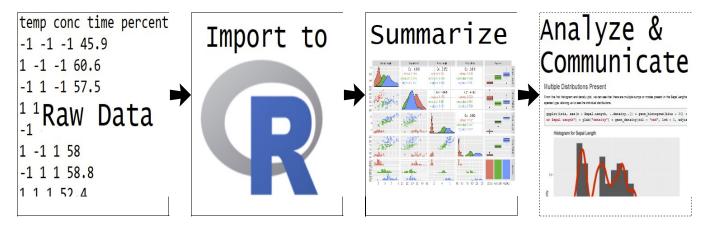
# NC STATE UNIVERSITY

# Importing Data: Basics & Delimited Data

#### What is this course about?

Basic use of R for reading, manipulating, and plotting data!



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Basic use of R for reading, manipulating, and plotting data!

- · read and write basic R programs
- · import well formatted data into R
- · do basic data manipulation in R
- · produce common numerical and graphical summaries in R
- · describe a use case of an analysis done in R

#### Where do we start?

- · Common raw data formats
- · Comma Separated Value (CSV) files
- · Asides: R projects and R packages
- · Read 'clean' delimited data
- · Excel, SAS, & SPSS data
- · Resources for JSON, XML, databases, and APIs

# **Importing Data**

#### How to read in data depends on raw/external data type!

- · Lecture focus: Delimited data
  - Delimiter Character (such as a , ) that separates data entries

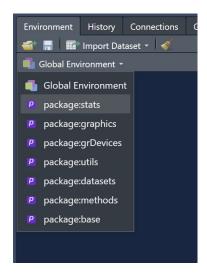
Treatment, Sex, Age, Duration, Pain	temp conc time percent	"color	" "spine	" "width 28.3	" "sate:	11" 3050	"weight"	"у"	2012>4>12>MIN>LAA>D.J. Reyburn
P,F,68,1,No		4	3	22.5	0	1550	0		2012>4>12>SD>ARI>Marty Foster
B,M,74,16,No	-1 -1 -1 45.9	2	1	26 24.8	9	2300 2100	1		2012>4>12>WSH>CIN>Mike Everitt
P,F,67,30,No	1 -1 -1 60.6	4	3	26	4	2600	1		2012>4>12>PHI>MIA>Jeff Nelson
		2	1	23.8	0	2100 2350	0		
P,M,66,26,Yes	-1 1 -1 57.5	4	2	24.7	0	1900	0		2012>4>12>CHC>MIL>Fieldin Culbreth
B,F,67,28,No		4	1	23.7	0 0	1950 2150	0 0		2012>4>12>LAD>PIT>Wally Bell
B,F,77,16,No	1 1 -1 58.6	4	3	24.3	0	2150	0		2012>4>12>TEX>SEA>Doug Eddings
	NAME OF THE PARTY	3	3	25.8	0	2650	0		
A,F,71,12,No	-1 -1 1 53.3	5	3	28.2 21	11 a	3050 1850	9		2012>4>12>COL>SF>Ron Kulpa
B,F,72,50,No	Service and Act appropriate	3	1	26	14	2300	1		2012>4>12>DET>TB>Mark Carlson
	1 -1 1 58	2	1	27.1	8	2950	1		2012>4>13>NYY>LAA>Mike DiMuro
B,F,76,9,Yes		3	3	25.2 29	1	2000 3000	1		
Λ M 71 17 Voc	-1 1 1 58.8	5	2	24 7	a	2200	a	ı	2012>4>13>COL>ART>Mark Wegner
Comma: usually .csv	Space: usually .txt or .dat	1		Tab:	usua	ally .t	sv or .tx	t	General: usually .txt or .dat

## Importing Delimited Data: Standard R Methods

- · When you open R a few packages are loaded
- · R package
  - Collection of functions/datasets/etc. in one place
  - Packages exist to do almost anything
  - List of CRAN approved packages on R's website
  - Plenty of other packages on places like GitHub

# Importing Delimited Data: Standard R Methods

· When you open R a few packages are loaded



• utils package has family of read. functions ready for use!

# Importing Delimited Data: Standard R Methods

#### Function and purpose:

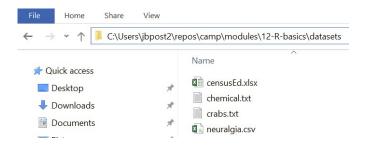
Type of Delimeter	Function
Comma	read.csv()
Semicolon ( , for decimal)	read.csv2()
Tab	read.delim()
White Space/General	<pre>read.table(sep = "")</pre>

· Each function requires a path to the file

- Let's read in the 'neuralgia.csv' file
- · How does R locate the file?

#### Path to File

- · Let's read in the 'neuralgia.csv' file
- · How does R locate the file?
  - Can give full path name
    - ex: C:/Users/jbpost2/repos/camp/modules/12-R-basics/datasets /neuralgia.csv
    - ex: C:\\Users\\jbpost2\\repos\\camp\\modules\\12-R-basics\\datasets \\neuralgia.csv



- · Let's read in the 'neuralgia.csv' file
- · Use full local path

head(neuralgiaData)

```
##
    Treatment Sex Age Duration Pain
            Р
                   68
                             1
## 1
                F
                                 No
## 2
            В
                M 74
                            16
                                No
## 3
            Ρ
                F 67
                            30
                               No
## 4
                M 66
                            26 Yes
## 5
               F 67
                            28
                                No
## 6
            В
               F 77
                            16
                                 No
```

- · Let's read in the 'neuralgia.csv' file
- · Using full local path not recommended!
  - Can't share code without changing path...

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- · Can change working directory
  - Folder where R 'looks' for files
  - Supply **relative** path

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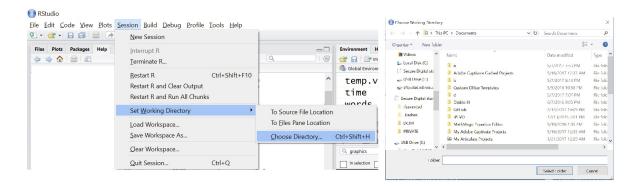
getwd()

## [1] "C:/Users/jbpost2/repos/camp/modules/12-R-basics/Module2\_ImportingData"

- · Can change working directory
  - Via code

```
setwd("C:/Users/jbpost2/repos/camp/modules/12-R-basics/datasets")
#or
setwd("C:\\Users\\jbpost2\\repos\\camp\\modules\\12-R-basics\\datasets")
```

- · Can change working directory
  - Via code
  - Via menus



- · Let's read in the 'neuralgia.csv' file
- Use relative path (.../ drops down a folder)

```
neuralgiaData <- read.csv("../datasets/neuralgia.csv")</pre>
```

- Working directory: ".../12-R-basics/Module2\_ImportingData"
- File location: ".../12-R-basics/datasets/neuralgia.csv
- As long others have the same folder structure, can share code with no path change needed!

- · Let's read in the 'neuralgia.csv' file
- · R can pull from URLs as well!

neuralgiaData <- read.csv("https://www4.stat.ncsu.edu/~online/datasets/neuralgia.csv")
head(neuralgiaData)</pre>

```
##
    Treatment Sex Age Duration Pain
## 1
             Р
                   68
                             1
                F
                                 No
## 2
            В
                M 74
                            16
                                 No
## 3
            Ρ
                            30
                F 67
                                No
                M 66
                            26 Yes
## 5
            В
                F 67
                            28
                                 No
                F 77
                            16
## 6
            В
                                No
```

```
read.csv() function
```

Tell R where the file lives via:

- · a full local path (not recommended)
- · a relative path
  - can set the working directory with setwd()
- pulling from URL

# Aside: RStudio Project

- · Often have many files associated with an analysis
- · With multiple analyses things get cluttered...

# Aside: RStudio Project

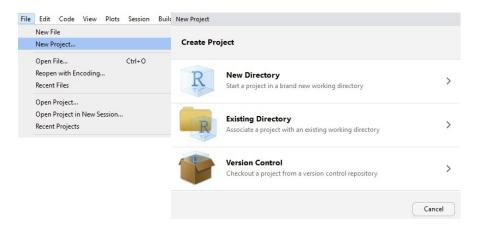
- · Often have many files associated with an analysis
- · With multiple analyses things get cluttered...
- · Want to associate different
  - environments
  - histories
  - working directories
  - source documents

#### with each analysis

· Can use "Project" feature in R Studio

## Aside: RStudio - Project

Easy to create! Use an existing folder or create one:



- · Easily switch between analyses!
- · Create one for today's lesson
- · Swap between projects using menu in top right

# **Reading Delimited Data**

- Functions from read. family work well
- · Concerns:
  - (formerly, prior to R 4.0) poor default function behavior
    - strings were read as factors

# **Reading Delimited Data**

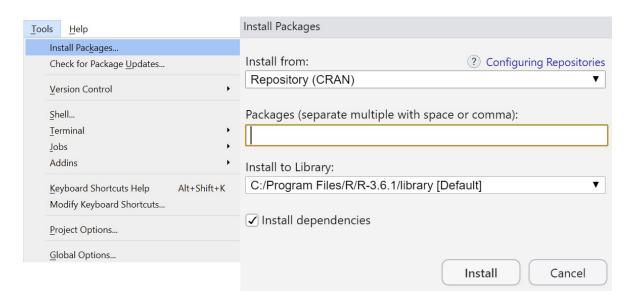
- · Functions from read. family work well
- · Concerns:
  - poor default function behavior
    - (formerly, prior to R 4.0) strings are read as factors
    - row & column names can be troublesome
  - Slower processing
  - (Slightly) different behavior on different computers

- · R package
  - Collection of functions in one place
  - Packages exist to do almost anything
  - List of CRAN approved packages on R's website
  - Plenty of other packages on places like GitHub
- "<u>TidyVerse</u>" collection of R packages that share common philosophies and are designed to work together!

- · First time using a package
  - Must install package (download files)
  - Can use code or menus

```
install.packages("readr")
#can do multiple packages at once
install.packages(c("readr", "readxl", "haven", "DBI", "httr"))
```

- · First time using a package
  - Must install package (download files)
  - Can use code or menus



- · Only install once!
- Each session: read in package using library() or require()

```
library("readr")
require("haven")
```

- · Difference if no package
  - library() throws an error
  - require() returns FALSE

```
library("notAPackage")

## Error in library("notAPackage"): there is no package called 'notAPackage'

require("notAPackage")

## Warning in library(package, lib.loc = lib.loc, character.only = TRUE,
## logical.return = TRUE, : there is no package called 'notAPackage'
```

- · Many packages to do things in R
- · How to choose?
  - Want 'fast' code
  - Want 'easy' syntax
  - Good default settings on functions
  - Nice set of examples and vignettes
- Enter: TidyVerse

Install the tidyverse package

install.packages("tidyverse")

Install the tidyverse package

install.packages("tidyverse")

· Load library

library(tidyverse)

Once library loaded, check help(filter)

- · Can call functions without loading full library with::
- · If not specified, most recently loaded package takes precedent

```
dplyr::filter(neuralgiaData, Treatment == "P")
```

##		Treatment	Sex	Age	Duration	Pain
##	1	P	F	68	1	No
##	2	P	F	67	30	No
##	3	P	M	66	26	Yes
##	4	P	F	64	1	Yes
##	5	P	M	74	4	No
##	6	P	M	70	1	Yes
##	7	P	M	83	1	Yes
##	8	P	M	77	29	Yes
##	9	P	F	79	20	Yes
##	10	P	M	78	12	Yes
##	11	P	M	66	4	Yes
##	12	P	F	65	29	No
##	13	P	M	60	26	Yes
##	14	P	F	72	27	No
##	15	P	F	70	13	Yes
##	16	P	F	68	27	Yes
##	17	P	M	68	11	Yes
##	18	P	M	67	17	Yes
##	19	P	F	67	1	Yes
##	20	P	F	72	11	Yes

Install packages first (download it)

· Can do more than one at a time

Load package with require() or library()

· Call without loading using ::

# **Reading Delimited Data**

We'll use the tidyverse!

#### Function and purpose:

Type of Delimeter	utils Function	readr Function
Comma	read.csv()	read_csv()
Semicolon ( , for decimal)	read.csv2()	read_csv2()
Tab	read.delim()	read_tsv()
General	<pre>read.table(sep = "")</pre>	read_delim()
White Space	<pre>read.table(sep = "")</pre>	<pre>read_table() read_table2()</pre>

## **Reading Delimited Data**

· Let's read in the 'neuralgia.csv' file

neuralgiaData2 <- readr::read\_csv("https://www4.stat.ncsu.edu/~online/datasets/neuralgia.csv")
## Parsed with column specification:
## cols(
## Treatment = col\_character(),
## Sex = col\_character(),
## Age = col\_double(),
## Duration = col\_double(),
## Pain = col\_character()
## Pain = col\_character()</pre>

· Let's read in the 'neuralgia.csv' file

neuralgiaData2

```
## # A tibble: 60 x 5
    Treatment Sex
                      Age Duration Pain
    <chr> <chr> <dbl>
                             <dbl> <chr>
## 1 P
                       68
                                 1 No
## 2 B
                       74
                                16 No
## 3 P
             F
                                30 No
                       67
## 4 P
             M
                       66
                                26 Yes
## 5 B
                       67
                                28 No
## # ... with 55 more rows
```

- · Notice: fancy printing!
- · Checking column type is a basic data validation step
- tidyverse data frames are called tibbles

```
class(neuralgiaData2)
## [1] "spec_tbl_df" "tbl_df" "tbl" "data.frame"
```

#### tibbles

• Behavior slightly different than a standard data frame. No simplification!

```
neuralgiaData[,1]

## [1] PBPPBBABBAAABAPAPAPBBAAABPBBPPAABBBAPB

## [39] BPPPABAPPABPPPBAPABA

## Levels: PBA

neuralgiaData2[,1]

## # A tibble: 60 x 1

## Treatment

## <chr>
## 1 P

## 2 B

## 3 P

## 4 P

## 5 B

## # ... with 55 more rows
```

#### tibbles

- · Behavior slightly different than a standard data frame. No simplification!
- Use either pull() or \$

· How did R determine the column types?

```
help(read csv)
```

- Other useful inputs:
  - skip = 0
  - col names = TRUE
  - na = c("", "NA")

- · Reading clean delimited data pretty easy!
- Let's read in the 'chemical.txt' file (space delimited)
- read\_table2() allows multiple white space characaters between entries

- · Reading *clean* delimited data pretty easy!
- · Let's read in the 'chemical.txt' file (space delimited)
- read table2() allows multiple white space characaters between entries

read table2("https://www4.stat.ncsu.edu/~online/datasets/chemical.txt")

```
## # A tibble: 19 x 4
##
       temp conc time percent
##
      <dbl> <dbl> <dbl>
                           <dbl>
##
   1
       -1
             -1
                   -1
                            45.9
      1
##
    2
             -1
                   -1
                            60.6
##
    3
      -1
             1
                   -1
                            57.5
##
    4
      1
              1
                   -1
                            58.6
    5 -1
             -1
                    1
                            53.3
##
   6
      1
             -1
                    1
                            58
##
    7
      -1
             1
                    1
                            58.8
##
##
    8
      1
              1
                    1
                            52.4
    9
      -2
              0
                    0
##
                            46.9
## 10
              0
        2
                    0
                            55.4
             -2
## 11
        0
                    0
                            55
              2
## 12
        0
                    0
                            57.5
## 13
              0
                   -2
                            56.3
## 14
        0
              0
                    2
                            58.9
## 15
              0
                            56.9
        0
                    0
## 16
        2
             -3
                    0
                            61.1
## 17
        2
             -3
                            62.9
                    0
## 18
      -1.4 2.6
                   0.7
                            60
## 19
      -1.4
              2.6
                    0.7
                            60.6
```

- · Reading *clean* delimited data pretty easy!
- Let's read in the 'crabs.txt' file (tab delimited)

- · Reading clean delimited data pretty easy!
- · Let's read in the 'crabs.txt' file (tab delimited)

```
read_tsv("https://www4.stat.ncsu.edu/~online/datasets/crabs.txt")
```

```
## # A tibble: 173 x 6
    color spine width satell weight
    <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
##
        3
          3 28.3
                             3050
## 2
       4 3 22.5
                                     0
                         0
                             1550
    2 1 26
4 3 24.8
                             2300
## 4
                             2100
                                     0
                        0
      4
            3 26
## 5
                             2600
                                     1
## # ... with 168 more rows
```

- · Reading clean delimited data pretty easy!
- Let's read in the 'umps2012.txt' file ('>' delimited)
- Notice no column names provided
  - Year Month Day Home Away HPUmpire

- · Reading *clean* delimited data pretty easy!
- Let's read in the 'umps2012.txt' file ('>' delimited)
- · Notice no column names provided
  - Year Month Day Home Away HPUmpire

```
read delim("https://www4.stat.ncsu.edu/~online/datasets/umps2012.txt", delim = ">",
          col names = c("Year", "Month", "Day", "Home", "Away", "HPUmpire"))
## # A tibble: 2,359 \times 6
##
     Year Month
               Day Home Away HPUmpire
##
    <dbl> <dbl> <dbl> <chr> <chr> <chr>
## 1 2012
             4 12 MIN
                               D.J. Reyburn
                          LAA
## 2 2012
            4 12 SD
                          ARI Marty Foster
\#\# 3 2012 4 12 WSH CIN Mike Everitt
## 4 2012
            4 12 PHI MIA Jeff Nelson
## 5 2012 4 12 CHC MIL Fieldin Culbreth
## # ... with 2,354 more rows
```

## Reading Fixed Field & Tricky Non-Standard Data

- read\_fwf()
  - reads in data where entries are very structured
- read\_file()
  - reads an entire file into a single string
- read\_lines()
  - reads a file into a character vector with one element per line
- Usually parse the last two with regular expressions :(

# Next Up!

· Read data from other sources

Type of file	Package	Function
Delimited	readr	<pre>read_csv(), read_tsv(),read_table(), read_delim()</pre>
Excel (.xls,.xlsx)	readxl	read_excel()
SAS (.sas7bdat)	haven	read_sas()
SPSS (.sav)	haven	read_spss()

· Resources for JSON, XML, databases, and APIs