Entering / cleaning data 1

Some odds and ends

Missing values

In R, $\mathbb{N}\mathbb{A}$ is used to represent a missing value in a vector. This value can show up in numerical or character vectors (or in vectors of some other classes):

```
c(1, 4, NA)

## [1] 1 4 NA

c("Jane Doe", NA)

## [1] "Jane Doe" NA
```

The \$ operator

We've talked about how you can combine vectors of the same length to create a dataframe.

To go the other direction (pull a column from a dataframe), you can use the \$ operator.

For example, say you have the following dataset and want to pull the color column as a vector:

```
## 1 red 1
## 2 blue 2
```

The \$ operator

You can pull the color column as a vector using the name of the dataframe, the dollar sign, and then the name of the column:

```
example df$color
## [1] red blue
## Levels: blue red
class(example df$color)
## [1] "factor"
(Note: You can use tab completion in RStudio after you put in
example_df$.)
```

Order of evaluation

In R, you can "nest" functions within a single call. Just like with math, the order that the functions are evaluated moves from the inner set of parentheses to the outer one.

For example, to print the structure of the dataframe from the previous example after creating it, you can run:

```
str(data.frame(color = c("red", "blue"), value = c(1, 2)))
## 'data.frame': 2 obs. of 2 variables:
## $ color: Factor w/ 2 levels "blue", "red": 2 1
## $ value: num 1 2
```

paste and paste0

If you want to paste together several character strings to make a length-one character vector, you can use the paste function to do that:

```
paste("abra", "ca", "dabra")
```

```
## [1] "abra ca dabra"
```

By default, spaces are used to separate each original character string in the final string.

paste and paste0

If you want to remove these spaces, you can use the sep argument in the paste function:

```
paste("abra", "ca", "dabra", sep = "")
```

[1] "abracadabra"

A short-cut function is paste0, which is identical to running paste with the argument sep = "":

```
paste0("abra", "ca", "dabra")
```

```
## [1] "abracadabra"
```

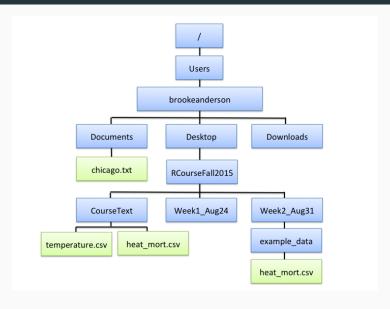
Getting data into R

Basics of getting data into R

Basic approach:

- Download data to your computer
- Make sure R is working in the directory with your data (getwd, setwd)
- Read data into R (functions in readr: read_csv, read_table, read_delim, read_fwf, etc.)
- Check to make sure the data came in correctly (dim, ncol, nrow, head, tail, str, colnames)

Computer directory structure



You can check your working directory anytime using getwd():

```
getwd()
```

```
## [1] "/Users/_gbanders/r_course/RProgrammingForResearch/slides"
```

You can use setwd() to change your directory.

To get to your home directory (for example, mine is "/Users/brookeanderson"), you can use the abbreviation ~.

For example, if you want to change into your home directory and print its name, you could run:

```
setwd("~")
getwd()
```

```
## [1] "/Users/brookeanderson"
```

Remember that, since ~ is a shortcut for my home directory, the following two calls would give me the same result:

```
setwd("~")
getwd()

## [1] "/Users/brookeanderson"

setwd("/Users/brookeanderson")
getwd()
```

```
## [1] "/Users/brookeanderson"
```

The most straightforward way to read in data is often to put it in your working directory and then read it in using the file name. If you're working in the directory with the file you want, you should see the file if you list files in the working directory:

list.files()

```
##
    [1] "CourseNotes_Week1.pdf"
##
       "CourseNotes_Week1.Rmd"
##
    [3]
        "CourseNotes Week10.pdf"
    [4]
       "CourseNotes Week10.Rmd"
##
##
    [5]
        "CourseNotes_Week11.pdf"
##
    [6]
       "CourseNotes Week11.Rmd"
##
    [7]
       "CourseNotes_Week12.pdf"
##
    [8]
        "CourseNotes_Week12.Rmd"
##
    [9]
        "CourseNotes Week13.pdf"
##
   [10]
       "CourseNotes_Week13.Rmd"
   [11] "CourseNotes Week14.pdf"
##
```

The "Files" pane in RStudio (often on the lower right) will also show you the files available in your current working directory.

This should line up with what you get if you run list.files().

Getting around directories

There are a few abbreviations you can use to represent certain relative or absolute locations when you're using setwd():

Shorthand	Meaning
~	Home directory
	Current working directory
	One directory up from current working directory (parent directory)
/	Two directories up from current working directory
/data	The 'data' subdirectory of the parent directory

Taking advantage of paste0

You can create an object with your directory name using paste0, and then use that to set your directory. We'll take a lot of advantage of this for reading in files.

The convention for paste0 is:

Taking advantage of paste0

Here's an example:

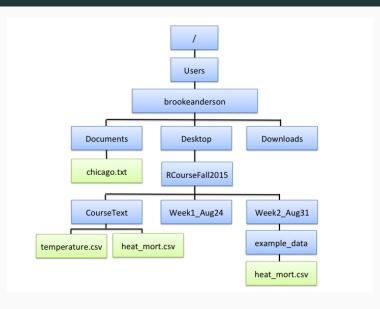
When you want to reference a directory or file, you can use one of two types of pathnames:

- Relative pathname: How to get to the file or directory from your current working directory
- Absolute pathname: How to get to the file or directory from anywhere on the computer

If directions worked like pathnames, here is how you would tell someone to get to this building:

- Relative pathname: Turn right on Center, then turn right when you
 get to the intersection, then go to the second building on your left
 (only works if you started on Prospect going west right before the
 intersection with Center).
- Absolute pathname: Go to 350 W. Lake Street, Fort Collins, CO, USA.

Computer directory structure



Say your current working directory was

/Users/brookeanderson/RProgrammingForResearch and you wanted to get into the subdirectory data. Here are examples using the two types of pathnames:

Absolute:

```
setwd("/Users/brookeanderson/RProgrammingForResearch/data")
```

Relative:

```
setwd("data")
```

Here are some other examples of relative pathnames:

If data is a subdirectory of your current parent directory:

```
setwd("../data")
```

If data is a subdirectory of your home directory:

```
setwd("~/data")
```

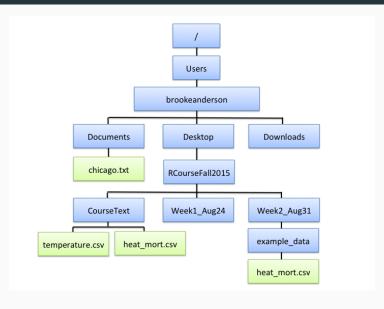
If data is a subdirectory of the subdirectory Ex of your current working directory:

```
setwd("Ex/data")
```

Both methods of writing filenames have their own advantages and disadvantages:

- Relative pathname: Which file you are indicating depends on which working directory you are in, which means that your code will break if you try to re-run it from a different working directory. However, relative pathways in your code make it easier for you to share a working version of a project with someone else. For most of this course, we will focus on using relative pathnames, especially when you start collaborating.
- Absolute pathname: No matter what working directory you're in, it is completely clear to your computer which file you mean when you use an absolute pathname. However, your code will not work on someone else's computer without modifications (because the structure of their computer's full directory will be different).

Computer directory structure



If you want to read in data from a file that is not in your working directory, there are three options:

- Move the data into your working directory (this can be done outside of R).
- Change your working directory so that you are working in the directory that has the data (e.g., with setwd()).
- Use a pathname, rather than a simple filename, to refer to the data file (this will be the recommended method for most of this course, although we'll practice other methods today).

In-course exercise

We'll take a break now to do the first part of the in-course exercise (Sections 2.6.1 and 2.6.2).

Reading data into R

What kind of data can you get into R?

The sky is the limit...

- Flat files
- Files from other statistical packages (SAS, Excel, Stata, SPSS)
- Tables on webpages (e.g., the table near the end of this page)
- Data in a database (e.g., SQL)
- Data stored in XML and ISON
- Really crazy data formats used in other disciplines (e.g., netCDF files from climate folks, MRI data stored in Analyze, NIfTI, and DICOM formats)
- Data through APIs (e.g., GoogleMaps, Twitter, many government agencies)
- Incredibly messy data using scan and readLines

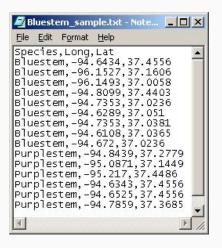
Types of flat files

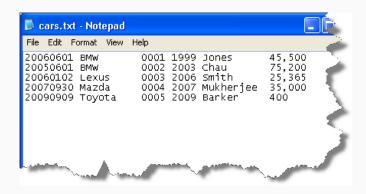
R can read in data from *a lot* of different formats. The only catch: you need to tell R how to do it.

To start, we'll look at flat files:

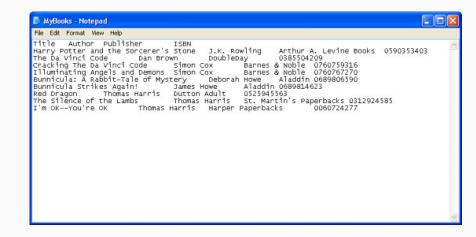
- 1. Fixed width files
- 2. Delimited files
 - ".csv": Comma-separated values
 - ".tab", ".tsv": Tab-separated values
 - Other possible delimiters: colon, semicolon, pipe ("|")

See if you can identify what types of files the following files are. . .





```
H|20110606|pizza.txt|
D|10|Chicken Pesto|20|23|30|5.5|7.4|9.9||
D|10|Meatball|10|53|60|6.5|8.4|10.9
D|10|Fire Cracker|3|13|60|5.8|7.9|11.9|
D|10|Spinach|1|2|5|5.5
D|10|BBQ Chicken|35|102
D|10|Vegetarian|5|13|28|4.
D|10|Mexican|11|33|36|5.
D|10|The Monaco|22|53
D|10|Chilli Prawn|5|5|6|
D|10|Chefs Special|8|18|40
D|10|Marinara|3|17|41|5.
D|10|Supreme|50|52|58|5.5|7
D|10|Margherita|9|19|87
D|10|Napoli|60|85|66|5.2|7.2|9.2
D|10|Caprice|31|32|38|5.5|7.4|9.3|
D|10|Ham and Pineapple|18|39|28|5.8|7.0|9.0|
T|16|
```



What type of file?

File Edit Format View Help Title, Subtitle, Larger Work, Contributor #1, Contributor #2, Contributor #3, Contributor #4, Genre, Publisher, Published Location, Date Published.Instrumentation.Kev.Location.Indiana Connection.Sheet Music Consortium, Notes, Complete """A"" You're Adorable", The alphabet song, Buddy Kaye, Sidney Lippman, Fred Wise, Popular standard, Laurel Music Corporation, "New York, NY", 1948, Voice and piano/guitar or ukulele,c Major,,None,Yes,Perry Como pictured on cover, "Aba Daba Honey Moon, The",,"""Two Weeks with Love"" Motion Picture",Arthur Fields,Walte Donovan,,, "Popular Standard, Movie Selection", Leo Feist Inc., "New York, NY",1942, Voice and Piano, C Minor, None, Yes,, Abi Bezunt, """Mamele"" Motion Picture", Abraham Ellstein, Molly Picon, ,, "Popular Standard Movie Selection", Metro Music Co., "New York, NY", 1939, Voice and Piano, E Minor, , None, No, Molly Picon pictured on cover, Abdul the Bulbul Ameer,,, Bob Kaai, Jim Smock,,, Popular Standard, Calumet Music Co. ."Chicago, IL".1935, "Voice, Piano, Hawaiian Guitar, Ukulele", G Major,,None,Yes,Ben Pollack pictured on cover, About A Quarter to Nine,,"""Go Into Your Dance"" Motion Picture",Harry Warren,Al Dubin,,"Popular Standard, Movie Selection",M. Witmark & Sons, "New York, NY",1935, "Voice, Piano, Guitar, Ukelele", E Minor, None, No, Al Jolson and Ruby Keeler pictured on cover. Absent,,, John. W. Metcalf, Catherine Young Glen,,, Popular Standard, Arthur P. Schmidt, "Boston, MA", 1899, Voice and Piano, G Major, None, Yes, The Academy Two-Step...Barclay Walker....Popular Standard.Carlin & Lennox, "Indianapolis, IN", , Piano, F Major, , Composer, No, , Ac-cent-tchu-ate the Positive, Mister In Between, """Here Come the Waves" Motion Picture", Harold Arlen, Johnny Mercer,,, "Popular Standard, Movie Selection", Edwin H. Morris & Co., "New York, NY", 1944, "Voice, Piano, Guitar", F Major, None, Yes, Bing Crosby and Betty Hutton pictured on cover, Across the Alley From the Alamo,,,, Joe Greene,,,, Popular Standard, Leslie Music

What type of file?

1000235 Baylog 1000236 Gallardo 1000237 Christian 1000238 Baufield 1000239 Frazier 1000240 Garrido 1000241 williams 1000242 Morel Padilla 1000244 Rosenberg 1000245 Blanchard 1000246 wiggins 1000247 miller 1000248 Coon 1000248 Coon	Nick Cathy Mike Daniel Robert Edward Zachary David Damian Wayne Phong S David Jeffrey Terry Walter Timothy
1000234 Faley	John Nick Cathy

Types of flat files

To figure out the structure of a flat file, start by opening it in a text editor.

As an alternative, you can open it in RStudio (right click on the file name and then choose "Open With" and "RStudio").

Reading in flat files

R can read any of the types of files we just looked at by using one of the functions from the readr package (e.g., read_table, read_fwf). Find out more about those functions with:

```
library(readr)
?read_table
?read_fwf
```

read.table family of functions

Some of the interesting options with the readr family of functions are:

Description	
What is the delimiter in the data?	
How many lines of the start of the file should you skip?	
What would you like to use as the column names?	
What would you like to use as the column types?	
How many rows do you want to read in?	
How are missing values coded?	

Many members of the readr package that read delimited files are doing the same basic thing. The only difference is what defaults they have for the separator (delim).

Some key members of the readr family for delimited data:

Function	Separator
read_csv	comma
read_csv2	semi-colon
read_table2	whitespace
read_tsv	tab

For any type of delimited flat files, you can also use the more general read_delim function to read in the file. However, you will have to specify yourself what the delimiter is (e.g., delim = "," for a comma-separated file).

The readr package also includes some functions for reading in fixed width files:

- read_fwf
- read_table

These allow you to specify field widths for each fixed width field, but they will also try to determine the field-widths automatically.

You will also see code that uses functions like read.csv and read.table to read in flat files. These are from the read.table family of functions, which are part of base R. The readr functions are very similar, but have some more sensible defaults, including in determining column classes.

```
## Parsed with column specification:
## cols(
## YEAR = col_integer(),
## GoogleKnowlege_Occupation = col_character(),
## Show = col_character(),
## Group = col_character(),
## Raw_Guest_List = col_character()
## )
```

Compared to the read.table family of functions, the read_* functions:

- Work better with large datasets: faster, includes progress bar
- Have more sensible defaults (e.g., characters default to characters, not factors)

The "tidyverse"

The readr package is part of the "tidyverse"— a collection of recent and developing packages for R, many written by Hadley Wickham.



The "tidyverse"



"A giant among data nerds"

https://priceonomics.com/hadley-wickham-the-man-who-revolutionized-r/

Reading in online flat files

If you're reading in data from a non-secure webpage (i.e., one that starts with http), if the data is in a "flat-file" format, you can just read it in using the web address as the file name:

```
## # A tibble: 5 x 4
                 nC
##
      pos
            nA
                      nG
    <int> <int> <int> <int>
##
## 1
      500
           307
                153
                     192
## 2
     1500
           310 169 207
    2500 319 167 177
## 3
## 4 3500 373 164
                     168
## 5 4500 330
                175
                     224
```

Reading in online flat files

##

With the readr family of functions, you can also read in data from a secure webpage (e.g., one that starts with https). This allows you to read in data from places like GitHub and Dropbox public folders:

<int>

2776

Date Day Cases Guinea

<chr> <int>

1 1/5/2015 289

Reading data from other files types

You can also read data in from a variety of other file formats, including:

File type	Function	Package
Excel	read_excel	readxl
SAS	read_sas	haven
SPSS	read_spss	haven
Stata	read_stata	haven

Saving / loading R objects

You can save an R object you've created as an .RData file using save():

```
save(ebola, file = "Ebola.RData")
list.files(pattern = "E")
```

```
## [1] "Ebola.RData"
```

This saves to your current working directory (unless you specify a different location).

Saving / loading R objects

Then you can re-load the object later using load():

```
rm(ebola)
ls()
## [1] "daily_show"
                            "dirpath_shortcuts"
## [3] "example df"
                            "ld genetics"
## [5] "my dir"
                            "read funcs"
## [7] "url"
load("Ebola.RData")
ls()
## [1] "daily_show"
                            "dirpath shortcuts"
## [3] "ebola"
                            "example_df"
## [5] "ld_genetics"
                            "my dir"
## [7] "read funcs"
                            "url"
```

Saving R objects

One caveat for saving R objects: some people suggest you avoid this if possible, to make your research more reproducible.

Imagine someone wants to look at your data and code in 30 years. R might not work the same, so you might not be able to read an .RData file. However, you can open flat files (e.g., .csv, .txt) and R scripts (.R) in text editors— you should still be able to do this regardless of what happens to R.

Potential exceptions:

- You have an object that you need to save that has a structure that won't work well in a flat file
- Your starting dataset is really, really large, and it would take a long time for you to read in your data fresh every time

In-course exercise

We'll take another break here to work on the next part of the In-Course Exercise (Section 2.6.3).

Data cleaning

Cleaning data

Common data-cleaning tasks include:

Task	dplyr function
Renaming columns	rename
Filtering to certain rows	filter
Selecting certain columns	select
Adding or changing columns	mutate

Cleaning data

As an example, let's look at the Daily Show data:

In cleaning up the data, we'll use dplyr functions, so we need to load that package:

```
library(dplyr)
```

Re-naming columns

A first step is often re-naming columns. It can be hard to work with a column name that is:

- long
- includes spaces
- includes upper case

Several of the column names in daily_show have some of these issues:

```
colnames(daily_show)

## [1] "YEAR"

## [2] "GoogleKnowlege_Occupation"

## [3] "Show"

## [4] "Group"

## [5] "Raw_Guest_List"
```

Renaming columns

To rename these columns, use rename. The basic syntax is:

If you want to change column names in the saved object, be sure you reassign the object to be the output of rename.

Renaming columns

To rename columns in the daily_show data, then, use:

Selecting columns

Next, you may want to select only some columns of the dataframe. You can use select for this. The basic structure of this command is:

```
## Generic code
select(dataframe, column_name_1, column_name_2, ...)
```

Selecting columns

For example, to select all columns except year (since that information is already included in date), run:

```
select(daily_show, job, date, category, guest_name)
```

```
## # A tibble: 2,693 x 4
##
                    job date category
##
                  <chr> <chr>
                                   <chr>>
                  actor 1/11/99 Acting
##
##
   2
               Comedian 1/12/99 Comedy
   3 television actress 1/13/99
##
                                  Acting
           film actress 1/14/99
##
   4
                                  Acting
##
   5
                  actor 1/18/99
                                  Acting
##
   6
                  actor 1/19/99
                                  Acting
   7
        Singer-lyricist 1/20/99 Musician
##
##
   8
                  model 1/21/99
                                  Media
                  actor 1/25/99
##
                                  Acting
      stand-up comedian 1/26/99
                                  Comedy
```

Selecting columns

The select function also provides some time-saving tools. For example, in the last example, we wanted all the columns except one. Instead of writing out all the columns we want, we can use – with the columns we don't want to save time:

```
daily_show <- select(daily_show, -year)
head(daily_show, 3)</pre>
```

Filtering to certain rows

Next, you might want to filter the dataset down so that it only includes certain rows. You can use filter to do that. The syntax is:

```
## Generic code
filter(dataframe, logical statement)
```

The logical statement gives the condition that a row must meet to be included in the output data frame. For example, you might want to pull:

- Rows from 2015
- Rows where the guest was an academic
- Rows where the job is not missing

Filtering to certain rows

For example, if you want to create a data frame that only includes guests who were scientists, you can run:

```
scientists <- filter(daily_show, category == "Science")
head(scientists)</pre>
```

```
## # A tibble: 6 x 4
##
               job date category
##
             <chr> <chr> <chr>
## 1
      neurosurgeon 4/28/03 Science
## 2
         scientist 1/13/04 Science
## 3
         physician 6/15/04 Science
## 4
            doctor 9/6/05 Science
         astronaut 2/13/06 Science
## 5
## 6 Astrophysicist 1/30/07 Science
## # ... with 1 more variables: guest name <chr>
```

Common logical operators in R

To build a logical statement to use in filter, you'll need to know some of R's logical operators:

Operator	Meaning	Example
==	equals	category == "Acting"
!=	does not equal	category != "Comedy
%in%	is in	<pre>category %in% c("Academic", "Science")</pre>
is.na()	is NA	is.na(job)
!is.na()	is not NA	!is.na(job)
&	and	<pre>year == 2015 & category == "Academic"</pre>
1	or	year == 2015 category == "Academic"

Add or change columns

You can change a column or add a new column using the mutate function. That function has the syntax:

Add or change columns

For example, the job column in daily_show sometimes uses upper case and sometimes does not:

```
head(unique(daily_show$job), 10)
```

```
## [1] "actor" "Comedian"
## [3] "television actress" "film actress"
## [5] "Singer-lyricist" "model"
## [7] "stand-up comedian" "actress"
## [9] "comedian" "Singer-songwriter"
```

Add or change columns

We could use the tolower function to make all listings lowercase:

mutate(daily_show, job = tolower(job))

```
## # A tibble: 2,693 x 4
##
                    job date category
##
                  <chr> <chr>
                                   <chr>>
                  actor 1/11/99 Acting
   1
##
##
               comedian 1/12/99 Comedy
##
   3 television actress 1/13/99
                                  Acting
           film actress 1/14/99
                                  Acting
##
   4
                  actor 1/18/99
##
   5
                                  Acting
##
   6
                  actor 1/19/99
                                  Acting
##
   7
        singer-lyricist 1/20/99 Musician
   8
                  model 1/21/99
                                  Media
##
##
                  actor 1/25/99 Acting
  10
##
      stand-up comedian 1/26/99 Comedy
## #
        with 2 683 more roug and 1 more variables.
```

Piping





Piping

If you look at the format of these dplyr functions, you'll notice that they all take a dataframe as their first argument:

Piping

Classically, you would clean up a dataframe in R by reassigning the dataframe object at each step:

```
daily_show <-read_csv("../data/daily_show_guests.csv",
                       skip = 4)
daily show <- rename(daily show,
                      job = GoogleKnowlege_Occupation,
                      date = Show,
                      category = Group,
                      guest_name = Raw_Guest_List)
daily show <- select(daily show, -YEAR)</pre>
daily_show <- mutate(daily_show, job = tolower(job))</pre>
daily_show <- filter(daily_show, category == "Science")</pre>
```

Piping

"Piping" lets you clean this code up a bit. It can be used with any function that inputs a dataframe as its first argument. It "pipes" the dataframe created right before the pipe (%>%) into the function right after the pipe.

Piping

With piping, the same data cleaning looks like:

```
## Parsed with column specification:
## cols(
## YEAR = col_integer(),
## GoogleKnowlege_Occupation = col_character(),
## Show = col_character(),
## Group = col_character().
```

dplyr versus base R

Just so you know, all of these actions also have alternatives in base R:

dplyr	Base R equivalent
rename	Reassign colnames
select	Square bracket indexing
filter	subset
mutate	Use \$ to change / create columns

You will see these alternatives used in older code examples.

Dates in R

Date class

One final common task in cleaning data is to change the class of some of the columns. This is especially common for dates, which will usually be read in as characters or factors.

Vector classes

Here are a few common vector classes in R:

Class	Example	
character	"Chemistry", "Physics", "Mathematics"	
numeric	10, 20, 30, 40	
factor	Male [underlying number: 1], Female [2]	
Date	"2010-01-01" [underlying number: 14,610]	
logical	TRUE, FALSE	

Vector classes

To find out the class of a vector, you can use class():

```
class(daily_show$date)
```

```
## [1] "character"
```

Note: You can use str to get information on the classes of all columns in a dataframe. It's also printed at the top of output from dplyr functions.

[1] "Date"

```
To convert a vector to the Date class, you can use as.Date():
daily_show <- mutate(daily_show,</pre>
                     date = as.Date(date, format = "%m/%d/%y"))
head(daily_show, 3)
## # A tibble: 3 \times 4
##
             job date category
##
          <chr> <date> <chr>
## 1 neurosurgeon 2003-04-28 Science
## 2
       scientist 2004-01-13 Science
## 3 physician 2004-06-15 Science
## # ... with 1 more variables: guest name <chr>
class(daily show$date)
```

Once you have an object in the Date class, you can do things like plot by date, calculate the range of dates, and calculate the total number of days the dataset covers:

```
range(daily_show$date)

## [1] "2003-04-28" "2015-04-23"

diff(range(daily_show$date))

## Time difference of 4378 days
```

The only tricky thing is learning the abbreviations for the format option. You use this option to specify the format of the date **before** you change it to a Date class. Here are some common date format abbreviations:

Abbreviation	Meaning
%m	Month as a number (e.g., 1, 05)
%B	Full month name (e.g., August)
%b	Abbreviated month name (e.g., Aug)
%у	Two-digit year (e.g., 99)
%Y	Four-digit year (e.g., 1999)

Here are some examples:

Current format of date	format =
10/23/2008	"%m/%d%Y"
08-10-23	"%y-%m-%d"
Oct. 23 2008	"%b. %d %Y"
October 23, 2008	"%B %d, %Y"

[1] "2008-10-13"

In many cases you can use functions from the lubridate package to parse dates more easily.

The ymd function from lubridate can be used regardless of the format, as long as the date elements are in the order: year, month, day. For example:

```
library(lubridate)
ymd("2008-10-13")
## [1] "2008-10-13"
vmd("'08 Oct 13")
## [1] "2008-10-13"
ymd("'08 Oct 13")
```

The lubridate package has similar functions for other date orders or for date-times. For example:

- dmy
- mdy
- ymd_h
- ymd_hm

We could have used these to transform the date in daily_show:

```
daily_show <- read_csv("../data/daily_show_guests.csv",</pre>
                        skip = 4) \% \%
  rename(job = GoogleKnowlege_Occupation,
         date = Show,
         category = Group,
         guest_name = Raw_Guest_List) %>%
  select(-YEAR) %>%
  mutate(date = mdy(date)) %>%
  filter(category == "Science")
head(daily_show, 2)
```

```
## # A tibble: 2 x 4
## job date category
## <chr> <date> <chr>
## 1 neurosurgeon 2003-04-28 Science
## 2 scientist 2004-01-13 Science
```

The lubridate package also includes functions to pull out certain elements of a date. For example, we could use wday to create a new column with the weekday of each show:

```
## # A tibble: 5 \times 3
##
          date show_day
                                    guest_name
##
        <date>
                  <ord>
                                         <chr>>
## 1 2003-04-28
                    Mon
                               Dr Sanjay Gupta
## 2 2004-01-13
                   Tues
                               Catherine Weitz
## 3 2004-06-15
                   Tues
                                Hassan Ibrahim
## 4 2005-09-06
                   Tues
                               Dr. Marc Siegel
## 5 2006-02-13 Mon Astronaut Mike Mullane
```

Functions in lubridate for pulling elements from a date include:

- wday
- mday
- yday
- month
- quarter
- year