Chapter 16 Answers

Highlights

```
• Review these functions
```

- data.frame
- head
- tail
- summary
- transform
- subset

16.1 What is returned by the following R commands? (Waking hours from wikipedia.)

```
creatures = c("dog","cat","armadillo","human")
friendly = c(TRUE,TRUE,FALSE,TRUE)
diet = c("cats","mice","termites","Twinkies(tm)")
waking.hours = c(13.9, 11.5, 5.9, 16.0)
creature.data = data.frame (friendly, diet, waking.hours, row.names=creatures)
creature.data
```

```
##
             friendly
                               diet waking.hours
## dog
                 TRUE
                               cats
                                             13.9
## cat
                 TRUE
                               mice
                                             11.5
                                              5.9
## armadillo
                FALSE
                           termites
## human
                 TRUE Twinkies(tm)
                                             16.0
```

creatures[creature.data\$waking.hours < 12 & !creature.data\$friendly]</pre>

```
## [1] "armadillo"
```

16.2 Write a single R command that alphabetizes the rows of the data frame creature.data from Exercise 16.1 by creature name.

```
creature.data[sort(creatures),]
```

```
##
             friendly
                               diet waking.hours
## armadillo
                FALSE
                           termites
                  TRUE
                                              11.5
## cat
                               mice
                  TRUE
                                             13.9
## dog
                               cats
                 TRUE Twinkies(tm)
                                             16.0
## human
```

```
# also (perhaps the separate object creatures was not created)
creature.data[sort(row.names(creature.data)),]
```

```
##
             friendly
                               diet waking.hours
## armadillo
                FALSE
                           termites
                                              5.9
                 TRUE
                                             11.5
## cat
                               mice
## dog
                 TRUE
                                             13.9
                               cats
                 TRUE Twinkies(tm)
## human
                                             16.0
```

```
# for fun:
summary(creature.data)
```

```
##
    friendly
                              diet
                                      waking.hours
## Mode :logical
                                :1
                                     Min. : 5.90
                    cats
## FALSE:1
                    mice
                                : 1
                                     1st Qu.:10.10
## TRUE:3
                    termites
                                :1
                                     Median :12.70
```

```
##
                     Twinkies(tm):1
                                       Mean
                                               :11.82
                                       3rd Qu.:14.43
##
##
                                               :16.00
16.3 Use the R subset function to create a data frame consisting of just the creature name and diet associated
with friendly creatures who are awake more than 12 hours a day from the data drame creature.data from
Exercise 16.1.
subset(creature.data,friendly & (waking.hours > 12),c(diet))
                  diet
## dog
                  cats
## human Twinkies(tm)
16.4 Consider the data frame 'creature.data from Exercise 16.1.
  • Extract the waking hours for a dog using two different R commands
creature.data["dog","waking.hours"]
## [1] 13.9
# subset extracts a new data frame,
# adding the [1,1] indexes extracts the cell value
subset(creature.data, "dog" == row.names(creature.data), c(waking.hours))[1,1]
## [1] 13.9
  • Extract the waking hours for all creatures using two different R commands
creature.data$waking.hours
## [1] 13.9 11.5 5.9 16.0
creature.data[["waking.hours"]]
## [1] 13.9 11.5 5.9 16.0
subset(creature.data,TRUE,c(waking.hours))[,1]
## [1] 13.9 11.5 5.9 16.0
  • Create a data frame that consists only of the rows for dogs and armadillos using two different R
     commands
subset(creature.data, "dog" == row.names(creature.data) | "armadillo" == row.names(creature.data), colnam
##
             friendly
                            diet waking.hours
## dog
                  TRUE
                                          13.9
                            cats
## armadillo
                 FALSE termites
                                          5.9
subset(creature.data, "dog" == row.names(creature.data) | "armadillo" == row.names(creature.data),)
##
             friendly
                            diet waking.hours
## dog
                  TRUE
                            cats
                                          13.9
## armadillo
                 FALSE termites
                                           5.9
subset(creature.data,(waking.hours < 16 & waking.hours>12) | waking.hours<6,)</pre>
##
             friendly
                            diet waking.hours
```

• Create a data frame that consists only of friendly creatures

cats

13.9

5.9

TRUE

FALSE termites

dog

armadillo

subset(creature.data,friendly,) ## friendly diet waking.hours ## dog TRUE cats 13.9 TRUE ## cat mice 11.5 ## human TRUE Twinkies(tm) 16.0 16.5 Consider a data frame named a with 4 rows and 3 columns populated with positive integers and -1 where -1 denotes a missing value. Write an R command that replaces each -1 with NA. a = data.frame(col1=1:4,col2=seq(10,40,by=10),col3=seq(100,400,by=100)) ## col1 col2 col3 ## 1 1 10 100 2 20 200 ## 2 ## 3 3 30 300 ## 4 400 4 40 a[1,1] = a[2,2] = a[3,3] = a\$col2[4] = -1## col1 col2 col3 ## 1 -1 10 100 ## 2 2 -1 200 ## 3 3 30 -1 4 ## 4 -1 400 $a[a==-1] \leftarrow NA$ a ## col1 col2 col3 ## 1 NA10 100 ## 2 2 NA200 ## 3 3 30 NA ## 4 400 16.6 Create the objects v, m, a, 1, and d as a vector, matrix, array, list, and data frame. Apply the functions class, typeof, and mode to each of the objects and summarize the results in a table. v = 1:12m = matrix(v,3,4)a = array(v,c(2,2,3))1 = list(v,m,a)d = subset(creature.data,friendly,) st = data.frame(class(m), class(a), class(1), class(d)), class = c(class(v), c(typeof(v), typeof(m), typeof(a), typeof(1), typeof(d)), typeof = c(mode(v), mode(m), mode(a), mode(1), mode(d)), row.names = c("v", "m", "a", "1", "d")) st ## class typeof mode## v integer integer numeric ## m matrix integer numeric array integer numeric ## a

```
## l list list list
## d data.frame list list
```

16.7 Create a data frame named w that consists of the following three named columns:

- x, the first four positve integers,
- y, the abbreviations of the first four months,
- z, the first names of the Beatles.

```
## 1 1 Jan George
## 2 2 Feb John
## 3 3 Mar Paul
## 4 4 Apr Ringo
```

Remove the second column of ${\tt w}$ in the following two fashions:

- set w\$y to NULL, that is, w\$y = NULL,
- subset the columns to keep, that is, w[c("x","z")].

```
w_ = w
w_$y = NULL

w_- = w
w__ [c("x","z")]
```

```
## 1 1 George
## 2 2 John
## 3 3 Paul
## 4 4 Ringo
```

Show that the first technique alters the data frame wbut the second technique does not alter the data frame w.

```
# the first method was an assignment operation, resulting in a change \mathbf{w}_{\perp}
```

```
## x z
## 1 1 George
## 2 2 John
## 3 3 Paul
## 4 4 Ringo
```

the second method was an extraction operation, with no permanent damage done to the data frame \mathtt{W}_{--}

```
## x y z
## 1 1 Jan George
## 2 2 Feb John
## 3 3 Mar Paul
## 4 4 Apr Ringo
```