Basics of Data Frames STAT 133

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Data Frames

Data Frame

A data.frame is the primary data structure that R provides for handling tabular data sets (eg spreadsheet like).

Function data.frame()

The data.frame() function allows us to create data frames

Creating a Data Frame

```
# data frame
df <- data.frame(
  name = c('Anakin', 'Padme', 'Luke', 'Leia'),
  gender = c('male', 'female', 'male', 'female'),
  height = c(1.88, 1.65, 1.72, 1.50),
  weight = c(84, 45, 77, 49)
)</pre>
```

by default, data.frame() converts strings into factors

Simple data frame df

```
## name gender height weight
## 1 Anakin male 1.88 84
## 2 Padme female 1.65 45
## 3 Luke male 1.72 77
## 4 Leia female 1.50 49
```

Inspecting Data Frames

- dimensions (number of rows and columns)
- type of data in each column
- row names and column names
- missing data
- overall summary of each variable

Overall structure

A summary of the structure can be obtained with str()

```
# structure of a data frame
str(df)

## 'data.frame': 4 obs. of 4 variables:
## $ name : Factor w/ 4 levels "Anakin","Leia",..: 1 4 3 2
## $ gender: Factor w/ 2 levels "female","male": 2 1 2 1
## $ height: num 1.88 1.65 1.72 1.5
## $ weight: num 84 45 77 49
```

Function str()

str() applied on data frames provides:

- number of rows
- number of variables
- name of each column
- ▶ mode (i.e. type) of each column (e.g. num, int, chr, factor)
- number of levels for factor variables

str() is good for visual inspection, but doesn't give you direct access to the displayed information.

Basic Information of Data Frames

| Function | Description |
|-----------------------|-------------------------------------|
| dim() | dimensions (rows and columns) |
| nrow() | number of rows |
| <pre>ncol()</pre> | number of columns |
| names() | name of columns |
| <pre>colnames()</pre> | name of columns |
| rownames() | names of rows |
| dimnames() | list with names of rows and columns |

Basic Information of Data Frames

```
dim(df) # dimensions in a two element vector
## [1] 4 4
nrow(df)
## [1] 4
ncol(df)
## [1] 4
```

Basic Information of Data Frames

```
colnames(df)
## [1] "name" "gender" "height" "weight"
rownames(df)
## [1] "1" "2" "3" "4"
dimnames(df) # names in a list
## [[1]]
## [1] "1" "2" "3" "4"
##
## [[2]]
## [1] "name" "gender" "height" "weight"
```

Function object.size()

To know how much memory space is allocated for a data frame (or any other R object) we use object.size()

```
object.size(df)
## 2136 bytes
```

Functions head() and tail

Inspect the first and last rows, respectively:

```
# first 3 rows
head(df, n = 3)
## name gender height weight
## 1 Anakin
           male 1.88 84
## 2 Padme female 1.65 45
## 3 Luke male 1.72 77
# last 2 rows
tail(df, n = 2)
##
    name gender height weight
## 3 Luke male 1.72
                       77
## 4 Leia female 1.50 49
```

Function summary()

There's also the function summary(), which provides a descriptive summary of each column

```
summary(df)
              gender height weight
##
      name
## Anakin:1 female:2 Min. :1.500 Min. :45.00
## Leia :1 male :2 1st Qu.:1.613 1st Qu.:48.00
## Luke :1
                     Median :1.685
                                   Median :63.00
## Padme :1
                     Mean :1.688
                                   Mean :63.75
##
                      3rd Qu.:1.760
                                   3rd Qu.:78.75
##
                      Max. :1.880
                                   Max. :84.00
```

Elementary Manipulations

Accessing Elements

There are many different ways in which the elements of a "data.frame" can be accessed (i.e. retrieved, selected).

- accessing individual cells
- accessing sets of cells
- accessing entire rows
- accessing entire columns

Notation System Reminder

Notation system to extract values from data frames

- ▶ to extract values use brackets: []
- inside the brackets specify indices for rows and columns
- each index is separated by comma
- row indices can be numbers or logicals
- column indices can be numbers, logicals, or names

Single Cells

Using row and column indices to access a single cell

```
# first cell 1,1
df[1, 1]
## [1] Anakin
## Levels: Anakin Leia Luke Padme
# cell 3,4
df[3, 4]
## [1] 77
# last cell
df[4, 4]
## [1] 49
```

Various Cells

Using vectors of row and column indices to access various cells

```
# various adjacent cells
df[1:3, 2:4]
## gender height weight
## 1 male 1.88 84
## 2 female 1.65 45
## 3 male 1.72 77
# various adjacent cells
# (permuted order)
df [4:1, 3:2]
## height gender
## 4 1.50 female
## 3 1.72 male
## 2 1.65 female
## 1 1.88 male
```

Various Cells

Using vectors of row and column indices to access various cells

```
# non-adjacent cells
df[c(2, 4), c(1, 3)]

## name height
## 2 Padme 1.65
## 4 Leia 1.50
```

Various Cells

Using excluding indices

```
# excluding various adjacent cells
df[-c(1:2), -c(2:3)]

## name weight
## 3 Luke 77
## 4 Leia 49
```

Retrieving Rows

Selecting rows

```
# first row
df[1,]
## name gender height weight
## 1 Anakin male 1.88 84
# rows 1 to 3
df[1:3,]
## name gender height weight
## 1 Anakin male 1.88 84
## 2 Padme female 1.65 45
## 3 Luke male 1.72 77
```

Retrieving Rows

Selecting rows (by excluding indices)

```
# all rows except first one
df[-1,]
    name gender height weight
##
## 2 Padme female 1.65 45
## 3 Luke male 1.72 77
## 4 Leia female 1.50 49
# rows except 2 to 3
df[-c(2:3),]
##
      name gender height weight
## 1 Anakin male 1.88 84
## 4 Leia female 1.50 49
```

Retrieving Columns

Selecting columns

```
# 1st column (as a vector)
df[ , 1]

## [1] Anakin Padme Luke Leia
## Levels: Anakin Leia Luke Padme
```

Using argument drop=FALSE

```
# 1st column (as a column)
df[ , 1, drop = FALSE]

##     name
## 1 Anakin
## 2 Padme
## 3 Luke
## 4 Leia
```

Retrieving Columns

```
# columns 1 to 3
df[ , 1:3]
## name gender height
## 1 Anakin male 1.88
## 2 Padme female 1.65
## 3 Luke male 1.72
## 4 Leia female 1.50
# columns 2, 4
df[, c(2,4)]
## gender weight
## 1 male
          84
## 2 female 45
## 3 male 77
## 4 female 49
```

Retrieving Columns

Selecting columns by excluding indices

```
# all columns but 2, 4,
df[, -c(2,4)]

## name height
## 1 Anakin   1.88
## 2 Padme   1.65
## 3 Luke   1.72
## 4 Leia   1.50
```

Accessing Single Columns

Besides using numeric indices, we can also access a single column using its name and following different syntax options:

- ▶ df[,"name"]
- ▶ df[["name"]]
- ▶ df\$name or df\$"name"
- df["name"]

Column by Name

```
# equivalent ways to retrieve one column
df[ ,"name"]
## [1] Anakin Padme Luke Leia
## Levels: Anakin Leia Luke Padme
df[["gender"]]
## [1] male female male female
## Levels: female male
df$height
## [1] 1.88 1.65 1.72 1.50
```

Columns by Name

Retrieve various columns by name:

```
# vector of names
df[, c("name", "gender", "height")]

## name gender height
## 1 Anakin male 1.88
## 2 Padme female 1.65
## 3 Luke male 1.72
## 4 Leia female 1.50
```

Adding New Elements

A typical data frame modification consists in adding new elements, that is, new rows and columns.

Adding One Column

Adding a single column to a data frame

```
# adding 'a_vector' as a 'new' column
df$eyecolor <- c('blue', 'brown', 'blue', 'brown')

df

## name gender height weight eyecolor
## 1 Anakin male 1.88 84 blue
## 2 Padme female 1.65 45 brown
## 3 Luke male 1.72 77 blue
## 4 Leia female 1.50 49 brown</pre>
```

Adding One Column

Using column binding cbind() to add a column to a data frame

```
haircolor <- c('blond', 'brown', 'blond', 'brown')</pre>
# binding a column
df <- cbind(df, haircolor)</pre>
df
##
      name gender height weight eyecolor haircolor
## 1 Anakin male 1.88
                          84
                                 blue
                                         blond
## 2 Padme female 1.65 45
                                brown brown
## 3 Luke male 1.72 77 blue blond
## 4 Leia female 1.50 49
                                brown brown
```

Adding One Column

Remember the recycling rule:

```
# "human" will be recycled!
df$species <- "human"
df
##
      name gender height weight eyecolor haircolor species
## 1 Anakin
           male 1.88
                         84
                               blue
                                       blond
                                              human
## 2 Padme female 1.65
                         45
                                      brown human
                               brown
## 3 Luke
           male 1.72
                         77
                               blue
                                       blond human
## 4 Leia female 1.50
                         49
                               brown
                                       brown
                                              human
```

Adding Several Columns

Equivalent ways to add several columns to a data frame

```
# adding vectors x and y
df[, c("x", "y")] <- cbind(1:4, 5:8)

##    name gender height weight eyecolor haircolor species x y
## 1 Anakin male 1.88 84 blue blond human 1 5
## 2 Padme female 1.65 45 brown brown human 2 6
## 3 Luke male 1.72 77 blue blond human 3 7
## 4 Leia female 1.50 49 brown brown human 4 8</pre>
```

Adding Several Columns

Equivalent ways to add several columns to a data frame

```
# adding vectors u and v
uv <- cbind(u = 1:4, v = 5:8)

df <- cbind(df, uv)</pre>
```

Removing Columns

Removing columns with the NULL object

```
# removing x and y
df$x <- NULL
df$y <- NULL
df
     name gender height weight eyecolor haircolor species
##
## 1 Anakin
            male
                  1.88
                         84
                               hlue
                                       blond
                                              human
## 2 Padme female 1.65
                         45
                              brown
                                      brown human
## 3 Luke male 1.72
                         77
                               blue blond human
## 4 Leia female 1.50
                         49
                              brown brown
                                              human
```

Removing Columns

Removing columns by reassignment

```
# removing columns 5, 6, ...

df <- df[ , 1:4]

df

## name gender height weight

## 1 Anakin male 1.88 84

## 2 Padme female 1.65 45

## 3 Luke male 1.72 77

## 4 Leia female 1.50 49
```

- Another operation is adding rows
- ▶ This can be done with row binding rbind()
- When adding rows to a data frame, we need to take into account the mode of each column
- If all columns have the same mode, then we can add a vector
- ▶ If columns have different modes, then we need to add data.frames

Be careful when adding vector rows to data frames!

```
# new vector
newone <- c("Han", 'male', 1.8, 80)
# trying to add a vector to data frame
rbind(df, newone)
## Warning in '[<-.factor'('*tmp*', ri, value = "Han"):</pre>
invalid factor level, NA generated
##
  name gender height weight
## 1 Anakin male 1.88
                           84
## 2 Padme female 1.65 45
## 3 Luke male 1.72 77
## 4 Leia female 1.5 49
## 5 <NA> male 1.8 80
```

Since columns in df are of different modes, we must create a new row "vector" in data.frame format

```
# creating a data frame "vector"
han <- data.frame(
  name = "Han",
  gender = 'male',
  height = 1.8,
  weight = 80)
han

## name gender height weight
## 1 Han male 1.8 80</pre>
```

Use row binding rbind() to add one or more rows:

```
# adding 'han' with rbind()
df <- rbind(df, han)

df

##     name gender height weight
## 1 Anakin male    1.88    84
## 2 Padme female    1.65    45
## 3 Luke male    1.72    77
## 4 Leia female    1.50    49
## 5 Han male    1.80    80</pre>
```

Arranging Columns

A less common, but equally important type of data.frame modification involves rearranging or moving its columns.

The common approach to move columns is to define a vector with the column names in the desired order, and then redefine the current data frame.

Rearranging columns

```
# rearranging columns
df[ , c(1, 4, 3, 2)]

##    name weight height gender
## 1 Anakin     84     1.88     male
## 2 Padme     45     1.65 female
## 3 Luke     77     1.72     male
## 4 Leia     49     1.50 female
## 5 Han     80     1.80     male
```

Column Names

Changing column names

```
# change first column
names(df)[1] <- "Name"</pre>
# change weight
names(df)[4] <- "wgt"
df
##
      Name gender height wgt
## 1 Anakin male 1.88 84
## 2 Padme female 1.65 45
## 3 Luke male 1.72 77
## 4 Leia female 1.50 49
## 5 Han male 1.80 80
```

Column Names

Changing column names

```
# rename first column
names(df)[1] <- "name"</pre>
# rename weight
names(df)[4] <- "weight"</pre>
df
##
      name gender height weight
## 1 Anakin male 1.88
                           84
## 2 Padme female 1.65 45
## 3 Luke male 1.72 77
## 4 Leia female 1.50
                          49
## 5 Han male 1.80
                           80
```

Exercise

Creating new column

```
# height x weight
df$htwt <- df$height * df$weight
# gender and species
df$new <- paste(df$name, df$gender, sep = "_")</pre>
df
      name gender height weight htwt
##
                                            new
## 1 Anakin male 1.88 84 157.92 Anakin_male
## 2 Padme female 1.65 45 74.25 Padme_female
## 3 Luke male 1.72 77 132.44 Luke male
## 4 Leia female 1.50 49 73.50 Leia_female
## 5 Han male 1.80 80 144.00
                                      Han male
```

Subsetting using comparisons (logical vectors TRUE, FALSE)

```
df$name == "Luke"

## [1] FALSE FALSE TRUE FALSE FALSE

# Luke's info
df[df$name == "Luke", ]

## name gender height weight htwt new
## 3 Luke male 1.72 77 132.44 Luke_male
```

Subsetting using comparisons (logical vectors TRUE, FALSE)

```
df$gender == "male"
## [1]
      TRUE FALSE TRUE FALSE
                            TRUF.
# male subjects
df[df$gender == "male", ]
##
      name gender height weight htwt
                                          new
## 1 Anakin
            male 1.88 84 157.92 Anakin_male
## 3 Luke male 1.72 77 132.44 Luke_male
## 5 Han male 1.80 80 144.00 Han_male
```

Subsetting with composed statements

```
# male with height > 1.75
df[df$gender == "male" & df$height > 1.75, ]

## name gender height weight htwt new
## 1 Anakin male 1.88 84 157.92 Anakin_male
## 5 Han male 1.80 80 144.00 Han_male
```

Subsetting statements can become very verbose

```
# male with height > 1.75 and weight > 80

df[df$gender == "male"
   & df$height > 1.75
   & df$weight > 80, ]

## name gender height weight htwt new
## 1 Anakin male 1.88 84 157.92 Anakin_male
```

Subsetting with subset()

To reduce verbose subsetting statements we can use subset()

```
# male with height > 1.75
subset(df, gender == "male" & height > 1.75)
## name gender height weight htwt
                                           new
## 1 Anakin male 1.88 84 157.92 Anakin_male
## 5 Han male 1.80 80 144.00 Han male
# male with height > 1.75 and weight > 80
subset(df, gender == "male" & height > 1.75 & weight > 80)
## name gender height weight htwt
                                           new
## 1 Anakin male 1.88 84 157.92 Anakin_male
```

Subsetting with subset()

subset() also allows you to select columns according to a specified condition

Subsetting with subset()

subset() also allows you to select columns according to a specified condition

Ordering Rows

Remember sort() and order()

```
# sort() sorts the values
sort(df$weight)

## [1] 45 49 77 80 84

# order() gives you the position
order(df$weight)

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

When sorting rows, we want to work with the ordered positions

```
# sorting rows by weight
df[order(df$weight),]

## name gender height weight htwt new
## 2 Padme female 1.65 45 74.25 Padme_female
## 4 Leia female 1.50 49 73.50 Leia_female
## 3 Luke male 1.72 77 132.44 Luke_male
## 5 Han male 1.80 80 144.00 Han_male
## 1 Anakin male 1.88 84 157.92 Anakin_male
```

```
# sorting subjects by height
df[order(df$height), c('name', 'height')]

## name height
## 4 Leia 1.50
## 2 Padme 1.65
## 3 Luke 1.72
## 5 Han 1.80
## 1 Anakin 1.88
```

```
# sorting subjects by height in decreasing order
ht_sort <- order(df$height, decreasing = TRUE)
df[ht_sort, c('name', 'height')]

## name height
## 1 Anakin    1.88
## 5 Han    1.80
## 3 Luke    1.72
## 2 Padme    1.65
## 4 Leia    1.50</pre>
```