Investigating objects and data patterns using base R Managing and Manipulating Data Using R

Lecture outline

- 1. Investigate objects, base R
 - 1.1 Functions to describe objects
 - 1.2 Variables names
 - 1.3 View and print data
 - 1.4 Missing values
- 2. Subsetting using subset operators
 - 2.1 Subset atomic vectors using []
 - 2.2 Subsetting lists/data frames using []
 - 2.3 Subsetting lists/data frames using [[]] and \$
 - 2.4 Subsetting data frames with [] combined with \$
- 3. Subsetting using subset() function
- 4. Sorting data

Libraries we will use today

"Load" the package we will use today (output omitted)

```
library(tidyverse)
```

If package not yet installed, then must install before you load. Install in "console" rather than .Rmd file

- Generic syntax: install.packages("package_name")
- Install "tidyverse": install.packages("tidyverse")

Note: when we load package, name of package is not in quotes; but when we install package, name of package is in quotes:

- install.packages("tidyverse")
- library(tidyverse)

Investigate objects, base R

Load .Rdata data frames we will use today

Data on off-campus recruiting events by public universities

- Data frame object df_event
 - One observation per university, recruiting event
- Data frame object df_school
 - One observation per high school (visited and non-visited)

```
rm(list = ls()) # remove all objects in current environment
getwd()
#> [1] "C:/Users/ozanj/Documents/rclass1/lectures/patterns_base_r"
#load dataset with one obs per recruiting event
load(url("https://github.com/ozanj/rclass/raw/master/data/recruiting/recruit_ev
#load("../../data/recruiting/recruit_event_somevars.Rdata")
#load dataset with one obs per high school
load(url("https://github.com/ozanj/rclass/raw/master/data/recruiting/recruit_school_somevars.Rdata")
```

Functions to describe objects

Simple base R functions to describe objects

This section introduces some base R functions to describe objects (some of these you have seen before)

- list objects, list.files() and ls()
- remove objects, rm()
- object type, typeof()
- object length (number of elements), length()
- object structure, str()
- number of rows and columns, ncol() and nrow

I use the functions typeof(), length(), str() anytime I encounter a new object

▶ Helps me understand the object before I start working with it

Listing objects

Files in your working directory

list.files() function lists files in your current working directory

if you run this code from .Rmd file, working directory is location .Rmd file is stored

Objects currently open in your R session

Listing objects currently open in your R session

1s() function lists objects currently open in R

```
x <- "hello!"
ls() # Objects open in R
#> [1] "df_event" "df_school" "x"
```

Removing objects currently open in your R session

rm() function removes specified objects open in R

```
rm(x)
ls()
#> [1] "df_event" "df_school"
```

Command to remove all objects open in R (I don't run it)

```
#rm(list = ls())
```

Base R functions to describe objects, typeof()

typeof() function determines the the internal storage type of an object (e.g., logical vector, integer vector, list)

- syntax
 - tyepof(x)
- arguments
 - x: any R object
- help:

?typeof

Examples

Recall that a data frame is an object where type is a list

```
typeof(c(TRUE,TRUE,FALSE,NA))
#> [1] "logical"
typeof(df_event)
#> [1] "list"
typeof(x = df_event)
#> [1] "list"
```

Base R functions to describe objects, length()

length() function determines the length of an R object

- for atomic vectors and lists, length() is the number of elements in the object
- syntax
 - length(x)
- arguments
 - x : any R object
- help:

?length

Example, length of an atomic vector is

```
length(c(TRUE,TRUE,FALSE,NA))
#> [1] 4
```

Example, length of a list or data frame

- length of a list is the number of elements
- data frame is a list
- length of a data frame = number of elements = number of variables

```
length(df_event) # = num elements = num columns
#> [1] 33
```

Base R functions to describe objects, str()

str() function compactly displays the structure of an R object

- "structure" includes type, length, and attribute of object and also nested objects
- syntax: str(object)
- arguments (partial)
 - bject : any R object
 - max.level: max level of nesting to display nested structures; default NA = all levels
- help: ?str

Example, atomic vectors

```
str(c(TRUE,TRUE,FALSE,NA))
#> logi [1:4] TRUE TRUE FALSE NA
str(object = c(TRUE,TRUE,FALSE,NA))
#> logi [1:4] TRUE TRUE FALSE NA
```

Example, lists/data frames (output omitted)

```
x <- list(c(1,2), list("apple", "orange"), list(2, 3)) # list
str(x)
str(df_event) # data frame</pre>
```

Base R functions to describe objects, ncol() and nrow()

ncol() nrow() and dim() functions

- Description
 - ncol() = number of columns; nrow() = number of rows
- syntax: ncol(x) nrow(x) dim(x)
- arguments
 - x: a vector, array, data frame, or NULL
- value/return:
 - if object x is an atomic vector: ncol() and nrow() returns NULL
 - if object x is a list but not a data frame: ncol() and nrow() returns NULL
 - ▶ if object x is a data frame: ncol() and nrow() returns integer of length 1

Example, object is a data frame

```
ncol(df_event) # num columns = num elements = num variables
#> [1] 33
nrow(df_event) # num rows = num observations
#> [1] 18680
# can wrap ncol() or nrow() within str() to see what functions return
#str(ncol(df_event))
```

Example, object is atomic vector or list that is not a data frame (output omitted)

```
ncol(c(TRUE,TRUE,FALSE,NA)) # atomic vector
x <- list(c(1,2), list("apple", "orange"), list(2, 3)) # list
nrow(x)</pre>
```

Base R functions to describe objects, dim()

dim() function returns the dimensions of an object (e.g., number of rows and columns)

- syntax: dim(x)
- arguments
 - x: a vector, array, data frame, or NULL
- value/return:
 - if object x is a data frame: dim() returns integer of length 2
 - first element = number of rows; second element = number of columns
 - if object x is an atomic vector: dim() returns NULL
 - if object x is a list but not a data frame: dim() returns NULL

Example, object is a data frame

```
dim(df_event) # shows number rows by columns
#> [1] 18680     33

str(dim(df_event)) # can wrap dim() within str() to see what functions return
#> int [1:2] 18680 33
```

Example, object is atomic vector or list that is not a data frame (output omitted)

```
dim(c(TRUE,TRUE,FALSE,NA)) # atomic vector
x <- list(c(1,2), list("apple", "orange"), list(2, 3)) # list
dim(x)</pre>
```

Variables names

names() function

names() function gets or sets the names of elements of an object

- > syntax:
 - pet the names of an object: names(x)
 - > set the names of an object: names(x) <- value
- arguments (partial)
 - x : an R object
 - value: a character vector with same length as object x or NULL
- value/return
 - names(x) returns a character vector of legnth = length(x) in which each element is the name of the element of x

Example, get names (of atomic vector)

```
a <- c(v1=1,v2=2,3,v4="hi!") # named atomic vector

a

#> v1 v2 v4

#> "1" "2" "3" "hi!"

length(a)

#> [1] 4

names(a)

#> [1] "v1" "v2" "" "v4"

length(names(a)) # investigate length of object names(a)

#> [1] 4

str(names(a)) # investigate structure of object names(a)

#> chr [1:4] "v1" "v2" "" "v4"
```

names() function

names() function gets or sets the names of elements of an object

- > syntax:
 - pet the names of an object: names(x)
 - > set the names of an object: names(x) <- value
- arguments (partial)
 - x : an R object
 - value: a character vector with same length as object x or NULL
- value/return
 - names(x) returns a character vector of legnth = length(x) in which each element is the name of the element of x

Example, set names (of atomic vector)

```
names(a) <- NULL # set names of vector a to NULL
a
#> [1] "1" "2" "3" "hi!"
names(a)
#> NULL

names(a) <- c("var1","var2","var3","var4") # set names of vector a
a
#> var1 var2 var3 var4
#> "1" "2" "3" "hi!"
names(a)
#> [1] "var1" "var2" "var3" "var4"
```

Applying names() function to a data frame

Recall that a data frame is an object where type is a list and each element is named

- each element is a variable
- each element name is a variable name

Example (output omitted)

```
names(df_event)
```

Investigate the object names(df_event)

```
typeof(names(df_event)) # type = character vector
#> [1] "character"
length(names(df_event)) # length = number of variables in data frame
#> [1] 33
str(names(df_event)) # structure of names(df_event)
#> chr [1:33] "instnm" "univ_id" "instst" "pid" "event_date" "event_type" ...
```

We can even assign a new object based on names(df_event)

```
names_event <- names(df_event)
typeof(names_event) # type = character vector
#> [1] "character"
length(names_event) # length = number of variables in data frame
#> [1] 33
str(names_event) # structure of names(df_event)
#> chr [1:33] "instnm" "univ_id" "instst" "pid" "event_date" "event_type"
```

Variable names

Refer to specific named elements of an object using this syntax:

object_name\$element_name

When object is data frame, refer to specific variables using this syntax:

- data_frame_name\$varname
- ▶ This approach to isolating variables very useful for investigating data

```
#df_event$instnm
typeof(df_event$instnm)
#> [1] "character"
typeof(df_event$med_inc)
#> [1] "double"
```

Variable names

Data frames are lists with following criteria:

- each element of list is (usually) a vector; each element of list is a variable
- length of data frame = number of variables

```
length(df_event)
#> [1] 33
nrow(df_event)
#> [1] 18680
#str(df event)
```

- each element of the list (i.e., variable) has the same length
 - Length of each variable is equal to number of observations in data frame

```
typeof(df_event$event_state)
#> [1] "character"
length(df_event$event_state)
#> [1] 18680
str(df event$event state)
typeof(df event$med inc)
#> [1] "double"
length(df_event$med_inc)
#> [1] 18680
str(df_event$med inc)
#> num [1:18680] 71714 89122 70137 70137 71024 ...
```

Variable names

The object df_school has one obs per high school

- variable visits_by_100751 shows number of visits by University of Alabama to each high school
- like all variables in a data frame, the var visits_by_100751 is just a vector

We perform calculations on a variable like we would on any vector of same type

```
v <- c(2,4,6)
typeof(v)
#> [1] "double"
length(v)
#> [1] 3
sum(v)
#> [1] 12
```

View and print data

Viewing and printing, data frames

Many ways to view/print a data frame object. Here are three ways:

- 1. Simply type the object name (output omitted)
 - number of observations and rows printed depend on YAML header settings and on object "attributes" (attributes discussed in future week)

df_event

2. Use the View() function to view data in a browser

View(df_event)

3. head() to show the first n rows

#?head
head(df_event, n=5)

Viewing and printing, data frames

obj_name[<rows>,<cols>] to print specific rows and columns of data frame

particularly powerful when combined with sequences (e.g., 1:10)

Examples (output omitted):

Print first five rows, all vars

```
df_event[1:5, ]
```

Print first five rows and first three columns

```
df_event[1:5, 1:3]
```

Print first three columns of the 100th observation

```
df_event[100, 1:3]
```

Print the 50th observation, all variables

```
df_event[50,]
```

Viewing and printing, variables within data frames

Recall that:

b obj_name\$var_name print specifics elements (i.e., variables) of a data frame

```
df_event$zip
```

each element (i.e., variable) of data frame is an atomic vector with length = number of observations

```
typeof(df_event$zip)
#> [1] "character"
length(df_event$zip)
#> [1] 18680
```

each element of a variable is the value of the variable for one observation

Print specific elements (i.e., observations) of variable based on element position

- syntax: obj_name\$var_name[<element position>]
- vectors don't have "rows" or "columns"; they just have elements
- > syntax be combined with sequences (e.g., print first 10 observations)

Viewing and printing, variables within data frames

Print specific elements (i.e., observations) of variable based on element position

```
syntax: obj_name$var_name[<element position>]
```

Example, print individual elements

```
df_event$zip[1:5] # print obs 1-5 of variable for event zip code
#> [1] "01002" "01007" "01020" "01020" "01027"

df_event$zip[1] # print obs 1 of variable for event zip code
#> [1] "01002"

df_event$zip[5] # print obs 5 of variable for event zip code
#> [1] "01027"

df_event$zip[c(1,3,5)] # print obs 5 of variable for event zip code
#> [1] "01002" "01020" "01027"
```

Print specific elements of multiple variables using combine function c()

- > syntax:
 - c(obj_name\$var1_name[<element position>], obj_name\$var2_name[<element positions]
- Example: print first five observations of variables "event_state" and
 "event type"

```
c(df_event$event_state[1:5],df_event$event_type[1:5])
#> [1] "MA" "MA" "MA" "MA" "public hs"
#> [7] "public hs" "public hs" "public hs" "public hs"
```

Exercise

Create a printing exercise using the df_school data frame

- Use obj_name[<rows>,<cols>] to print the first 5 rows and 3 columns of data frame
- 2. Use head() to print first 4 observations
- 3. Use obj_name\$var_name[1:10] to print the first 10 observations of a variable
- Use combine() to print the first 3 observations of variables "school_type" & "name"

 Use obj_name[<rows>,<cols>] to print the first 5 rows and 3 columns of data frame

2. Use head() to print first 4 observations

```
head(df_school, n=4)
#> # A tibble: 4 x 26
#> state code school type ncessch name address city zip code pct white
#> <chr> <chr
#> 1 AK public 020000~ Beth~ 1006 R~ Beth~ 99559
                                                                                             11.8
#> 2 AK public 020000~ Ayaq~ 106 Vi~ Konq~ 99559
                                                                                               0
                   public 020000~ Kwiq~ 108 Vi~ Kwiq~ 99622
#> 3 AK
#> 4 AK
                 public
                                     020000~ Nels~ 118 Vi~ Toks~ 99637
#> # ... with 18 more variables: pct black <dbl>, pct hispanic <dbl>,
#> # pct asian <dbl>, pct amerindian <dbl>, pct other <dbl>, num fr lunch <dbl
#> # total_students <dbl>, num_took_math <dbl>, num_prof_math <dbl>,
         num took rla <dbl>, num prof rla <dbl>, avgmedian inc 2564 <dbl>,
#> #
       visits by 110635 <int>, visits by 126614 <int>, visits by 100751 <int>,
#> #
#> #
       inst 110635 <chr>, inst 126614 <chr>, inst 100751 <chr>
```

3. Use obj_namevar_name[1:10]$ to print the first 10 observations of a variable

```
df_school$name[1:10]
#> [1] "Bethel Regional High School" "Ayagina'ar Elitnaurvik"
#> [3] "Kwigillingok School" "Nelson Island Area School"
#> [5] "Alakanuk School" "Emmonak School"
#> [7] "Hooper Bay School" "Ignatius Beans School"
#> [9] "Pilot Station School" "Kotlik School"
```

Use combine() to print the first 3 observations of variables "school_type" & "name"

Missing values

Missing values

Missing values have the value NA

NA is a special keyword, not the same as the character string "NA"

use is.na() function to determine if a value is missing

is.na() returns a logical vector

```
is.na(5)
#> [1] FALSE
is.na(NA)
#> [1] TRUE
is.na("NA")
#> [1] FALSE
typeof(is.na("NA")) # example of a logical vector
#> [1] "logical"
nvector \leftarrow c(10.5.NA)
is.na(nvector)
#> [1] FALSE FALSE TRUE
typeof(is.na(nvector)) # example of a logical vector
#> [1] "logical"
svector <- c("e", "f", NA, "NA")</pre>
is.na(svector)
#> [1] FALSE FALSE TRUE FALSE
```

Missing values are "contagious"

What does "contagious" mean?

operations involving a missing value will yield a missing value

```
7>5

#> [1] TRUE

7>NA

#> [1] NA

sum(1,2,NA)

#> [1] NA

0==NA

#> [1] NA

2*c(0,1,2,NA)

#> [1] 0 2 4 NA

NA*c(0,1,2,NA)

#> [1] NA NA NA NA
```

Functions and missing values example, table()

table() function is useful for investigating categorical variables

```
str(df_event$event_type)
#> chr [1:18680] "public hs" ...
table(df_event$event_type)
#>
#> 2yr college 4yr college other private hs public hs
#> 951 531 2001 3774 11423
```

Functions and missing values example, table()

By default table() ignores NA values

```
#?table
str(df_event$school_type_pri)
#> int [1:18680] NA NA NA NA 1 1 NA 1 NA ...
table(df_event$school_type_pri)
#>
#> 1 2 5
#> 3765 8 1
```

useNA argument controls if table includes counts of NA s. Allowed values:

```
never ("no") [DEFAULT VALUE]
```

```
only if count is positive ("ifany");
even for zero counts ("always")"
```

```
nrow(df_event)
#> [1] 18680
table(df_event$school_type_pri, useNA="always")
#>
#> 1 2 5 <NA>
#> 3765 8 1 14906
```

Broader point: Most functions that create descriptive statistics have options about how to treat missing values'

▶ When investigating data, good practice to always show missing values

Subsetting using subset operators

Subsetting to Extract Elements

"Subsetting" refers to isolating particular elements of an object

Subsetting operators can be used to select/exclude elements (e.g., variables, observations)

- ▶ there are three subsetting operators: [] , \$, [[]]
- these operators function differently based on vector types (e.g, atomic vectors, lists, data frames)

Wichham refers to number of "dimensions" in R objects

An atomic vector is a 1-dimensional object that contains n elements

```
x <- c(1.1, 2.2, 3.3, 4.4, 5.5)

str(x)

#> num [1:5] 1.1 2.2 3.3 4.4 5.5
```

Lists are multi-dimensional objects

 Contains n elements; each element may contain a 1-dimensional atomic vector or a multi-dimensional list. Below list contains 3 dimensions

```
list <- list(c(1,2), list("apple", "orange"))
str(list)
#> List of 2
#> $: num [1:2] 1 2
#> $:List of 2
#> ..$: chr "apple"
#> ..$: chr "orange"
```

Data frames are 2-dimensional lists

- ▶ each element is a variable (dimension=columns)
- within each variable, each element is an observation (dimension=rows)

```
ncol(df_school)

#> [1] 26

nrow(df_school)

#> [1] 21301
```

Subset atomic vectors using []

Subsetting elements of atomic vectors

"Subsetting" a vector refers to isolating particular elements of a vector

- ▶ I sometimes refer to this as "accessing elements of a vector"
- > subsestting elements of a vector is similar to "filtering" rows of a data-frame
- is the subsetting function for vectors

Six ways to subset an atomic vector using []

- 1. Using positive integers to return elements at specified positions
- 2. Using negative integers to exclude elements at specified positions
- 3. Using logicals to return elements where corresponding logical is TRUE
- 4. Empty [] returns original vector (useful for dataframes)
- 5. Zero vector [0], useful for testing data
- If vector is "named," use character vectors to return elements with matching names

1. Using positive integers to return elements at specified positions (subset atomic vectors using [])

Create atomic vector x

```
(x <- c(1.1, 2.2, 3.3, 4.4, 5.5))

#> [1] 1.1 2.2 3.3 4.4 5.5

str(x)

#> num [1:5] 1.1 2.2 3.3 4.4 5.5
```

- [] is the subsetting function for vectors
 - contents inside [] can refer to element number (also called "position").
 - e.g., [3] refers to contents of 3rd element (or position 3)

```
x[5] #return 5th element
#> [1] 5.5

x[c(3, 1)] #return 3rd and 1st element
#> [1] 3.3 1.1

x[c(4,4,4)] #return 4th element, 4th element, and 4th element
#> [1] 4.4 4.4 4.4

#Return 3rd through 5th element
x[3:5]
#> [1] 3.3 4.4 5.5
```

2. Using negative integers to exclude elements at specified positions (subset atomic vectors using [])

Before excluding elements based on position, investigate object

```
x
#> [1] 1.1 2.2 3.3 4.4 5.5

length(x)
#> [1] 5
str(x)
#> num [1:5] 1.1 2.2 3.3 4.4 5.5
```

Use negative integers to exclude elements based on element position

```
x[-1] # exclude 1st element
#> [1] 2.2 3.3 4.4 5.5

x[c(3,1)] # 3rd and 1st element
#> [1] 3.3 1.1
x[-c(3,1)] # exclude 3rd and 1st element
#> [1] 2.2 4.4 5.5
```

3. Using logicals to return elements where corresponding logical is TRUE (subset atomic vectors using [])

```
x
#> [1] 1.1 2.2 3.3 4.4 5.5
```

When using x[y] to subset x, good practice to have length(x) == length(y)

```
length(x) # length of vector x
#> [1] 5
length(c(TRUE,FALSE,TRUE,FALSE,TRUE)) # length of y
#> [1] 5
length(x) == length(c(TRUE,FALSE,TRUE,FALSE,TRUE)) # condition true
#> [1] TRUE
x[c(TRUE,TRUE,FALSE,FALSE,TRUE)]
#> [1] 1.1 2.2 5.5
```

Recycling rules:

in x[y], if x is different length than y, R "recycles" length of shorter to match length of longer

```
length(c(TRUE,FALSE))
#> [1] 2
x
#> [1] 1.1 2.2 3.3 4.4 5.5
x[c(TRUE,FALSE)]
#> [1] 1.1 3.3 5.5
```

3. Using logicals to return elements where corresponding logical is TRUE (subset atomic vectors using [])

```
x
#> [1] 1.1 2.2 3.3 4.4 5.5
```

Note that a missing value (${\tt NA}$) in the index always yields a missing value in the output

```
x[c(TRUE, FALSE, NA, TRUE, NA)]
#> [1] 1.1 NA 4.4 NA
```

Return all elements of object x where element is greater than 3

```
x # pritn object X
#> [1] 1.1 2.2 3.3 4.4 5.5
x>3 # for each element of X, print T/F whether element value > 3
#> [1] FALSE FALSE TRUE TRUE TRUE
x[x>3]
#> [1] 3.3 4.4 5.5
```

CRYSTAL/PATRICIA - ADD ONE OR TWO MORE PRACTICAL EXAMPLES?

> ?THIS APPROACH IS USED A LOT ON REAL DATA WORK; WANT STUDENTS TO BE ABLE TO SEE HOW IT IS USED. SOMETHING LIKE:?

```
      names (df_school)

      #> [1] "state_code" "school_type" "ncessch"

      #> [4] "name" "address" "city"
```

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4. Empty [] returns original vector (subset atomic vectors using [])

```
x
#> [1] 1.1 2.2 3.3 4.4 5.5
x[]
#> [1] 1.1 2.2 3.3 4.4 5.5
```

This is useful for sub-setting data frames, as we will show below

5. Zero vector [0] (subset atomic vectors using [])

Zero vector, x[0]

R interprets this as returning element 0

```
x[0]
#> numeric(0)
```

Wickham states:

"This is not something you usually do on purpose, but it can be helpful for generating test data."

6. If vector is named, character vectors to return elements with matching names (subset atomic vectors using [])

Create vector y that has values of vector x but each element is named

```
x

#> [1] 1.1 2.2 3.3 4.4 5.5

(y <- c(a=1.1, b=2.2, c=3.3, d=4.4, e=5.5))

#> a b c d e

#> 1.1 2.2 3.3 4.4 5.5
```

Return elements of vector based on name of element

enclose element names in single '' or double "" quotes

```
#show element named "a"
y["a"]
#> a
#> 1.1

#show elements "a", "b", and "d"
y[c("a", "b", "d")]
#> a b d
#> 1.1 2.2 4.4
```

Subsetting lists/data frames using []

Subsetting lists using []

Using [] operator to subset lists works the same as subsetting atomic vector

Using [] with a list always returns a list

```
list_a <- list(list(1,2),3,"apple")</pre>
str(list a)
#> List of 3
#> $ :List of 2
#> ..$ : num 1
#> ..$ : num 2
#> $ : num 3
#> $ : chr "apple"
#create new list that consists of elements 3 and 1 of list_a
list_b \leftarrow list_a[c(3, 1)]
str(list b)
#> List of 2
#> $ : chr "apple"
#> $ :List of 2
#> ..$ : num 1
#> ..$ : num 2
#show elements 3 and 1 of object list a
#str(list a[c(3, 1)])
```

Subsetting data frames using []

Recall that a data frame is just a particular kind of list

- each element = a column = a variable
- Using [] with a list always returns a list
 - Using [] with a data frame always returns a data frame

Two ways to use [] to extract elements of a data frame

- use "single index" df_name[<columns>] to extract columns (variables) based on element position number (i.e., column number)
- use "double index" df_name[<rows>, <columns>] to extact particular rows and columns of a data frame

Subsetting data frames using [] to extract columns (variables) based on element position

Use "single index" df_name[<columns>] to extract columns (variables) based on element number (i.e., column number)

Examples [output omitted]

```
names(df_event)

#extract elements 1 through 4 (elements=columns=variables)
df_event[1:4]
df_event[c(1,2,3,4)]

str(df_event[1:4])
#extract columns 13 and 7
df_event[c(13,7)]
```

Subsetting Data Frames to extract columns (variables) and rows (observations) based on positionality

use "double index" syntax df_name[<rows>, <columns>] to extact particular rows and columns of a data frame

often combined with sequences (e.g., 1:10)

```
#Return rows 1-3 and columns 1-4
df event[1:3, 1:4]
#> # A tibble: 3 x 4
#> instnm univ id instst pid
\#> < chr> < int> < chr> < int>
#> 1 UM Amherst 166629 MA 57570
#> 2 UM Amherst 166629 MA 56984
#> 3 UM Amherst 166629 MA 57105
#Return rows 50-52 and columns 10 and 20
df event [50:52, c(10,20)]
#> # A tibble: 3 x 2
#> event state pct tworaces zip
#> <ch.r>
                          <d.h1.>
                          1.98
#> 1 MA
#> 2 MA
                       1.98
#> 3 MA
                          1.98
```

Subsetting Data Frames to extract columns (variables) and rows (observations) based on positionality

```
use "double index" syntax df_name[<rows>, <columns>] to extact particular rows
and columns of a data frame

recall that empty [] returns original object (output omitted)

#return original data frame
df_event[]

#return specific rows and all columns (variables)
df_event[1:5, ]

#return all rows and specific columns (variables)
df_event[, c(1,2,3)]
```

Use [] to extract data frame columns based on variable names

Selecting columns from a data frame by subsetting with $\ [\]$ and list of element names (i.e., variable names) enclose in quotes

"single index" approach extracts specific variables, all rows (output omittted)

```
df_event[c("instnm", "univ_id", "event_state")]
select(df_event,instnm,univ_id,event_state) # same same
```

"Double index" approach extracts specific variables and specific rows

syntax df_name[<rows>, <columns>]

Student exercises

Use subsetting operators from base R in extracting columns (variables), observations:

- Use both "single index" and "double index" in subsetting to create a new dataframe by extracting the columns instmm, event_date, event_type from df_event. And show what columns (variables) are in the newly created dataframe.
- 2. Use subsetting to return rows 1-5 of columns $\mbox{state_code}$, \mbox{name} , $\mbox{address}$ from $\mbox{df_school}.$

Solution to Student Exercises

Solution to 1

base R using subsetting operators

```
# single index
df_event_br <- df_event[c("instnm", "event_date", "event_type")]
#double index
df_event_br <- df_event[, c("instnm", "event_date", "event_type")]
names(df_event_br)
#> [1] "instnm" "event_date" "event_type"
```

Solution to 2

base R using subsetting operators

Subsetting lists/data frames using [[]] and \$

Subset single element from object using [[]] operator, atomic vectors

So far we have used [] to excract elements from an object

- ▶ Apply [] to atomic vector: returns atomic vector with elements you requested
- Apply [] to list: returns list with elements you requested
- [[]] also extract elements from an object
- ▶ Applying [[]] to atomic vector gives same result as []; that is, an atomic vector with element you request

```
(x <- c(1.1, 2.2, 3.3, 4.4, 5.5))
#> [1] 1.1 2.2 3.3 4.4 5.5

str(x[3])
#> num 3.3

str(x[[3]])
#> num 3.3
```

- ► Applying [[]] to a list
 - Understanding what [] vs. [[]] does to a list is very important but requires some explanation!

Subsetting lists using [] vs. [[]], introduce "train metaphor"

Advanced R chapter 4.3 by Wickham uses the "train metaphor" to explain a list vs. contents of a list and how this relates to [] vs. [[]]

Below code chunk makes a list named x that contains 3 elements

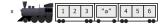
```
x \leftarrow list(1:3, "a", 4:6) # create list object x
```

In our train metaphor, object x is a train that contains 3 carriages



Subsetting lists using [] vs. [[]], introduce "train metaphor"

list object x is a train that contains 3 carriages



When we "subset a list" – that is, extract one or more elements from the list – we have two broad choices (image below)



1. Extracting elements using [] always returns a list, usually one with fewer elements

you can think of this as a train with fewer carriages

```
str(x[1]) # returns a list
#> List of 1
#> $ : int [1:3] 1 2 3
```

- 2. Extracting element using [[]] returns contents of particular carriage
 - I say applying [[]] to a list or data frame returns a simpler object that moves up one level of hierarchy

```
str(x[[1]]) # returns an atomic vector
#> int [1:3] 1 2 3
```

```
Subset lists using [] vs. [[]], deepen understanding of []
```

Rules about applying subset operator [] to a list

- Applying [] to a list always returns a list
- Resulting list contains 1 or more elements depending on what typed inside []

Here is a list object named x

```
x <- list(1:3, "a", 4:6)
```

Here is an image of a few "trains" that can be created by applying [] to x

And here is code to create the "trains" shown in above image (output omitted)

```
x[1:2]
x[-2]
x[c(1,1)]
x[0]
x[] # returns the original list; not shown in above train picture
```

```
Subset lists using [] vs. [[]], deepen understanding of [[]]
```

Rules about applying subset operator [[]] to a list

Can apply [[]] to return the contents of a single element of a list

Create list x and show "train" Image of applying x[1] vs. x[[1]]

```
x \leftarrow list(1:3, "a", 4:6)
```



Object created by x[1] is a list with one element (output omitted)

```
x[1]
str(x[1])
```

Object created by x[[1]] is a vector with 3 elements (output omitted)

- x[[1]] gives us "contents" of element 1
- Since element 1 contains a numeric vector, object created by x[[1]] is a numeric vector

```
x[[1]]
str(x[[1]])
```

```
Subset lists using [] vs. [[]] , deepen understanding of [[]]
```

Rules about applying subset operator [[]] to a list

Can apply [[]] to return the contents of a single element of a list

```
x <- list(1:3, "a", 4:6) # create list x
```

We cannot use [[]] to subset multiple elements of a list (output omitted)

e.g., we could write x[[2]] but not x[[2:3]]

```
x[[c(2)]] # this works, subset element 2 using [[]]
x[[c(2,3)]] # this doesn't work; subset element 2 and 3 using [[]]
x[c(2,3)] # this works; subset element 2 and 3 using []
```

```
Subset lists using [] vs. [[]], deepen understanding of [[]]
```

Like [] , can use [[]] to return contents of named elements specified using quotes

```
syntax: obj_name[["element_name"]]
```

Same list as before, but this time elements named

```
x <- list(var1=1:3, var2="a", var3=4:6)
```

Subset list x using [[]] element names vs. element position

We can do the same thing with data frames because data frames are lists

```
▶ e.g., df_event[["zip"]] returns contents of element named "zip"
```

object created by df_event[["zip"]] is character vector of length = 18,680

```
# df_event[["zip"]] # this works but long output
str(df_event[["zip"]])
#> chr [1:18680] "01002" "01007" "01020" "01020" "01027" "01027" "01027" ...
typeof(df_event[["zip"]])
#> [1] "character"
65,
```

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General rules of applying [] vs [[]] to (nested) objects

What we just learned about applying [] vs [[]] to lists applies more generally to "nested objects"

"nested objects" are objects with a hierarchical structure such that an element of an object contains another object

General rules of applying [] vs. [[]] to nested objects

- subset any object x using [] will return object with same data structure as x
- subset any object x using [[]] will return an object thay may or may not have same data structure of x
 - if object x is not a nested object, then applying [[]] to a single element of x will return object with same data structure as x
 - if object x has a nested data structure, then then applying [[]] to a single element of x will "move up one level of hierarchy" to extract the **contents** of element x

Subset lists/data frames using \$

```
x <- list(var1=1:3, var2="a", var3=4:6)
```

obj_name\$element_name is shorthand operator for obj_name[["element_name"]]

These three lines of code all give the same result

```
x[[1]]
#> [1] 1 2 3
x[["var1"]]
#> [1] 1 2 3
x$var1
#> [1] 1 2 3
```

df_name\$var_name: easiest way in base R to refer to variable in a data frame

these two lines of code are equivalent

```
str(df_event[["zip"]])
#> chr [1:18680] "01002" "01007" "01020" "01020" "01027" "01027" "01027" "01027" ...
str(df_event$zip)
#> chr [1:18680] "01002" "01007" "01020" "01020" "01027" "01027" "01027" ...
```

Subsetting data frames with [] combined with \$

Subsetting Data Frames with [] combined with \$

Combine [] with \$ to subset data frame same as filter() or subset()

Syntax: df_name[df_name\$var_name <condition>,]

- Note: Uses "double index" df_name[<rows>, <columns>] syntax
- ► Cannot use "single index" df_name[<columns>]

Examples (output omitted)

► All observations where the hich school received at least 1 visit from UC Berkeley (var= visits_by_110635) and all columns

```
df_school[df_school$visits_by_110635 >= 1, ]
```

▶ All obs where the high school received at least 1 visit from UC Berkeley and the first three columns

```
df_school[df_school$visits_by_110635 == 1, 1:3]
```

▶ All obs where the high school received at least 1 visit from UC Berkeley and variables "state_code" "school_type" "name"

```
df_school[df_school$visits_by_110635 == 1, c("state_code", "school_type", "name")
```

Subsetting Data Frames with [] combined with \$

Combine [] with \$ to subset data frame same as filter() or subset()

- Syntax: df_name[df_name\$var_name <condition>,]
- ▶ Can be combined with count() or nrow() to avoid printing many rows

Count obs where high schools received at least 1 visit by Bama (100751) and at least one visit by Berkeley (110635)

compare with filter() and subset() approaches

```
#[] combined with $ approach
count(df_school[df_school$visits_by_110635 >= 1
  & df school\$visits by 100751 >= 1, ])
#> # A tibble: 1 x 1
#>
         n
\#> \langle i,n,t,>
#> 1 247
count(df_school[df_school[["visits_by_110635"]] >= 1
  & df school[["visits by 100751"]] >= 1, ])
#> # A tibble: 1 x 1
#>
#> <int>
#> 1 247
df school[]
#> # A tibble: 21,301 x 26
                                                                             70 / 93
```

Subsetting Data Frames with [] and \$, NA Observations

When sub-setting via [] combined with \$, result will include:

- rows where condition is TRUE
- **as well as** rows with NA (missing) values for condition.

Task: How many events at public high schools with at least \$50k median household income

extracting observations via [] combined with \$

```
#num obs event type=="public hs" and med inc is missing
nrow(df event[df event$event type == "public hs"
  & is.na(df_event$med_inc)==1 , ])
#> [1] 75
#num obs event type=="public hs" & med inc is not NA & med inc >= $50,000
nrow(df event[df event$event type == "public hs"
  & is.na(df_event$med_inc)==0 & df_event$med_inc>=50000 , ])
#> [1] 9941
#num obs event type=="public hs" and med inc >= $50,000
nrow(df_event[df_event$event_type == "public hs"
  & df event$med inc>=50000 , ])
#> \[ 11 \] 10016
```

Subsetting Data Frames with [] and \$, NA Observations subset using [] combined with \$, result includes:

rows where condition TRUE; AND rows with NA for condition

Base R filter using subset() excludes rows with NA for condition

```
#num obs event type=="public hs" and med inc is missing
nrow(subset(df_event, event_type == "public hs" & is.na(med_inc)==1))
#> [1] 75
#num obs event type=="public hs" & med inc is not NA & med inc >= $50,000
nrow(subset(df_event, event_type == "public hs" & is.na(med_inc)==0
 & med inc>=50000))
#> [1] 9941
#num obs event tupe=="public hs" & med inc >= $50,000
nrow(subset(df event, event type == "public hs"
 & med_inc>=50000))
#> [1] 9941
```

Tidyverse filter() excludes rows with NA for condition.

```
#num obs event type=="public hs" and med inc is missing
nrow(filter(df_event, event_type == "public hs", is.na(med_inc)==1))
#> [1] 75
#num obs event type=="public hs" & med inc is not NA & med inc >= $50,000
nrow(filter(df_event, event_type == "public hs", is.na(med_inc)==0, med_inc>=50
#> [1] 9941
#num obs event type=="public hs" & med inc >= $50,000
```

prov(filter(df eyent eyent type == "public be" med inc>=50000))

Subsetting Data Frames with [] and \$, NA Observations

To exclude rows where condition is NA if subset using [] combined w/ \$

- ▶ use which() to ask only for values where condition evaluates to TRUE
- which() returns position numbers for elements where condition is TRUE

```
#?which
c(TRUE,FALSE,NA,TRUE)
#> [1] TRUE FALSE NA TRUE
str(c(TRUE,FALSE,NA,TRUE))
#> logi [1:4] TRUE FALSE NA TRUE
which(c(TRUE,FALSE,NA,TRUE))
#> [1] 1 4
```

Task: Count events at public HS with at least \$50k median household income?

Student Exercises

Subsetting Data Frames with (1) [] and \$; (2) subset() and filter():

- 1. Show how many public high schools in California with at least 50% Latinx (hispanic in data) student enrollment from df_school.
- 2. Show how many out-state events at public high schools with more than \$30K median from df_event (do not forget to exclude missing values).

Solution to 1

base R using [] and \$

base R using subset() function

tidyverse using filter() function

use is.na to exclude NA

Solution to 2:

#> <int>

#> 1 7781

```
base R using [] and $ (NA included)
```

& event inst =="Out-State"& df event\$med inc > 30000))

#> [1] 7784 tidyverse using filter() function (NA excluded)

base R using subset() function (NA excluded)
nrow(subset(df_event, event_type == "public hs"

Subsetting using subset() function

Subset function

The subset() is a base R function and easiest way to "filter" observations

- can also used subset() to select variables
- Like tidyverse filter(), subset() can be combined with:
 - with assignment (<-) to create new objects
 - with count() to count number of observations that satisfy criteria

?subset

Syntax [when object is data frame]: subset(x, subset, select, drop = FALSE)

- x is object to be subset
- subset is the logical expression(s) (evaluates to TRUE/FALSE) indicating elements (rows) to keep
- select indicates columns to select from data frame (if argument is not used default will keep all columns)
- drop to preserve original dimensions [SKIP]
 - cane take values TRUE or FALSE; default is FALSE
 - only need to worry about dataframes when subset output is single column

Using df_school , show all public high schools that are at least 50% Latinx (var= $pct_hispanic$) student enrollment in California

Using tidyverse filter() [output omitted]

```
filter(df_school, school_type == "public", pct_hispanic >= 50,
    state_code == "CA")

filter(df_school, school_type == "public" & pct_hispanic >= 50
    & state_code == "CA") # same as above
```

Using base R, subset() [output omitted]

```
#public high schools with at least 50% Latinx student enrollment
subset(df_school, school_type == "public" & pct_hispanic >= 50
    & state_code == "CA")
```

Count all CA public high schools that are at least 50% Latinx

Can wrap filter() or subset() within count() to count number of observations that satisfy criteria

```
#filter()
count(filter(df_school, school_type == "public", pct_hispanic >= 50,
   state code == "CA"))
#> # A tibble: 1 x 1
#> n.
#> <int>
#> 1 713
count(filter(df_school, school_type == "public" & pct_hispanic >= 50
   & state_code == "CA"))
#> # A tibble: 1 x 1
#> n.
\#> \langle i,n,t,>
#> 1 713
#subset()
count(subset(df school, school type == "public" & pct hispanic >= 50
     & state_code == "CA"))
#> # A tibble: 1 x 1
#>
\#> \langle i,n,t,>
#> 1 713
```

Note that both filter() and subset() identify the number of observations for which the condition is TRUE

```
count(filter(df_school, TRUE))
#> # A tibble: 1 x 1
#> n.
#> <int>
#> 1 21301
count(subset(df_school, TRUE))
#> # A tibble: 1 x 1
#> n.
\#> \langle i,n,t,>
#> 1 21301
count(filter(df school, FALSE))
#> # A tibble: 1 x 1
#> n.
#> <int>
#> 1
count(subset(df_school, FALSE))
#> # A tibble: 1 x 1
#>
#> <int>
#> 1
```

Count all CA public high schools that are at least 50% Latinx and received at least 1 visit from UC Berkeley (var= visits_by_110635)

```
#filter()
count(filter(df_school, school_type == "public", pct_hispanic >= 50,
   state_code == "CA", visits_by_110635 >= 1))
#> # A tibble: 1 x 1
#>
\#> \langle i,n,t,>
#> 1 100
#subset()
count(subset(df school, school type == "public" & pct hispanic >= 50
  & state_code == "CA" & visits_by_110635 >= 1))
#> # A tibble: 1 x 1
#>
\#> \langle i,n,t,>
#> 1 100
```

 ${\tt subset()}$ can also use ${\tt \%in\%}$ operator, which is more efficient version of ${\tt OR}$ operator |

 Count number of schools from MA, ME, or VT that received at least one visit from University of Alabama (var= visits_by_100751)

```
#filter()
count(filter(df school, state code %in% c("MA", "ME", "VT"),
  visits_by_100751 >= 1))
#> # A tibble: 1 x 1
#>
\#> \langle i,n,t,>
#> 1 108
#subset()
count(subset(df school, state code %in% c("MA", "ME", "VT")
  & visits_by_100751 >= 1))
#> # A tibble: 1 x 1
#>
\#> \langle i,n,t,>
#> 1 108
```

Use the select argument within subset() to keep selected variables

```
syntax: select = c(var_name1,var_name2,...,var_name_n)
```

Subset all CA public high schools that are at least 50% Latinx **AND** only keep variables name and address

```
subset(df_school, school_type == "public" & pct_hispanic >= 50
           & state_code == "CA", select = c(name, address))
#> # A tibble: 713 x 2
#> n.a.me
                            address
#> <chr>
                          \langle chr \rangle
#> 1 Tustin High 1171 El Camino Real
#> 2 Bell Gardens High 6119 Agra St.
#> 3 Santa Ana High 520 W. Walnut
#> 4 Warren High
                        8141 De Palma St.
#> 5 Hollywood Senior High 1521 N. Highland Ave.
#> 6 Venice Senior High 13000 Venice Blvd.
                 1201 Brewster Ave.
#> 7 Sequoia High
#> 8 Santa Barbara Senior High 700 E. Anapamu St.
#> 9 Santa Paula High 404 N. Sixth St.
#> 10 Azusa High
                          240 N. Cerritos Ave.
#> # ... with 703 more rows
```

Combine subset() with assignment (<-) to create a new data frame

Create a new date frame of all CA public high schools that are at least 50% Latinx AND only keep variables <code>name</code> and <code>address</code>

```
df_school_v2 <- subset(df_school, school_type == "public" & pct_hispanic >= 50
 & state code == "CA", select = c(name, address))
head(df school v2, n=5)
#> # A tibble: 5 x 2
#> name
                       address
#> <chr>
          <chr>
#> 1 Tustin High 1171 El Camino Real
#> 2 Bell Gardens High 6119 Agra St.
#> 3 Santa Ana High 520 W. Walnut
#> 4 Warren High 8141 De Palma St.
#> 5 Hollywood Senior High 1521 N. Highland Ave.
nrow(df school v2)
#> [1] 713
```

Student Exercises

Compare tidyverse to subset() from base R in extracting columns (variables), observations:

- Use both base R and tidyverse to create a new dataframe by extracting the columns instnm, event_date, event_type from df_event. And show what columns (variables) are in the newly created dataframe.
- Use both base R and tidyverse to create a new dataframe from df_school that includes out-of-state public high schools with 50%+ Latinx student enrollment that received at least one visit by the University of California Berkeley (var= visits_by_110635). And count the number of observations.
- Use both base R and tidyverse to count the number of public schools from CA, FL or MA that received one or two visits from UC Berkeley from df_school.
- 4. Use base R to subset all public out-of-state high schools visited by University of California Berkeley that enroll at least 50% Black students, and only keep variables "state_code", "name" and "zip_code".

```
Solution to 1
```

```
base R using subset() function
df_event_br <- subset(df_event, select=c(instnm, event_date, event_type))</pre>
names(df event br)
#> [1] "instnm" "event_date" "event_type"
tidvverse using select() function
df_event_tv <- select(df_event, instnm, event_date, event_type)</pre>
names(df event tv)
#> [1] "instnm" "event date" "event type"
Solution to 2
base R using subset() function
df_school_br <- subset(df_school, state_code != "CA" & school_type == "public"</pre>
                         & pct_hispanic >= 50 & visits_by_110635 >=1 )
nrow(df school br)
#> \[ \int 17 \] 10
```

tidyverse using filter() function

```
df_school_tv <- filter(df_school, state_code != "CA" & school_type == "public"</pre>
                        & pct_hispanic >= 50 & visits_by_110635 >=1 )
nrow(df school tv)
```

Solution to 3

base R using subset() function

tidyverse using filter() function

Solution to 4

base R using subset() function

```
subset(df_school, school_type == "public" & state_code != "CA"
      & visits_by_100751 >= 1 & pct_hispanic >= 50,
      select = c(state_code, name, zip_code))
#> # A tibble: 73 x 3
#> state_code name
                                               zip code
#> <chr> <chr>
                                               <chr>>
#> 1 AZ Aqua Fria High School
                                               85323
#> 2 AZ Desert Edge High School
                                               85338
#> 3 AZ Tempe High School
                                               85281
#> 4 AZ
              Westview High School
                                               85353
#> 5 AZ
              Apollo High School
                                               85302
#> 6 AZ
              South Mountain High School
                                               85040
#> 7 AZ
              Tolleson Union High School
                                               85353
#> 8 CO
              THORNTON HIGH SCHOOL
                                               80229
#> 9 CO
              MARTIN LUTHER KING JR. EARLY COLLEGE 80249
#> 10 CO BATTLE MOUNTAIN HIGH SCHOOL
                                               81620
#> # ... with 63 more rows
```

Sorting data

Base R sort() for vectors

sort() is a base R function that sorts vectors

Syntax: sort(x, decreasing=FALSE, ...)

- where x is object being sorted
- By default it sorts in ascending order (low to high)
- Need to set decreasing argument to TRUE to sort from high to low

```
#?sort()
x<- c(31, 5, 8, 2, 25)
sort(x)
#> [1] 2 5 8 25 31
sort(x, decreasing = TRUE)
#> [1] 31 25 8 5 2
```

Base R order() for dataframes

order() is a base R function that sorts vectors

- Syntax: order(..., na.last = TRUE, decreasing = FALSE)
- where ... are variable(s) to sort by
- ▶ By default it sorts in ascending order (low to high)
- ▶ Need to set decreasing argument to TRUE to sort from high to low

Descending argument only works when we want either one (and only) variable descending or all variables descending (when sorting by multiple vars)

 use - when you want to indicate which variables are descending while using the default ascending sorting

```
df_event[order(df_event$event_date), ]
df_event[order(df_event$event_date, df_event$total_12), ]
#sort descending via argument
df_event[order(df_event$event_date, decreasing = TRUE), ]
df_event[order(df_event$event_date, df_event$total_12, decreasing = TRUE), ]
#sorting by both ascending and descending variables
df_event[order(df_event$event_date, -df_event$total_12), ]
```

Compare tidyverse to base r, sorting

```
-Create a new dataframe from df_events that sorts by ascending by event_date, ascending event_state, and descending pop_total .

tidyverse

df_event_tv <- arrange(df_event, event_date, event_state, desc(pop_total))

base R using order() function

df_event_br1 <- df_event[order(df_event$event_date, df_event$event_state, -df_event$pop_total), ]
```