Lists

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Lists

- A vector, all elements must be of the same mode.
- A list structure can combine objects of different types.
- For those familiar with Python, an R list is similar to a Python dictionary or, for that matter, a Perl hash.
- C programmers may find it similar to a C struct.
- The list plays a central role in R, forming the basis for data frames, object-oriented programming, and so on.

Creating Lists

- Technically, a list is a vector.
- Ordinary vectors are termed atomic vectors, since their components cannot be broken down into smaller components.
- In contrast, lists are referred to as recursive vectors.
- Let's consider an employee database.
- For each employee, we wish to store the name, salary, and a Boolean indicating union membership.
- Since we have three different modes here: character, numeric, and logical.

We could create a list to represent our employee,

Joe.

```
Console C:/R Home/ 🖒
> j <- list(name="Joe", salary=55000, union=T)</pre>
$name
[1] "Joe"
$salary
[1] 55000
$union
[1] TRUE
```

```
Console C:/R Home/ 🖒
> jalt <- list("Joe", 55000, T)</pre>
> jalt
[[1]]
[1] "Joe"
[[2]]
[1] 55000
[[3]]
    TRUE
```

 Actually, the component names: called tags in the R literature: such as salary are optional.

 However, it is generally considered clearer and less error-prone to use names instead of numeric indices.

 Names of list components can be abbreviated to whatever extent is possible without causing

ambiguity.

Since lists are vectors, they can be created via

vector().

List Indexing

```
Console C:/R Home/ 😞
$name
[1] "Joe"
$salary
[1] 55000
$union
[1] TRUE
> j$salary
[1] 55000
> j[["salary"]]
[1] 55000
> j[[2]]
[1] 55000
```

 You can access a list component in several different ways.

- We can refer to list components by their numerical indices, treating the list as a vector.
- However, note that in this case, we use double brackets([[]]) instead of single ones.
- So, there are three ways to access an individual component c of a list 1st and return it in the data type of c:
 - 1st\$c
 - lst[["c"]]
 - lst[[i]], where i is the index of c within lst.

- An alternative techniques listed is to use single brackets rather than double brackets.
 - lst["c"]
 - lst[i], where i is the index of c within lst.
- Both single-bracket and double-bracket indexing access list elements in vector-index fashion.
- But there is an important difference from ordinary (atomic) vector indexing.
- If single brackets [] are used, the result is another list: a *sublist* of the original.

```
Console C:/R Home/ 🖒
> j[1:2]
$name
[1] "Joe"
$salary
[1] 55000
> j2 <- j[2]
> j2
$salary
[1] 55000
> class(j2)
[1] "list"
> str(j2)
List of 1
 $ salary: num 55000
```

• The subsetting operation returned another list consisting of the first two components of the original list j.

```
Console C:/R Home/ 🖒
> j
$name
[1] "Joe"
$salary
[1] 55000
Sunion
[1] TRUE
> j[[1:2]]
Error in j[[1:2]] : subscript out of bounds
> j2a <- j[[2]]
> j2a
[1] 55000
> class(j2a)
[1] "numeric"
```

 By contrast, can use double brackets [[]] for referencing only a single component, with the result having the type of that component.

Adding and Deleting List Elements

```
Console C:/R Home/ 🖒
    <- list(a="abc", b=12)
$a
[1] "abc"
$b
[1] 12
> z$c <- "sailing" # Add a c component
> z
$a
[1] "abc"
$b
[1] 12
$c
    "sailing"
```

- The operations of adding and deleting list elements arise in a surprising number of contexts.
- New components can be added after a list is created.

Adding and Deleting List Elements (Cont.)

```
Console C:/R Home/ 🗇
> z[[4]] < -28
> z[5:7] <- c(FALSE, TRUE, TRUE)
$a
[1] "abc"
$b
[1] 12
$c
[1] "sailing"
[[4]]
[1] 28
[[5]]
[1] FALSE
[[6]]
[1] TRUE
[[7]]
[1] TRUE
```

 Adding components can also be done via a vector index.

Adding and Deleting List Elements (Cont.)

```
Console C:/R Home/ 🔊
> z$b <- NULL
$a
[1] "abc"
$c
[1] "sailing"
[[3]]
[1] 28
[[4]]
[1] FALSE
[[5]]
[1] TRUE
[[6]]
[1] TRUE
```

 You can delete a list component by setting it to NULL.

Adding and Deleting List Elements (Cont.)

• Since a list is a vector, you can obtain the number of components in a list via length().

```
Console C:/R Home/ 
> length(j)
[1] 3
>
```

Accessing List Components and Values

```
Console C:/R Home/ 🖒
$name
[1] "Joe"
$salary
[1] 55000
$union
[1] TRUE
> names(j)
[1] "name" "salary" "union"
```

• If the components in a list do have *tags*, as is the case with name, salary, and union for j, can obtain them via names ().

```
Console C:/R Home/ 
> ulj <- unlist(j)
> ulj
   name salary union
   "Joe" "55000" "TRUE"
>
> class(ulj)
[1] "character"
```

- To obtain the values, use unlist().
- The return value of unlist() is a vector, a vector of character strings.
- Note that the element names in this vector come from the components in the original list.

 On the other hand, if we were to start with numbers, we would get numbers.

So the output of unlist() in this case was a numeric vector.

• R chose the least common denominator: character strings.

Console C:/R Home/

```
Console C:/R Home/ 🖒
> w <- list(a=5, b="xyz")
> wu <- unlist(w)
> class(wu)
[1] "character"
> wu
    а
  "5" "xyz"
```

We can remove them by setting their names to

NULL.

```
Console C:/R Home/
> wu
  "5" "xyz"
 names(wu) <- NULL
> wu
[1] "5"
```

 We can also remove the elements' names directly with unname (), as follows.

```
> w <- list(a=5, b="xyz")
> wu <- unlist(w)
> wu
  "5" "xyz"
> wun <- unname(wu)</pre>
> wun
[1] "5" "xyz"
```

Applying Functions to Lists

 Two functions are handy for applying functions to lists: lapply and sapply.

Using the lapply() and sapply() Functions

- The function lapply() (for *list apply*) works like the matrix apply() function.
- Calls the specified function on each component of a list (or vector coerced to a list) and returning another list.

```
Console C:/R Home/ 
> lapply(list(1:3, 25:29), median)
[[1]]
[1] 2

[[2]]
[1] 27
```

• R applied median () to 1:3 and to 25:29, returning a list consisting of 2 and 27.

Using the lapply() and sapply() Functions (Cont.)

- In some cases, such as the example here, the list returned by lapply() could be simplified to a vector or matrix.
- This is exactly what sapply() (for simplified [l]apply) does.

Recursive Lists

```
Console C:/R Home/ 😞
> b <- list(u=5, v=12)
> c <- list(w=13)
> a <- list(b,c)
> a
[[1]]
[[1]]$u
[1] 5
[[1]]$v
[1] 12
[[2]]
[[2]]$w
[1] 13
> length(a)
```

- Lists can be recursive, meaning that you can have lists within lists.
- This code makes a into a twocomponent list, with each component itself also being a list.

Recursive Lists (Cont.)

```
Console C:/R Home/ 🖒
> c(list(a=1,b=2,c=list(d=5,e=9)))
$a
[1] 1
$b
[1] 2
$c
$c$d
[1] 5
$c$e
> c(list(a=1,b=2,c=list(d=5,e=9)),recursive=T)
      b c.d c.e
```

• The concatenate function c() has an optional argument recursive, which controls whether *flattening* occurs when recursive lists are combined.

Recursive Lists (Cont.)

- In the first case, we accepted the default value of recursive.
- That is FALSE.
- Which is obtained a recursive list, with the c component of the main list itself being another list.
- In the second call, with recursive set to TRUE.
- That is a single list as a result.
- Which is only the names look recursive.
- It's odd that setting recursive to TRUE gives a *nonrecursive* list.