

# Data management with R

# What we have covered

- Data management with Python
  - List/Dictionary/Pandas series/dataframe
  - Filtering
  - Grouping
  - Joining
  - Datetime management
- Introduction of machine learning
  - Supervised learning/unsupervised learning
  - Data mining process
  - Decision tree algorithm
  - Clustering algorithm

# Agenda

- What is R and Why R
- Vectorization
  - Recycling
- Getting help
- Workspace
- Working with packages
- Working with different types of data in R
  - Dealing with characters
  - Dealing with factors
  - Dealing with dates
- Working with different data structures in R
- Different basic looping functions in R (`apply()`, `lapply()`, `sapply()`, `tapply()`)
- Importing and exporting data from/into other formats (e.g. csv and excel)
- Other mathematical functions
- Case study

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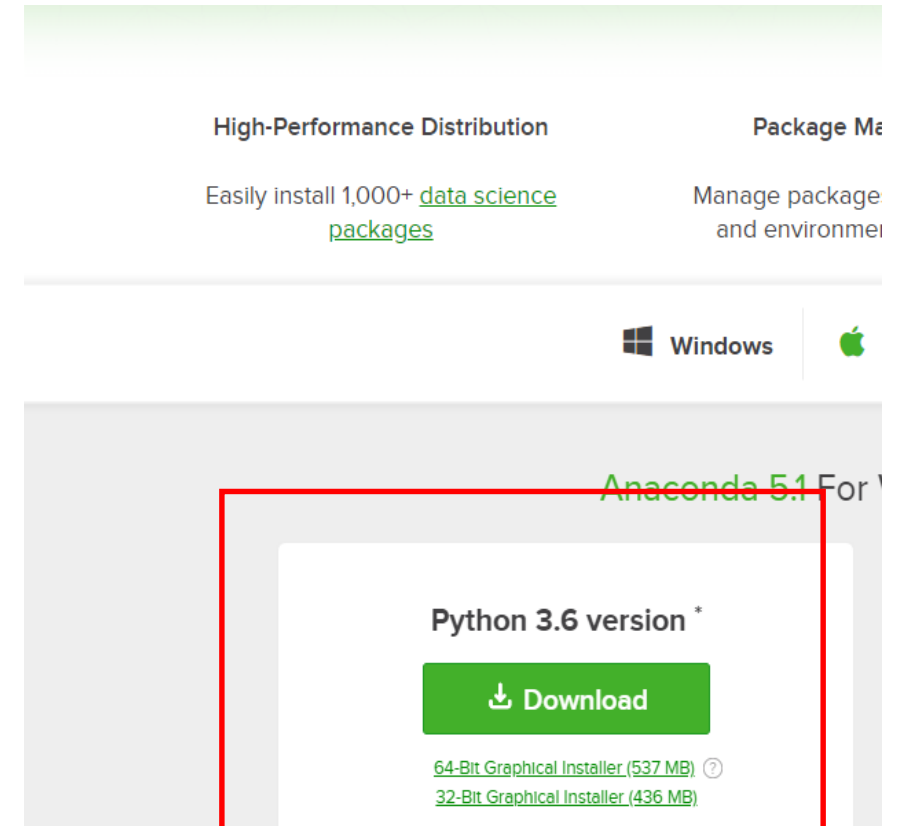
# What is R

- R is a comprehensive statistical platform, offering all manner of data-analytic techniques. Just about any type of data analysis can be done in R
- R contains advanced statistical routines not yet available in other packages
- R has the state-of-art graphics capability. If you want to visualize complex data, R has the most comprehensive and power feature set available.
- <http://cran.r-project.org>
- R studio (<https://www.rstudio.com/>) is a good GUI platform with R



# Install Jupyter

- <https://www.anaconda.com/download/>
- `conda install -c r r-essentials`



# Why R

- Open source
- Flexible
- It comes with lots of useful modules which support different topics on data analytic
- Lots of good supporting material available
- It comes with very good features on the following items:
  - Data visualization support (e.g. ggplot2)
  - Reporting feature (e.g. Rmarkup)
  - Data management
  - Data analytic

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# Vectorization

```
X<-c(1,2,3)
```

```
Y<-c(4,5,6)
```

```
Z<-X+Y
```

```
Z
```

+ is a vector operation in R, and it will operate on the entire vectors at once

# Recycling

- When performing an operation on two or more vectors of unequal length, R will recycle elements of the shorter vector( s) to match the longest vector.

## Example

```
long <- 1: 10
```

```
short <- 1: 5
```

```
long ## [1] 1 2 3 4 5 6 7 8 9 10
```

```
short ## [1] 1 2 3 4 5
```

```
long + short ## [1] 2 4 6 8 10 7 9 11 13 15
```

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# Getting help

- To get the supporting material on the function `functionname` which is already loaded

`Help(functionname)` or

`?functionname`

- To get the supporting material on the function `functionname` which has been installed but not yet loaded

`help( functionname, package = "packagename")`.

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# Workspace

- To define the workspace use `setwd()` function
- To fetch the path of the workspace use `getwd()` function
- Note the forward slashes in the pathname of the `setwd()` command. R treats the backslash (`\`) as an escape character. Even when you are using R on a Windows platform, use forward slashes in pathnames.
- Example

```
Setwd("d:/test")
```

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# Working with packages

- Install package
  - `install.packages("packagename")`
- Load package
  - `library(packagename)`
- Get help on the use of the package
  - E.g. `help(package="caret")`
- Check which library packages have been installed
  - `library()`
- Check packages already loaded
  - `search()`
- List vignettes available for a specified package
  - `Vignette(package="packagename")`
- View specific vignette
  - E.g. `Vignette("caret")`



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# Working with different types of data in R

- Integer
- Numeric
- Character
- logical

# Check the type

- `is.numeric`
- `is.character`
- `is.logical`
- `as.numeric`
- `as.character`
- `as.logical`



To check if the data type is of type numeric, character or logical respectively

To convert the data type to the type numeric, character or logical respectively

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# Dealing with character

- Create string
- Counting number of elements or characters
- Subset string
- Set operator for character strings
- Regular expression
- Split string

# Create string-1/2

- Use Paste() function

## Example 1

```
a <- "learning to create" # create string a
```

```
b <- "character strings" # create string b
```

```
paste(a,b)
```

## Example 2

```
paste(" I", "love", "R", sep = "-")
```

```
## [1] "I-love-R"
```

## Create string-2/2

- # use paste0() to paste without spaces btwn characters

```
paste0(" I", "love", "R") ## [1] "IloveR"
```

```
# paste objects with different lengths
```

```
paste(" R", 1: 5, sep = " v1.")
```

```
## [1] "R v1.1" "R v1.2" "R v1.3" "R v1.4" "R v1.5"
```

# Counting number of elements or characters

- To count the number of elements in a string use `length()`:

```
length(" How many elements are in this string?")
```

```
## [1] 1
```

```
length( c(" How", "many", "elements", "are", "in", "this", "string?"))
```

```
## [1] 7
```

- To count the number of characters in a string use `nchar()`:

```
nchar(" How many characters are in this string?")
```

```
## [1] 39
```

```
nchar( c("How", "many", "characters", "are", "in", "this", "string?"))
```

```
## [1] 3 4 10 3 2 4 7
```



# Subset string

```
alphabet <- paste( LETTERS, collapse = "")
```

```
# extract 18th through last character
```

```
substring( alphabet, first = 18)
```

```
## [1] "RSTUVWXYZ"
```

```
# recursive extraction; specify start position only
```

```
substring( alphabet, first = 18: 24)
```

```
## [1] "RSTUVWXYZ" "STUVWXYZ" "TUVWXYZ" "UVWXYZ" "VWXYZ" "WXYZ"
```

```
## [7] "XYZ"
```

**Quiz:** What is the output from the following statement ?

```
substring( alphabet, first = 1: 5, last = 3: 7)
```

# Set operator for character strings-1/6

- Set Union
- Set Intersection
- Identifying different elements
- Test for element equality
- Test for exact equality
- Testing if elements are contained in a string
- Sorting a string

# Set operator for character strings-2/6

```
set_1 <- c("lagunitas", "bells", "dogfish", "summit", "odell")
set_2 <- c("sierra", "bells", "harpoon", "lagunitas", "founders")
#Set Union
union(set_1, set_2)
#Set Intersect
intersect(set_1, set_2)
# returns elements in set_1 not in set_2
setdiff(set_1, set_2)
## [1] "dogfish" "summit" "odell"
# returns elements in set_2 not in set_1
setdiff(set_2, set_1)
## [1] "sierra" "harpoon" "founders"
```

# Set operator for character strings-3/6

To test if two vectors contain the same elements regardless of order use `setequal()`:

```
set_3 <- c(" woody", "buzz", "rex")  
set_4 <- c(" woody", "andy", "buzz")  
set_5 <- c(" andy", "buzz", "woody")  
setequal( set_3, set_4)  
## [1] FALSE  
setequal( set_4, set_5) ## [1] TRUE
```

# Set operator for character strings-4/6

To test if two character vectors are equal in content and order use `identical()`:

```
set_6 <- c(" woody", "andy", "buzz")
```

```
set_7 <- c(" andy", "buzz", "woody")
```

```
set_8 <- c(" woody", "andy", "buzz")
```

```
identical( set_6, set_7)
```

```
## [1] FALSE
```

```
identical( set_6, set_8)
```

```
## [1] TRUE
```

# Set operator for character strings-5/6

To test if an element is contained within a character vector use `is.element()` or `%in%`:

```
good <- "andy"
```

```
bad <- "sid"
```

```
is.element( good, set_8)
```

```
## [1] TRUE
```

```
good %in% set_8
```

```
## [1] TRUE
```

**Quiz:** what is the output of the following statement

```
Bad %in% set_8
```

# Set operator for character strings-6/6

To sort a character vector use sort():

```
sort( set_8)
```

```
## [1] "andy" "buzz" "woody"
```

```
sort( set_8, decreasing = TRUE)
```

```
## [1] "woody" "buzz" "andy"
```

# Regular expression-1/2

```
# use the built in data set `state.division`  
head( as.character( state.division))  
## [1] "East South Central" "Pacific" "Mountain" ## [4] "West South  
Central" "Pacific" "Mountain"  
# find the elements which match the pattern  
grep(" North", state.division)  
## [1] 13 14 15 16 22 23 25 27 34 35 41 49  
# use 'value = TRUE' to show the element value  
grep(" North", state.division, value = TRUE)
```



# Regular expression-2/2

# can use the 'invert' argument to show the non-matching elements  
grep(" North | South", state.division, invert = TRUE)

```
## [1] 2 3 5 6 7 8 9 10 11 12 19 20 21 26 28 29 30 31 32 33 37 38 39 ##  
[24] 40 44 45 46 47 48 50
```

# String splitting

To split the elements of a character string use `str_split()`.

```
z <- "The day after I will take a break and drink a beer."
```

```
str_split( z, pattern = " ") ## [[ 1]] ## [1] "The" "day" "after" "I" "will" "take"
"a" "break" ## [9] "and" "drink" "a" "beer."
```

```
a <- "Alabama-Alaska-Arizona-Arkansas-California"
```

```
str_split( a, pattern = "-")
```

```
## [[ 1]] ## [1] "Alabama" "Alaska" "Arizona" "Arkansas" "California"
```

Note that the output of `strs_plit()` is a list. To convert the output to a simple atomic vector simply wrap in `unlist()`:

```
unlist( str_split( a, pattern = "-"))
```

```
## [1] "Alabama" "Alaska" "Arizona" "Arkansas" "California"
```

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# Dealing with Factors

- Factors are variables in R which take on a limited number of different values; such variables are often referred to as categorical variables.
- One of the most important uses of factors is in statistical modeling; since categorical variables enter into statistical models such as `lm` and `glm` differently than continuous variables, storing data as factors insures that the modeling functions will treat such data correctly.
- One can think of a factor as an integer vector where each integer has a label. In fact, factors are built on top of integer vectors using two attributes: the `class()` “factor”, which makes them behave differently from regular integer vectors, and the `levels()`, which defines the set of allowed values.

# Dealing with Factors

- Factor objects can be created with the `factor()` function:

```
# create a factor string
```

```
gender <- factor( c(" male", "female", "female", "male", "female"))
```

```
gender
```

```
## [1] male female female male female
```

```
## Levels: female male # inspect to see if it is a factor class
```

```
class( gender)
```

```
## [1] "factor"
```

```
# show that factors are just built on top of integers
```

```
typeof( gender)
```

```
## [1] "integer"
```

# Dealing with Factors

- # See the underlying representation of factor

```
unclass( gender)
```

```
## [1] 2 1 1 2 1
```

```
## attr(," levels")
```

```
## [1] "female" "male"
```

```
# what are the factor levels?
```

```
levels( gender)
```

```
## [1] "female" "male"
```

```
# show summary of counts
```

```
summary( gender)
```

```
## female male
```

```
## 3 2
```

# Ordering levels

- When creating a factor we can control the ordering of the levels by using the levels argument:
- # when not specified the default puts order as alphabetical

```
gender <- factor( c(" male", "female", "female", "male", "female"))
```

```
gender
```

```
## [1] male female female male female
```

```
## Levels: female male
```

```
# specifying order
```

```
gender <- factor( c(" male", "female", "female", "male", "female"), levels = c(" male", "female"))
```

```
gender
```

```
## [1] male female female male female
```

```
## Levels: male female
```

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# Dealing with Dates

To convert a string that is already in a date format (YYYY-MM-DD) into a date object use `as.Date()`:

```
x <- c(" 2015-07-01", "2015-08-01", "2015-09-01")
```

```
as.Date( x)
```

```
## [1] "2015-07-01" "2015-08-01" "2015-09-01"
```

# Dealing with Dates

Note that the default date format is YYYY-MM-DD; therefore, if your string is of different format you must incorporate the format argument. There are multiple formats that dates can be in; for a complete list of formatting code options in R type `?strptime` in your console.

```
y <- c(" 07/ 01/ 2015", "07/ 01/ 2015", "07/ 01/ 2015")
as.Date( y, format = "% m/% d/% Y")
## [1] "2015-07-01" "2015-07-01" "2015-07-01"
```

# Dealing with Dates: Date format

Symbol	Meaning	Example
%d	Day as a number (0-31)	01-31
%a	Abbreviated weekday	Mon
%A	Unabbreviated weekday	Monday
%m	Month(00-12)	00-12
%b	Abbreviated month	Jan
%B	Unabbreviated month	January
%y	Two-digit year	07
%Y	Four-digit year	2007

In R the default format for date value is yyyy-mm-dd

# Dealing with Dates: Example on date management in R

```
strDates <- c("01/05/1965", "08/16/1975")
```

```
dates <- as.Date(strDates, "%m/%d/%Y")
```

```
today <- Sys.Date()
```

```
format(today, format="%B %d %Y")
```

```
format(today, format="%A")
```

# Dealing with Dates

To create a sequence of dates we can leverage the `seq()` function.

As with numeric vectors, you have to specify at least three of the four arguments (from, to, by, and length.out).

```
seq( as.Date(" 2010-1-1"), as.Date(" 2015-1-1"), by = "years")
```

```
seq( as.Date(' 2015-09-15'), as.Date(' 2015-09-30'), by = "2 days")
```

```
seq( as.POSIXct(" 2015-1-1 0: 00"), as.POSIXct(" 2015-1-1 12: 00"), by =  
"hour")
```

# Using lubridate package to work with dates- 1/3

```
x <- c(" 2015-07-01", "2015-08-01", "2015-09-01")
```

```
y <- c(" 07/ 01/ 2015", "07/ 01/ 2015", "07/ 01/ 2015")
```

```
library( lubridate)
```

```
ymd( x)
```

```
## [1] "2015-07-01 UTC" "2015-08-01 UTC" "2015-09-01 UTC"
```

```
mdy( y) ## [1] "2015-07-01 UTC" "2015-07-01 UTC" "2015-07-01 UTC"
```

# Using lubridate package to work with dates- 2/3

```
x <- c(" 2015-07-01", "2015-08-01", "2015-09-01")
```

```
year( x)
```

```
## [1] 2015 2015 2015 # default is numerical value
```

```
month( x)
```

```
## [1] 7 8 9
```

```
# show abbreviated name
```

```
month( x, label = TRUE)
```

```
## [1] Jul Aug Sep ## 12 Levels: Jan < Feb < Mar < Apr < May < Jun < Jul < Aug < Sep < ... < Dec
```

```
# show unabbreviated name
```

```
month( x, label = TRUE, abbr = FALSE)
```

```
## [1] July August September ## 12 Levels: January < February < March < April < May < June < ... < December
```

```
wday( x, label = TRUE, abbr = FALSE)
```

```
## [1] Wednesday Saturday Tuesday ## 7 Levels: Sunday < Monday < Tuesday < Wednesday < Thursday < ... < Saturday
```

# Using lubridate package to work with dates- 3/3

```
# most recent daylight savings time
```

```
ds <- ymd_hms(" 2015-03-08 01: 59: 59", tz = "US/ Eastern")
```

```
# if we add a duration of 1 sec we gain an extra hour
```

```
ds + dseconds( 1)
```

```
## [1] "2015-03-08 03: 00: 00 EDT"
```



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# Data structure in R

- Vector
- List
- Matrix
- Data frames

# Basic data management technique

- Checking the structure of the data
  - Use `str()` function
- Adding elements into the data structure
- Adding attributes into the data structure
  - To check the existing attributes use `attributes()` function
- Subset the data structure
- Preserving or simplifying the data structure

# Adding attributes in vector

```
# assigning names to a pre-existing vector
names( v1) <- letters[ 1: length( v1)]
V1
## a b c d e f g h i j
## 8 9 10 11 12 13 14 15 16 17
attributes( v1)
## $ names ## [1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j"
# adding names when creating vectors
v2 <- c( name1 = 1, name2 = 2, name3 = 3)
v2
## name1 name2 name3
## 1 2 3
attributes( v2) ## $ names ## [1] "name1" "name2" "name3"
```

# Subsetting with vector

- The four main ways to subset a vector include combining square brackets [ ] with:
  - Positive integers
  - Negative integers
  - Logical values
  - Names

# Subsetting with vector

```
v1
## a b c d e f g h i j
## 8 9 10 11 12 13 14 15 16 17
v1[ 2]
## b
## 9
v1[ 2: 4]
## b c d
## 9 10 11
v1[ c( 2, 4, 6, 8)]
## b d f h
## 9 11 13 15
# note that you can duplicate index positions
v1[ c( 2, 2, 4)]
## b b d
## 9 9 11
```

# Subsetting with vector

```
v1[-1]
```

```
## b c d e f g h i j
```

```
## 9 10 11 12 13 14 15 16 17
```

```
v1[-c( 2, 4, 6, 8)]
```

```
## a c e g i j
```

```
## 8 10 12 14 16 17
```

# Creating List-Example

```
l <- list( 1: 3, "a", c( TRUE, FALSE, TRUE), c( 2.5, 4.2))
```

```
str( l)
```

```
## List of 4
```

```
## $ : int [1: 3] 1 2 3
```

```
## $ : chr "a"
```

```
## $ : logi [1: 3] TRUE FALSE TRUE
```

```
## $ : num [1: 2] 2.5 4.2
```

```
# a list containing a list
```

```
l <- list( 1: 3, list( letters[ 1: 5], c( TRUE, FALSE, TRUE)))
```

```
str( l)
```

```
## List of 2
```

```
## $ : int [1: 3] 1 2 3
```

```
## $ :List of 2
```

```
## .. $ : chr [1: 5] "a" "b" "c" "d" ...
```

```
## .. $ : logi [1: 3] TRUE FALSE TRUE
```



# Adding elements into List-1/2

How to add the element vector into the existing list ?

```
l2 <- list( l1, c( 2.5, 4.2))
```

```
str( l2)
```

```
## List of 2
```

```
## $ :List of 3
```

```
## .. $ : int [1: 3] 1 2 3
```

```
## .. $ : chr "a"
```

```
## .. $ : logi [1: 3] TRUE FALSE TRUE
```



Correct ?

# Adding elements into List-2/2

```
l3 <- append( l1, list( c( 2.5, 4.2)))
```

```
str( l3)
```

```
## List of 4
```

```
## $ : int [1: 3] 1 2 3
```

```
## $ : chr "a"
```

```
## $ : logi [1: 3] TRUE FALSE TRUE
```

```
## $ : num [1: 2] 2.5 4.2
```

Alternatively, we can also add a new list component by utilizing the '\$' sign and naming the new item:

```
l3 $ item4 <- "new list item"
```

```
str( l3)
```

```
## List of 5
```

```
## $ : int [1: 3] 1 2 3
```

```
## $ : chr "a" ## $ : logi [1: 3] TRUE FALSE TRUE ## $ : num [1: 2] 2.5 4.2 ## $ item4: chr "new list item"
```

# Adding attributes to list-1/5

```
# adding names to a pre-existing list
```

```
names(l1) <- c("item1", "item2", "item3")
```

```
str(l1)
```

```
## List of 3
```

```
## $ item1: int [1: 6] 1 2 3 4 5 6
```

```
## $ item2: chr [1: 4] "a" "dding" "to a" "list"
```

```
## $ item3: logi [1: 3] TRUE FALSE TRUE attributes( l1)
```

```
## $ names ## [1] "item1" "item2" "item3"
```

```
# adding names when creating lists
```

```
l2 <- list( item1 = 1: 3, item2 = letters[ 1: 5], item3 = c( T, F, T, T))
```

```
str(l2)
```

```
## List of 3
```

```
## $ item1: int [1: 3] 1 2 3
```

```
## $ item2: chr [1: 5] "a" "b" "c" "d" ...
```

```
## $ item3: logi [1: 4] TRUE FALSE TRUE TRUE attributes( l2)
```

```
## $ names ## [1] "item1" "item2" "item3"
```

# To subset the list-2/5

- To subset lists we can utilize the single bracket [ ], double brackets [[ ]], and dollar sign \$ operators.
- Each approach provides a specific purpose and can be combined in different ways to achieve the following subsetting objectives:
  - Subset list and preserve output as a list
  - Subset list and simplify output
  - Subset list to get elements out of a list
  - Subset list with a nested list

# To subset the list-3/5

```
# extract first list item
```

```
l2[ 1]
```

```
## $ item1
```

```
## [1] 1 2 3
```

```
# same as above but using the item's name
```

```
l2[" item1"]
```

```
## $ item1
```

```
## [1] 1 2 3
```

```
# extract multiple list items l2[ c( 1,3)]
```

```
## $ item1
```

```
## [1] 1 2 3
```

```
## ## $ item3
```

```
## [1] TRUE FALSE TRUE TRUE
```

# To subset the list-4/5

- To extract one or more list items while simplifying the output use the `[[ ]]` or `$` operator:

# extract first list item and simplify to a vector

```
l2[[ 1]]
```

```
## [1] 1 2 3
```

# same as above but using the item's name

```
l2[[" item1"]]
```

```
## [1] 1 2 3
```

# same as above but using the ``$`` operator

```
l2 $ item1
```

```
## [1] 1 2 3
```

# To subset the list-5/5

- To extract individual elements out of a specific list item combine the [[ (or \$) operator with the [ operator:

# extract third element from the second list item

```
l2[[ 2]][ 3]
```

```
## [1] "c"
```

# same as above but using the item's name

```
l2[[" item2"]][ 3]
```

```
## [1] "c"
```

# same as above but using the ` \$ ` operator

```
l2 $ item2[ 3]
```

```
## [1] "c"
```

# Dealing with Matrix->Create matrix

- # numeric matrix

```
m1 <- matrix( 1: 6, nrow = 2, ncol = 3)
```

```
m1
```

```
##      [, 1] [, 2] [, 3]
```

```
##[1,] 1      3      5
```

```
## [2,] 2      4      6
```



# Adding row or column in matrix

```
v1 <- 1:4
```

```
v2 <- 5:8
```

```
cbind( v1, v2)
```

```
##      v1 v2
```

```
## [1,] 1 5
```

```
## [2,] 2 6
```

```
## [3,] 3 7
```

```
## [4,] 4 8
```

```
rbind( v1, v2)
```

```
##      [, 1] [, 2] [, 3] [, 4]
```

```
## v1 1 2 3 4
```

```
## v2 5 6 7 8
```

# Adding attributes in matrix

```
# add column names
```

```
colnames( m2) <- c(" col1", "col2", "col3")
```

```
# add row names
```

```
rownames( m2) <- c(" row1", "row2", "row3", "row4")
```

# To subset matrix-1/3

- To subset matrices we use the [ operator;
- however, since matrices have 2 dimensions we need to incorporate subsetting arguments for both row and column dimensions.
- A generic form of matrix subsetting looks like: `matrix[ rows, columns]`.

# To subset matrix-2/3

m2

##		col_1	col_2	col_3
## row_1	1	5	9	
## row_2	2	6	10	
## row_3	3	7	11	
## row_4	4	8	12	

# subset for rows 1 and 2 but keep all columns

m2[ 1: 2, ]

# subset for columns 1 and 3 but keep all rows

m2[ , c( 1, 3)]

# subset for both rows and columns

m2[ 1: 2, c( 1, 3)]

# To subset matrix-3/3

- Note that subsetting matrices with the [ operator will simplify the results to the lowest possible dimension.
- To avoid this you can introduce the drop = FALSE argument:

# simplifying results in a named vector

```
m2[, 2]
```

```
## row_1 row_2 row_3 row_4
```

```
## 5 6 7 8
```

# preserving results in a 4x1 matrix

```
m2[, 2, drop = FALSE]
```

```
## col_2
```

```
## row_1 5
```

```
## row_2 6
```

```
## row_3 7
```

```
## row_4 8
```

# Create data frame-1/2

```
df <- data.frame( col1 = 1: 3, col2 = c(" this", "is", "text"), col3 = c( TRUE, FALSE, TRUE), col4 = c( 2.5, 4.2, pi))
```

```
# assess the structure of a data frame
```

```
str( df)
```

```
## 'data.frame': 3 obs. of 4 variables:
```

```
## $ col1: int 1 2 3
```

```
## $ col2: Factor w/ 3 levels "is"," text"," this": 3 1 2
```

```
## $ col3: logi TRUE FALSE TRUE
```

```
## $ col4: num 2.5 4.2 3.14
```

```
# number of rows
```

```
nrow( df)
```

```
## [1] 3
```

```
# number of columns
```

```
ncol( df)
```

```
## [1] 4
```

# Create data frame-2/2

- Note how col2 in df was converted to a column of factors.
- This is because there is a default setting in `data.frame()` that converts character columns to factors.
- We can turn this off by setting the `stringsAsFactors = FALSE` argument:  

```
df <- data.frame( col1 = 1: 3, col2 = c(" this", "is", "text"), col3 = c( TRUE, FALSE, TRUE), col4 = c( 2.5, 4.2, pi), stringsAsFactors = FALSE)
```

# To subset data frame

- Data frames possess the characteristics of both lists and matrices:
- if you subset with a single vector, they behave like lists and will return the selected columns with all rows;
- if you subset with two vectors, they behave like matrices and can be subset by row and column:



# Missing value-1/5

- A common task in data analysis is dealing with missing values.
- In R, missing values are often represented by NA or some other value that represents missing values (i.e. 99).

# Missing value-2/5

```
# vector with missing data
```

```
x <- c( 1: 4, NA, 6: 7, NA)
```

```
x
```

```
## [1] 1 2 3 4 NA 6 7 NA
```

```
is.na( x)
```

```
## [1] FALSE FALSE FALSE FALSE TRUE FALSE FALSE TRUE
```

```
# data frame with missing data
```

```
df <- data.frame( col1 = c( 1: 3, NA), col2 = c(" this", NA," is", "text"), col3 = c( TRUE, FALSE, TRUE, TRUE), col4 = c( 2.5, 4.2, 3.2, NA), stringsAsFactors = FALSE)
```

```
# identify NAs in full data frame
```

```
is.na( df)
```

```
## col1 col2 col3 col4
```

```
## [1,] FALSE FALSE FALSE FALSE
```

```
## [2,] FALSE TRUE FALSE FALSE
```

```
## [3,] FALSE FALSE FALSE FALSE
```

```
## [4,] TRUE FALSE FALSE TRUE
```

```
# identify NAs in specific data frame column
```

```
is.na( df $ col4)
```

```
## [1] FALSE FALSE FALSE TRUE
```

# Missing value-3/5

- To identify the location or the number of NAs we can leverage the `which()` and `sum()` functions:

# identify location of NAs in vector

**`which( is.na( x))`**

## [1] 5 8

# identify count of NAs in data frame

**`sum( is.na( df))`**

## [1] 3

# Missing value-4/5

- # recode missing values with the mean

```
x[ is.na( x)] <- mean( x, na.rm = TRUE)
```

```
round( x, 2)
```

```
## [1] 1.00 2.00 3.00 4.00 3.83 6.00 7.00 3.83
```

```
# data frame that codes missing values as 99
```

```
df <- data.frame( col1 = c( 1: 3, 99), col2 = c( 2.5, 4.2, 99, 3.2)) # change  
99s to NAs
```

```
df[ df == 99] <- NA
```

# Missing value-5/5

- First, to find complete cases we can leverage the `complete.cases()` function which returns a logical vector identifying rows which are complete cases.

```
df[ complete.cases( df), ]
```

```
df[! complete.cases( df), ]
```

`na.omit()` would be the shorthand alternative

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  - Dealing with factors
  - Dealing with dates
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# Apply function

- The apply family consists of vectorized functions which minimize your need to explicitly create loops.
- These functions will apply a specified function to a data object and their primary difference is in the object class in which the function is applied to (list vs. matrix, etc) and the object class that will be returned from the function.
- `apply()` (Input:matrix or dataframe)
- `lapply()` (input:list Output:list)
- `sapply()`
- `tapply()`

# Apply()

- The syntax for `apply()` is as follows

**`apply( x, MARGIN, FUN, ...)`**

- where `x` is the matrix, dataframe
- array `MARGIN` is a vector giving the subscripts which the function will be applied over. E.g., for a matrix `1` indicates rows, `2` indicates columns, `c( 1, 2)` indicates rows and columns.
- `FUN` is the function to be applied
- `...` is for any other arguments to be passed to the function



# Apply()-Example

#To calculate the mean for every column of the dataset mtcars

```
apply( mtcars, 2, mean)
```

# get column quantiles (notice the quantile percents as row names)

```
apply( mtcars, 2, quantile, probs = c( 0.10, 0.25, 0.50, 0.75, 0.90))
```

# lapply()

- The lapply() function does the following simple series of operations:
- it loops over a list, iterating over each element in that list it applies a function to each element of the list (a function that you specify) and returns a list (the l is for “list”).
- The syntax for lapply() is as follows
  - where x is the list FUN is the function to be applied
  - ... is for any other arguments to be passed to the function
- syntax of lapply function

lapply( x, FUN, ...)

# lapply-Example

```
data <- list( item1 = 1: 4, item2 = rnorm( 10), item3 = rnorm( 20, 1),  
item4 = rnorm( 100, 5))
```

```
# get the mean of each list item
```

```
lapply( data, mean)
```

# lapply-Example

```
# list of R's built in beaver data
```

```
beaver_data <- list( beaver1 = beaver1, beaver2 = beaver2)
```

```
# get the mean of each list item
```

```
lapply( beaver_data, function( x) round( apply( x, 2, mean), 2))
```

```
## $ beaver1
```

```
## day time temp activ
```

```
## 346.20 1312.02 36.86 0.05
```

```
## ## $ beaver2
```

```
## day time temp activ
```

```
## 307.13 1446.20 37.60 0.62
```

# supply()

- The supply() function behaves similarly to lapply();
- the only real difference is in the return value.
- supply() will try to simplify the result of lapply() if possible.
- Essentially, supply() calls lapply() on its input and then applies the following algorithm:
- If the result is a list where every element is length 1, then a vector is returned
- If the result is a list where every element is a vector of the same length (> 1), a matrix is returned.
- If neither of the above simplifications can be performed then a list is returned

# tapply()

- tapply() is used to apply a function over subsets of a vector.
- It is primarily used when we have the following circumstances:
- A dataset that can be broken up into groups (via categorical variables - aka factors)
- We desire to break the dataset up into groups Within each group, we want to apply a function

# tapply()

- The arguments to tapply() are as follows:
- x is a vector
- INDEX is a factor or a list of factors (or else they are coerced to factors)  
FUN is a function to be applied
- ... contains other arguments to be passed FUN
- simplify, should we simplify the result?
- syntax of tapply function

```
tapply( x, INDEX, FUN, ..., simplify = TRUE)
```

# tapply()-Example

- `tapply( mtcars $ mpg, mtcars $ cyl, mean)`
- `apply( mtcars, 2, function( x) tapply( x, mtcars $ cyl, mean))`



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# Importing data from a delimited text file-1/2

```
# First, save the following 4 lines in a file named
# "studentgrades.csv" in the current working directory
  StudentID,First,Last,Math,Science,Social Studies
  011,Bob,Smith,90,80,67
  012,Jane,Weary,75,,80
  010,Dan,"Thornton, III",65,75,70
  040,Mary,"O'Leary",90,95,92

# Next, read the data into R using the read.table() function
grades <- read.table("studentgrades.csv", header=TRUE,
                     row.names="StudentID", sep=",")
grades # print data frame
str(grades) # view data frame structure
```

# Importing data from a delimited text file-2/2

# Alternatively, import the data while specifying column classes

```
grades <- read.table("studentgrades.csv", header=TRUE,  
                    row.names="StudentID", sep=";",  
                    colClasses=c("character", "character", "character",  
                                "numeric", "numeric", "numeric"))
```

```
grades # print data frame
```

```
str(grades) # view data frame structure
```

# Importing/exporting data from/to Excel

```
library(xlsx)
```

```
workbook <- "C:/myworkbook.xlsx"
```

```
mydataframe <- read.xlsx(workbook,1)
```

```
write.xlsx(workbook,"test.xlsx")
```

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# Mathematical functions in R

## Common mathematical functions used

Function	Description
<code>abs(x)</code>	Absolute values <code>abs(-4)</code> is 4
<code>signif(x,digits=n)</code>	Rounds x to the specified number of decimal digits <code>signif(3.475,digits=2)</code> returns 3.5
<code>mean(x)</code>	The mean value of a vector <code>Mean(c(1,2,3,4))</code> returns 2.5
<code>median(x)</code>	The median value of a vector <code>median(c(1,2,3,4))</code> returns 2.5
<code>Sd(x)</code>	The standard deviation of a vector <code>Sd(c(1,2,3,4))</code> returns 1.29
<code>Quantile(x,probs)</code>	Quantiles where x is the numeric vector, where quantiles are desired and probs is a numeric vector with probabilities in [0,1] #75 <sup>th</sup> and 25 <sup>th</sup> percentiles of x <code>Y &lt;- quantile(x, c(0.3,0.84))</code>
<code>Range(x)</code>	The range of the numeric values within the numeric vector <code>X &lt;- c(1,2,3,4)</code> <code>Range(x)</code> returns <code>c(1,4)</code> <code>Diff(range(x))</code> returns 3
<code>Scale(x,center=TRUE,scale=TRUE)</code>	Column center (center=TRUE) or standardize (center=TRUE,scale=TRUE) data object x

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# Case study

- In this example you are studying how men and women differ in the ways they lead their organizations.
- Typical questions might be:
  - Do men and women in management positions differ in the degree to which they defer to superiors ?
  - Does this vary from country to country, or are these gender differences universal
- One way to address these questions is to have bosses in multiple countries rate their managers on deferential behavior



# Sample dataset used in this example

Manager	Date	Country	Gender	Age	q1	q2	q3	q4	q5
1	10/24/14	US	M	32	5	4	5	5	5
2	10/28/14	US	F	45	3	5	2	5	5
3	10/01/14	UK	F	25	3	5	5	5	2
4	10/12/14	UK	M	39	3	3	4		
5	05/01/14	UK	F	99	2	2	1	2	1

# Requirements on the data processing

- The five ratings (q1 to q5) need to be combined, yielding a single mean deferential score from each manager
- In surveys, respondents often skip questions. For example, the boss rating manager 4 skipped questions 4 and 5. You need a method of handling incomplete data. You also need to recode values like 99 for age to missing
- There may be hundreds of variables in a dataset, but you may only be interested in a few. To simplify matters you will want to create a new dataset with only the variables of interest
- Past research suggests that leadership behavior may change as a function of the managers' age. To examine this you may want to recode the current values of ages into a new categorical age grouping (for example, young, middle-aged, elder)
- Leadership behavior may change over time. You might want to focus on deferential behavior during the recent global financial crisis. To do so, you may want to focus on deferential behavior during the recent global financial crisis. You may want to limit the study to data gathered during a specific period of time (say, January 1, 2009 to December 31, 2009)

# Create the dataframe

```
manager <- c(1,2,3,4,5)
date <- c("10/24/08","10/28/08","10/1/08","10/12/08","5/1/09")
gender <- c("M","F","F","M","F")
age <- c(32,45,25,39,99)
q1 <- c(5,3,3,3,2)
q2 <- c(4,5,5,3,2)
q3 <- c(5,2,5,4,1)
q4 <- c(5,5,5,NA,2)
q5 <- c(5,5,2,NA,1)
leadership <- data.frame(manager,date,gender,age,q1,q2,q3,q4,q5,
                          stringsAsFactors=FALSE)
```

# Creating new attributes

## Three methods are available

Example

```
mydata<-data.frame(x1 = c(2, 2, 6, 4),  
                  x2 = c(3, 4, 2, 8))
```

### Method 1

```
mydata$sumx <- mydata$x1 + mydata$x2  
mydata$meanx <- (mydata$x1 + mydata$x2)/2
```

### Method 2

```
attach(mydata)  
mydata$sumx <- x1 + x2  
mydata$meanx <- (x1 + x2)/2  
detach(mydata)
```

# Missing values

- In R, missing values are represented by the symbol **NA**

## 1. **Checking if it is missing**

- Use `is.na()` function to check if it is NA value

## 2. **Recoding values into missing**

Example:

```
leadership$age[age==99] <-NA
```

## 3. **Excluding missing values from analysis**

- Using “`na.rm=TRUE`”

Example:

```
x <- c(1, 2, NA, 3)
```

```
y <- sum(x, na.rm=TRUE)
```

# Example: Create new categorical attribute

```
leadership <- within(leadership,{  
  agecat <- NA  
  agecat[age > 75] <- "Elder"  
  agecat[age >= 55 & age <= 75] <- "Middle Aged"  
  agecat[age < 55] <- "Young" })
```