## 2. Base R - Subsetting

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## Subsetting

R's subsetting operators are fast and powerful. Mastering them allows you to succinctly perform complex operations in a way that few other languages can match.

— Hadley Wickham "Advanced R"

#### **Overview**

- Subsetting operations allow you to process data stored in atomic vectors and lists
- R provides a range of flexible approaches that can be used to subset data.
- Learning outcomes:
  - The four main ways to subset a vector, namely, by positive integer, the minus sign, logical vectors, and vector element names.
  - The role of the [[ operator for processing lists, and how to distinguish this from the [ operator.
  - The use of the \$ operator
  - How to use the for loop structure to iterate through a list
  - How to use the if statement when processing lists, and how that statement differe from the ifelse() function covered in Chapter Chapter @ref(ch2).
  - Additional R functions that allow you to process vectors
  - How to solve all three test exercises.

#### **Atomic Vectors**

- Let's model the daily number of customers arriving at a restaurant as a *Poisson distribution*, with a mean ( $\lambda$ ) of one hundred customers per day ( $\lambda$ =100).
- $\bullet$  The Poisson distribution describes a discrete random variable, and the mean and variance are both equal to  $\lambda$
- In R, the rpois() function can be used to generate random numbers from a Poisson distribution, with mean ( $\lambda$ ).

```
# set the seed to ensure replication
set.seed(111)
# Generate the count data, assume a Poisson distribution
customers <- rpois(n = 10, lambda = 100)
names(customers) <- paste0("D",1:10)
customers</pre>
```

```
D1
        Ω2
            D.3
                 D4
                     D5
                         D6
                             D7
                                  D8
                                      D9 D10
##
        96
            97
                 98 101
                         85
                             98 118 102 94
## 102
```

## **Subsetting vectors - Positive Integers**

- For vectors in R, the operator [ is used to subset vectors
- Positive integers will subset atomic vector elements at given locations
- To extract the  $n^{th}$  item from a vector x the term x[n] is used
- This can also apply to a sequence, starting at n and finising at m can be extracted from the vector x as x[n:m]
- Indices can also be generated using the combine function c(), which is then passed in to subset a vector.

```
customers[1]
## D1
## 102
customers[1:5]
## D1 D2 D3 D4 D5
```

96

## 102

97 98 101

## **Subsetting vectors - Negative Integers**

- Negative integers, expressed as a vector, can be used to exclude elements from a vector
- One or more elements can be excluded

```
customers[-1]
##
   Π2
       D.3
           D4 D5
                   D6 D7
                           D8
                              D9 D10
##
   96
       97
           98 101 85 98 118 102 94
customers[-c(1,length(customers))]
##
   D2.
       D3
           D4
              D5
                   D6
                      D7
                           D8
                               D9
           98 101
##
   96
       97
                   85
                       98 118 102
customers[-(2:(length(customers)-1))]
```

## D1 D10 ## 102 94

## **Subsetting vectors - Logical Vectors**

- Logical vectors can be used to subset a vector
- This allows for the use of relational and logical operators.
- when a logical vector is used to subset a vector, only the corresponding cells of the logical vector element that contain TRUE will be retained in the operation.

```
##
      D1
                    D3
                           D4
                                  D5
                                        D6
                                               D7
                                                      D8
                                                             D9
                                                                   D10
    TRUE FALSE FALSE FALSE TRUE FALSE FALSE
##
                                                    TRUF.
                                                           TRUE FALSI
##
    D1
        D5
             Вď
                  Д9
```

• The two statements can be combined into the one expression.

```
(customers[customers > 100])
```

```
## D1 D5 D8 D9
## 102 101 118 102
```

102 101 118 102

## Recycling

- A nice feature of subsetting with logical vectors is that the logical vector size does not have to equal the size of the target vector.
- When the length of the logical vector is less than the target vector, R will *recycle* the logical vector by repeating the sequence of values until all the target values have been subsetted.

```
# Subset every third element from the vector
customers[c(TRUE,FALSE,FALSE)]
```

```
## D1 D4 D7 D10
## 102 98 98 94
```

## **Subsetting vectors - By Element Names**

- If a vector has named elements usually set via the function names(), then elements can subsetted through their name.
- This is convenient if you want to retrieve an element but do not necessarily want to know its exact indexed location.

```
customers
       D2
           D3 D4 D5 D6 D7
                               Вď
##
   D1
                                   D9 D10
           97 98 101 85 98 118 102 94
## 102
       96
# Show the value from day 10
customers["D10"]
## D10
   94
##
customers[c("D1","D10")]
```

## **Subsetting Lists**

In a similar manner to our exploration of atomic vectors, we first generate a simulated manufacturing data for two products, A and B.

```
# A small products database. Main list has two products
products <- list(</pre>
             A=list(product="A",
                     sales=12000,
                     quarterly=list(quarter=1:4,
                                sales=c(6000,3000,2000,1000)))
             B=list(product="B",
                     sales=8000.
                     quarterly=list(quarter=1:4,
                                sales=c(2500,1500,2800,1200)))
```

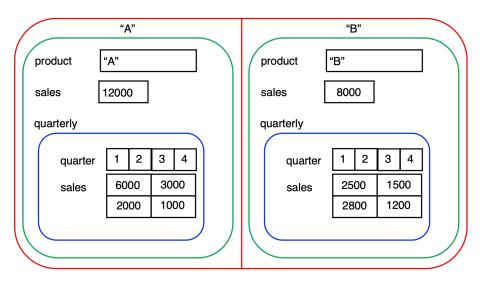
### **Exploring the list structure**

```
str(products)
## List of 2
   $ A:List of 3
##
## ..$ product : chr "A"
## ..$ sales : num 12000
## ..$ quarterly:List of 2
## ....$ quarter: int [1:4] 1 2 3 4
## ....$ sales : num [1:4] 6000 3000 2000 1000
##
   $ B:List of 3
## ..$ product : chr "B"
    ..$ sales : num 8000
##
##
    ..$ quarterly:List of 2
    ....$ quarter: int [1:4] 1 2 3 4
##
```

##

....\$ sales : num [1:4] 2500 1500 2800 1200

## Visualising the List



**Figure 1:** Visualisation of the products data structure

#### Some observations on the list

- At its core, the list is simply a vector of two named elements, and this is highlighted with the red lines.
- We can check this with R code to show (1) the length of the vector and (2) the name of each element.

```
# Show the vector length (2 elements)
length(products)

## [1] 2
# Show the names of each element
names(products)

## [1] "A" "B"
```

#### Some observations on the list

- However, even though there are just two elements in the list, each element has a significant internal structure.
- Each element contains a list, highlighted in green. This list contains three elements:
  - the product name, a character atomic vector,
  - the sales, a numeric atomic vector
  - an element named quarterly, which is another list (coloured blue).
- This third list contains two atomic vector elements:
  - the quarter number (1, 2, 3 and 4), and
  - the corresponding sales amount for each quarter, for the product.
- Note that the sum of the sales vector in this list equals the amount in the sales vector in the previous list.

## **Subsetting lists**

- Subsetting lists is more challenging than subsetting atomic vectors.
- There are three methods that can be used, and to illustrate the core idea we define a list (11) that contains three named elements.

```
# Create a simple list vector

11 <- list(a="Hello",b=1:5,c=list(d=c(T,T,F),e="Hello World"))
# Show the structure
str(11)

## List of 3
## $ a: chr "Hello"
## $ b: int [1:5] 1 2 3 4 5
## $ c:List of 2
## ..$ d: logi [1:3] TRUE TRUE FALSE</pre>
```

..\$ e: chr "Hello World"

## Subsetting with [

- The single square bracket '[', when applied to a list, will always return a list.
- The same indexing method used for atomic vectors can also be used for filtering lists, namely: positive integers, negative integers, logical vectors, and the element name. Here are examples of filtering a list using each of these methods.

```
# extract the first element of the list
str(l1[1])

## List of 1
## $ a: chr "Hello"
str(l1["a"])
```

\$ a: chr "Hello"

## List of 1

### Subsetting with [

Some additional examples:

```
# extract the first two list elements
str(11[1:2])
## List of 2
## $ a: chr "Hello"
## $ b: int [1:5] 1 2 3 4 5
str(l1[c(T,F)])
## List of 2
## $ a: chr "Hello"
## $ c:List of 2
## ..$ d: logi [1:3] TRUE TRUE FALSE
```

## ..\$ e: chr "Hello World"

## **Extracting list contents with [[**

- The single bracket [ return a list, but in many cases this is not sufficent for analysis
- We will need to access the data within the list (which can be an atomic vector, and also a list).
- For example, finding the value of element a or element b.
- To do this, we must use the '[[' operator, which extracts the *contents* of a list at a given location (i.e. element 1, 2, .., N), where N is the list length.

```
# extract the contents of the first list element
11[[1]]
## [1] "Hello"
11[["a"]]
```

## [1] "Hello"

## **Extracting list contents with** [[

```
# extract the contents of the second list element
11[["b"]]
## [1] 1 2 3 4 5
# extract the contents of the third list element (a list!)
str(l1[["c"]])
## List of 2
## $ d: logi [1:3] TRUE TRUE FALSE
## $ e: chr "Hello World"
11[["c"]][["d"]]
## [1] TRUE TRUE FALSE
11[[3]][[1]]
```

## [1] TRUE TRUE FALSE

### **Extracting list contents with \$**

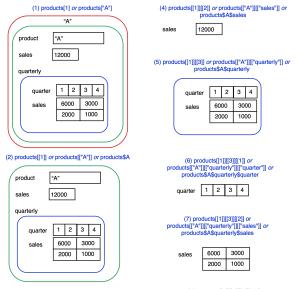
- There is a convenient alternative to the [[ operator, and this is the tag operator \$ which can be used once a list element is named.
- For example, for our list 11 the terms 11[[1]], 11[["a"] and 11\$a are the same, and in the general case 1[["y]] is equivalent to 1\$y

```
## [1] 1 2 3 4 5
# extract the contents of the third list element (a list!)
str(l1$c)
## List of 2
## $ d: logi [1:3] TRUE TRUE FALSE
## $ e: chr "Hello World"
```

11\$b

11\$c\$d

### Visualising subsets of products



# Code for (1) - get the first list element

```
# Example (1) - get the first element of the list as a list
ex1.1 <- products[1]
ex1.2 <- products["A"]
str(ex1.1)
## List of 1
## $ A:List of 3
## ..$ product : chr "A"
## ..$ sales : num 12000
## ..$ quarterly:List of 2
## ....$ quarter: int [1:4] 1 2 3 4
## ....$ sales : num [1:4] 6000 3000 2000 1000
```

# Code for (2) - get the contents of the first list element

```
# Example (2) - get the contents of the first list element
ex2.1 <- products[[1]]
ex2.2 <- products[["A"]]
ex2.3 <- products$A
str(ex2.1)
## List of 3
##
   $ product : chr "A"
## $ sales : num 12000
## $ quarterly:List of 2
## ..$ quarter: int [1:4] 1 2 3 4
## ..$ sales : num [1:4] 6000 3000 2000 1000
```

# Code for (3) - get the product name from the 1st list element

```
# Example (3) - get the product name for the first product
ex3.1 <- products[[1]][[1]]
ex3.2 <- products[["A"]][["product"]]
ex3.3 <- products$A$product
str(ex3.1)
## chr "A"</pre>
```

## Code for (4) - get annual sales of Product A

```
# Example (4) - get the annual sales for the first product
ex4.1 <- products[[1]][[2]]
ex4.2 <- products[["A"]][["sales"]]
ex4.3 <- products$A$sales
str(ex4.1)</pre>
```

num 12000

## Code for (5) - get the list for quarterly sales data

```
# Example (5) - get as a list, the detailed quarterly sales
ex5.1 <- products[[1]][[3]]
ex5.2 <- products[["A"]][["quarterly"]]
ex5.3 <- products$A$quarterly
str(ex5.1)</pre>
```

```
## List of 2
```

## \$ quarter: int [1:4] 1 2 3 4

## \$ sales : num [1:4] 6000 3000 2000 1000

## Code for (6) - get the quarters vector

```
# Example (6) - get the quarters
ex6.1 <- products[[1]][[3]][[1]]
ex6.1 <- products[["A"]][["quarterly"]][["quarter"]]
ex6.1 <- products$A$quarterly$quarter
str(ex6.1)
## int [1:4] 1 2 3 4</pre>
```

# Code for (7) - get the quarterly sales vector

num [1:4] 6000 3000 2000 1000

```
# Example (7) - get the quarterly sales
ex7.1 <- products[[1]][[3]][[2]]
ex7.1 <- products[["A"]][["quarterly"]][["sales"]]
ex7.1 <- products$A$quarterly$sales
str(ex7.1)</pre>
```

# Code for (8) - subset the quarterly sales for product A

```
# Example (8) - get the quarterly sales for the first two quarex8.1 <- products[[1]][[3]][[2]][1:2]
ex8.2 <-products[["A"]][["quarterly"]][["sales"]][1:2]
ex8.3 <-products$A$quarterly$sales[1:2]
str(ex8.1)</pre>
```

num [1:2] 6000 3000

## **Updating and adding new elements**

```
# Increase the sales of product A by 10,000
products$A$sales <- products$A$sales + 10000
# Add a new field to product A
products$A$type <- "Food"
str(products$A)

## List of 4
## $ product : chr "A"
## $ sales : num 22000</pre>
```

\$ quarterly:List of 2

\$ type : chr "Food"

## ..\$ quarter: int [1:4] 1 2 3 4

## ..\$ sales : num [1:4] 6000 3000 2000 1000

##

# Some observations on list subsetting

- Clearly, for list manipulation, the tag operator is the most programmer-friendly, so it is recommended to use this, and also try and ensure that the list elements are named
- Indexing using [[ by positive integer is very useful for looping structures, we will see an example of this shortly
- Functions such as those in the package purrr provide efficient and flexible ways to iterate through lists.

## **Iteration using Loops**

- Iteration is fundamental to all programming languages, and R is no exception.
- There are a number of basic looping structures than can be used in R, and we will focus on one of these, the for loop. The general structure is 'for(var in seq)expr, where:
  - var is a name for a variable that will change its value for each loop iteration
  - seq is an expression that evaluates to a vector
  - expr which is an expression, which can be either a simple expression, or a compound expression of the form {expr1; expr2}, which is effectively a number of lines of code with two curly braces.
- A convenient method to iterate over a vector (a list or an atomic vector), is to use the function seq\_along() which returns the indices of a vector.

# **Example loop structure (Atomic Vector)**

```
set.seed(100)
(v \leftarrow sample(1:6,10,replace = T))
## [1] 2 6 3 1 2 6 4 6 6 4
seq_along(v)
## [1] 1 2 3 4 5 6 7 8 9 10
n_six <- 0
for(i in seq_along(v)){
  n six \leftarrow n six + as.integer(v[i] == 6)
n six
```

## [1] 4

## **Example loop structure using [[ (List)**

Our goal is to find the average sales for the two products, and for this we can use a list.

```
sum sales <- 0
for(i in seq_along(products)){
  cat("Sales of product ",i," = ",products[[i]]$sales,"\n")
  sum sales <- sum sales+products[[i]]$sales</pre>
}
## Sales of product 1 = 22000
## Sales of product 2 = 8000
(avr_sales <- sum_sales / length(products))</pre>
```

## [1] 15000

#### The if statement

- if(cond) expr which evaluates expr if the condition cond is true
- if(cond) true.expr else false.expr, which evaluates true.expr if the condition is true, and otherwise evaluates false.expr

```
# create a test vector
v < -1:10
lv <- vector(mode="logical",length(v))</pre>
for(i in seq_along(v)){
  if(v[i] > mean(v))
    lv[i] <- TRUE
  else
    lv[i] <- FALSE
v[lv]
```

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#### Mini-Case Star Wars Movies

- The CRAN package repurrrsive is used
- It has lists on films (sw\_films), people (sw\_people), planets (sw\_planets), species (sw\_species), starships (sw\_starships) and species (sw\_species).
- Here we focus on the sw\_films list, which contains seven elements, and each element is a list that contains fourteen elements.

```
library(repurrrsive)
length(sw_films)
```

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

### The first movie

##

\$:List of 14

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```
str(sw_films[1])
## List of 1
```

```
## ..$ title : chr "A New Hope"
## ..$ episode_id : int 4
## ..$ opening crawl: chr "It is a period of civil war.\r\nl
```

```
## ..$ director : chr "George Lucas"
## ..$ producer : chr "Gary Kurtz, Rick McCallum"
```

```
## ..$ release_date : chr "1977-05-25"
## ..$ characters : chr [1:18] "http://swapi.co/api/people
## ..$ planets : chr [1:3] "http://swapi.co/api/planets
```

```
## ..$ starships : chr [1:8] "http://swapi.co/api/starshi
## ..$ vehicles : chr [1:4] "http://swapi.co/api/vehicle
## $ species : chr [1:5] "http://swapi.co/api/species
```

2. Base R - Subsetting

## Movie Directors for 1st and last movie

```
# Get the first film name and movie director
sw films[[1]][[1]]
## [1] "A New Hope"
sw_films[[1]][[4]]
## [1] "George Lucas"
# Get the last film name and movie director
sw films[[length(sw films)]][[1]]
## [1] "The Force Awakens"
sw_films[[length(sw_films)]][[4]]
```

## [1] "J. J. Abrams"

## Movie Directors for 1st and last movie

```
# Get the first film name and movie director
sw films[[1]]$title
## [1] "A New Hope"
sw_films[[1]]$director
## [1] "George Lucas"
# Get the last film name and movie director
sw films[[length(sw films)]]$title
## [1] "The Force Awakens"
sw_films[[length(sw_films)]]$director
```

## [1] "J. J. Abrams"

# Find all movies directed by George Lucas

- A for-loop structure (along with seq\_along()) is used to iterate over the entire loop and mark those elements as either a match (TRUE) or not a match (FALSE). This information is stored in an atomic vector.
- Before entering the loop, we create a logical vector variable (is\_target) of size seven (the same size as the list), and this will store information on whether a list item should be marked for further processing.
- For each list element we extract the directors name and check if it
  matches the target ("George Lucas"), and store this value in the
  corresponding element of is\_target.
- The vector is\_target can then be used to filter the original sw\_films list and retain all the movies directed by George Lucas.

#### **Code Solution**

```
# Search for movies by George Lucas and store these in a new
target <- "George Lucas"
# Create a logical vector that will hold information for posi
is target <- vector(mode="logical",length = length(sw films))
# Iterate through the entire sw films list (of 7)
for(i in seq_along(sw_films)){
  is_target[i] <- sw_films[[i]]$director == target</pre>
is_target
## [1] TRUE TRUE TRUE TRUE FALSE FALSE
target_list <- sw_films[is_target]</pre>
length(target list)
```

## [1] 4

## An additional task

In this case, we look to extract the movie titles into a new data structure, in this case an atomic vector.

```
## [1] "A New Hope" "Attack of the Clones" "The Phan
```

## [4] "Revenge of the Sith"

# Creating a new list

- One feature of R you will discover is that there are often may ways to achieve the same outcome.
- For example, another way to access the movies of George Lucas would be to re-arrange the *list of lists* into a single list, where each list element is an atomic vector of values (each of size seven).
- The process for creating the new data structure is:
  - Create a new list (sw\_films1) of elements you wish to store (for example, movie title, episode\_id and director) from the original list. This new list initially contains empty vectors.
  - Loop through the sw\_films list and append each movie title and director to the corresponding element of sw\_films1

## Sample solution

```
sw films1 <- list(title=c(), episode id=c(), director=c())</pre>
for(i in seq along(sw films)){
  sw_films1$title <- c(sw_films1$title,</pre>
                              sw films[[i]]$title)
  sw_films1$episode_id <- c(sw_films1$episode_id,</pre>
                              sw_films[[i]]$episode_id)
  sw_films1$director <- c(sw_films1$director,</pre>
                              sw films[[i]]$director)
str(sw films1)
```

```
## $ title : chr [1:7] "A New Hope" "Attack of the Clones
## $ episode id: int [1:7] 4 2 1 3 6 5 7
```

## \$ episode\_id: int [1:7] 4 2 1 3 6 5 7

## \$ director : chr [1:7] "George Lucas" "George Lucas" "George Lucas"

## List of 3

### **Parallel Vectors**

- Notice that we now have one list, and this list has three elements, each an atomic vector of size seven.
- These can be viewed as *parallel vectors*, where each vector is the same size, and the i-th element of each vector are related.
- This feature can be exploited to filter related atomic vectors using logical vector subsetting.

```
cat(sw_films1$title[1],"-",sw_films1$episode_id[1],"-",
    sw_films1$director[1],"\n")
```

```
## A New Hope - 4 - George Lucas
sw_films1$title[sw_films1$director=="George Lucas"]
```

```
## [1] "A New Hope" "Attack of the Clones" "The Phar
## [4] "Revenge of the Sith"
```

## **Useful R Functions**

R Function	Description
as.list()	Coerces the input argument into a list.
<pre>paste0()</pre>	Converts arguments to character strings and then concatenates (with no spaces)
rpois()	Generates up to n random numbers from a Poisson distribution with mean lambda
seq_along()	Generates a regular sequence that can be used to iterate over vectors
which()	Give the TRUE indices of a logical object

### Exercise 1

- Filter the list sw\_people (87 elements), contained in repurrrsive to include only those whose height is *not unknown*, and name this list sw\_people1.
- Use an atomic vector has\_height to filter the list, and populate this vector using a loop structure.
- This new list (sw\_people1) should have 81 elements.

```
sum(has_height)
## [1] 81
length(sw_people1)
```

## [1] 81

### Exercise 2

- Using a for loop over the filtered list sw\_people1 from exercise 2, create a list of people whose height is greater than or equal to 225 inches.
- The resulting vector should grow as matches are found, as we do not know in advance how many people will be contained in the result.
- Use the command characters <- c() to create the initial empty result vector.
- The if expression may be useful here too, and make sure that the height value is converted to a numeric value before evaluating.

The following result should be obtained.

```
# These are the characters whose height is >= 225
characters
```

```
## [1] "Chewbacca" "Yarael Poof" "Lama Su" "Tarfful"
```

### Exercise 3

- Based on the list sw\_species, and given that each species has a classification, create the following tabular summary, again using a loop to iterate through the list.
- Make use of the table() function

```
# A tabular summary of the types of species
t_species
```

```
## c_species
## amphibian artificial gastropod insectoid mammal
## 6 1 1 1 1 16
## reptilian sentient unknown
## 1 1 6
```

# **Lecture Summary**

- The four main ways to subset a vector, namely, by positive integer, the minus sign, logical vectors, and vector element names.
- The role of the [[ operator for processing lists, and how to distinguish this from the [ operator.
- The use of the \$ operator
- How to use the for loop structure to iterate through a list
- How to use the if statement when processing lists, and how that statement differs from the ifelse() function
- Additional R functions that allow you to process vectors
- How to solve all three test exercises.