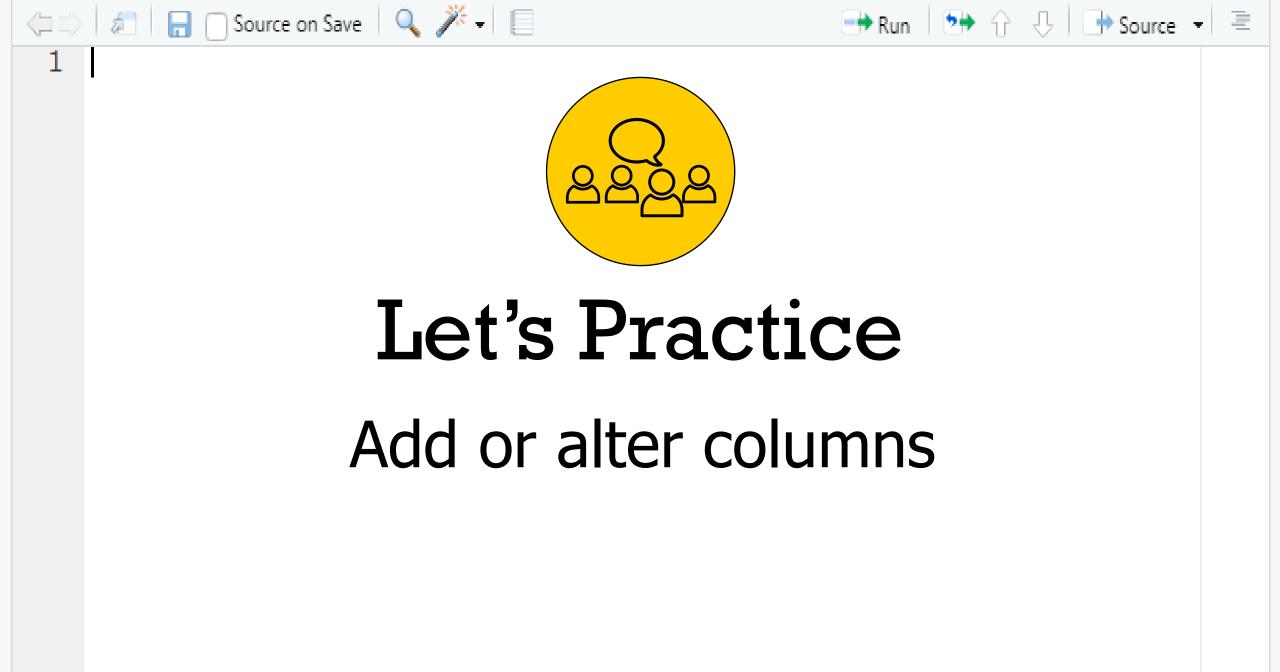


## Specify where the new column goes



When creating a new column with mutate() the column will be placed on the end (right) of the data frame. Use the arguments .before or .after to specify a different location.

```
> mutate(CBC Barrie, Total = Y2019 + Y2020, .before = Y2019)
        Common Name Total Y2019 Y2020
      Accipiter sp. 2 1 1
 American Black Duck 131 56 75
  American Coot 0 0 0
  mutate(CBC Barrie, Total = Y2019 + Y2020, .after = Common Name)
       Common Name Total Y2019 Y2020
      Accipiter sp. 2 1 1
 American Black Duck 131 56 75
  American Coot 0 0 0
```





## Challenge 4



Using mutate(), which code would create a new column, Count\_Sq, representing the Count value squared (Count^2) to the right of Count?

- a) > mutate(CBC\_data\_cleaned, Count\_Sq = Count^2)

# Streamline data manipulation: Piping Ceci n'est pas un pipe.



## Piping %>%



Conducting multiple functions on a data frame requires creating new data frames or overwriting the original.

Piping allows multiple functions to be computed in sequence from a single data set.

```
> CBC_Barrie_Dif <- CBC_data %>%
+ filter(Location = "Barrie") %>%
+ mutate(Difference = Y2019 - Y2020)
```



## Piping framework



# Previously applying functions followed the framework:

function(data, arguments)

```
> filter(CBC_data, Location = "Barrie")
```

# Piping rewrites this framework as:

data %>% function(arguments)

```
> CBC data %>% filter(Location = "Barrie")
```

#### Quickly add %>% with:

- Ctrl + Shift + M on Windows
- Cmd + Shift + M on Mac



## Writing pipelines



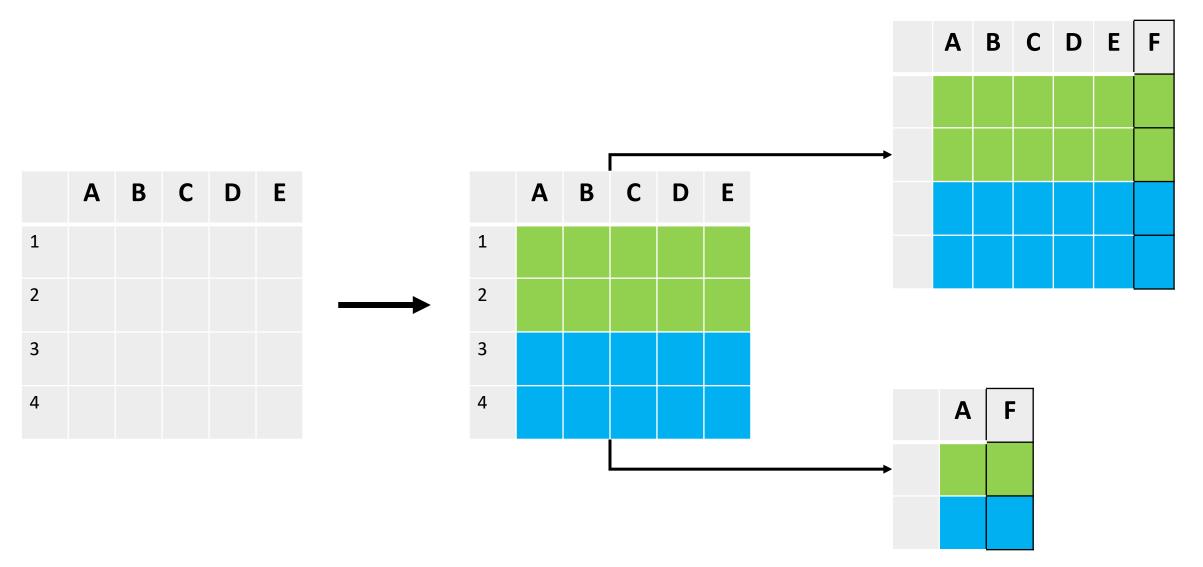
The pipe operator, \$>\$, is put at the END of each dataset or function. The output of each function is then applied to the next function.

Piping improves code readability, limits the need to create or overwrite objects, and reduces user commands.



## Aggregating data







## Aggregating data



Piping allows for quick calculations of values per group with group\_by() followed by either mutate() or summarise().

```
> data %>%
+ group_by(...) %>%
+ mutate(...)
→
```





#### Aggregating data: group by() %>% mutate()



```
> data %>%
+ group_by(...) %>%
+ mutate(...)
```

Computes new values by group for each row (observation) within the group.

```
> head(CBC Data, 3)
 Location Year Common Name Count
                                  Flag
1 Peterborough 2020 Cooper's Hawk 9
                                   <NA>
2 Peterborough 2017 Cooper's Hawk 12 HC
3 Peterborough 2019 Cooper's Hawk 7 <NA>
> CBC Data %>%
     group_by(Location, Year) %>%
     mutate(N Spp = n())
 Location Year Common Name Count Flag N Spp
1 Peterborough 2020 Cooper's Hawk 9
                                   <NA>
                                          54
2 Peterborough 2017 Cooper's Hawk 12 HC
                                          51
3 Peterborough 2019 Cooper's Hawk
                                 <NA>
                                          47
```



#### Aggregating data: group by() %>% summarise()



Computes a summary for each group and removes all unconsidered data and individual rows (observations).

```
> head(CBC Data, 3)
Location Year Common Name Count Flag
1 Peterborough 2020 Cooper's Hawk 9 <NA>
2 Peterborough 2017 Cooper's Hawk 12 HC
3 Peterborough 2019 Cooper's Hawk 7 <NA>
> CBC Data %>%
  group by (Location, Year) %>%
     summarise(N Spp = n())
  Location Year N Spp
1 Peterborough 2017 51
 Peterborough 2018 55
 Peterborough 2019 47
 Peterborough 2020
                  54
```





## Let's Practice

Aggregating data

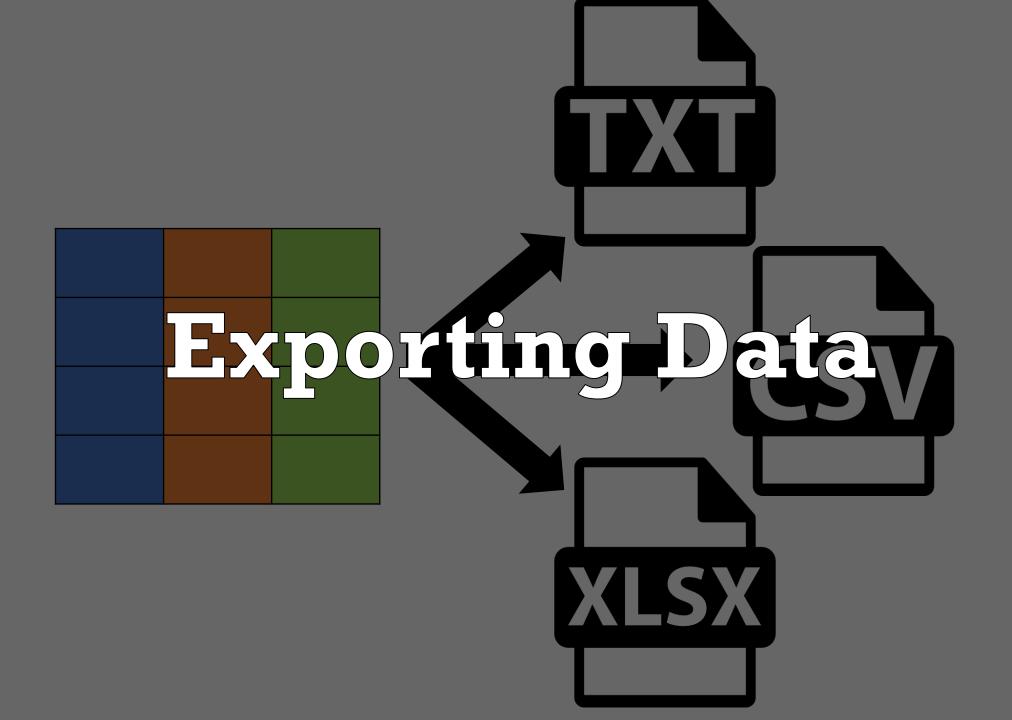


## Challenge 5



What use of group\_by() , mutate(), and/or summarise() would return
the following output:

```
CBC data cleaned %>%
                                   a)
                                              summarise(Avg BeakWidth = mean(Beak.Width))
                  Avg_BeakWidth
  Habitat
                           \langle db1 \rangle
  <chr>>
                                       CBC data cleaned %>%
1 Coastal
                            8.66
                                             mutate(Avg BeakWidth = mean(Beak.Width))
                            7.40
2 Forest
3 Grassland
                           11.2
4 Human Modified
                            8.45
                                       CBC data cleaned %>%
5 Marine
                           12.4
                                              group by (Habitat) %>%
                            9.95
6 Riverine
                                              summarise(Avg BeakWidth = mean(Beak.Width))
7 Shrubland
                            6.15
8 Wetland
                           14.5
                            7.31
9 Woodland
                                       CBC data cleaned %>%
                                              group by (Habitat) %>%
                                    d)
                                              mutate(Avg BeakWidth = mean(Beak.Width))
```





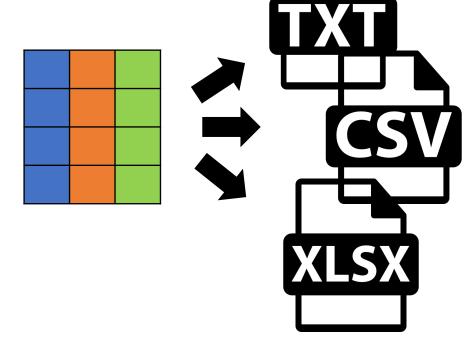
## **Export data**



Once data has been manipulated, it can be exported in all multiple formats (depending on the data).

R can read data from all common file types:

- Text (.txt)
- Comma separated values (.csv)
- Excel worksheet (.xlsx)
- SPSS dataset (.por)
- SAS Data (.xpt)
- Stata (.dta)
- Raster (.tiff, .grd, .bil, .asc, .sdat, .rst, .nc, .envi, .img)
- Shapefiles (.shp)
- JSON (.json)



See readr and forgein packages functions to export other file types



#### **Export functions**



#### **Text files** are exported with write.table()

```
> write.table(x = data, file = "file.txt", sep="\t",
+ row.names = TRUE, col.names = TRUE)
```



#### Comma separated values are exported with write.csv()

```
> write.csv(x = data, file = "file.csv", row.names = TRUE)
```



#### **Excel Worksheets (.xlsx)** are exported with write.xlsx()

- > library(openxlsx)
- > write.xlsx(x = data, file = "file.xlsx")

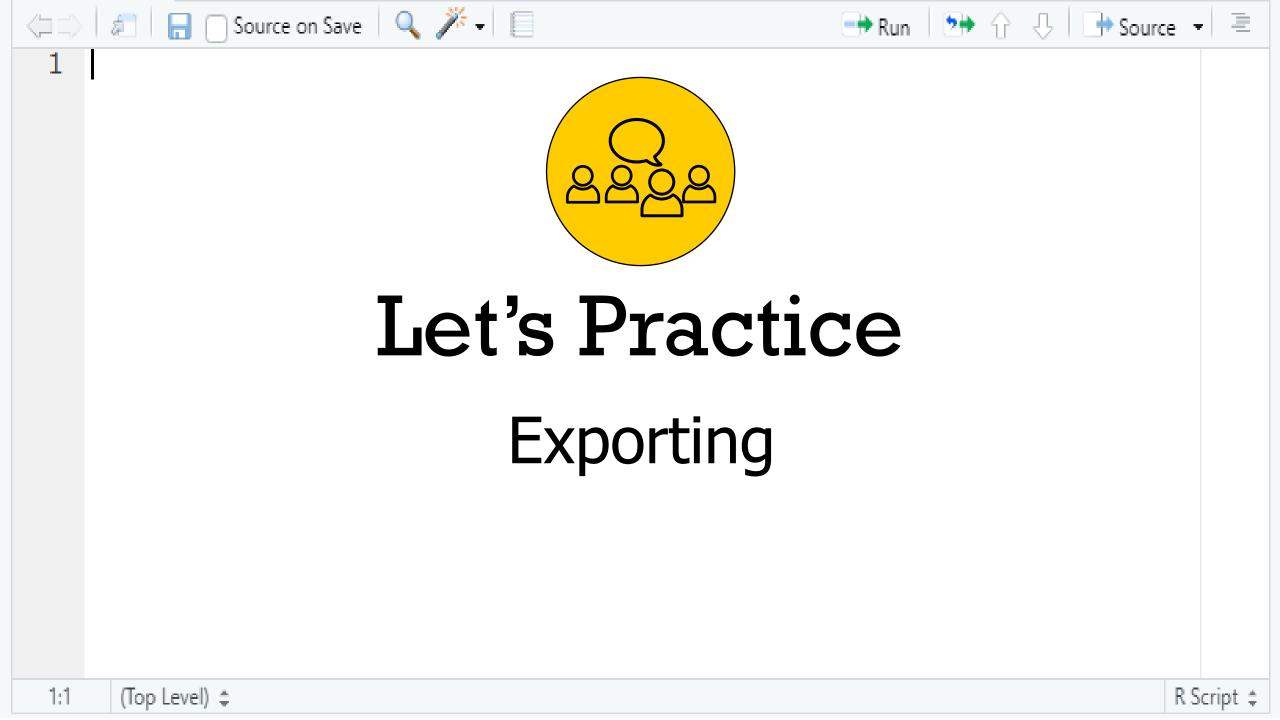


#### R environment data (.rda):

- > save(list = ls(), "file.rda")
- Export the entire R environment

> save(data1, data2, "file.rda")

Export the only objects data1 and data2





- Bata Library 415 8:30 am to 4:30 pm, Monday to Friday
- madgichelp@trentu.ca
- https://www.trentu.ca/library/madgic