

R-Programming Structures:

R is a block structured language in the manner of the ALGOL descendant family such as C, C++, Python, Perl and so on.

Here the basic structure of R as a programming language in which we review the more details on loops and functions and other control statements.

In many scripting languages we do not declare variables in R. R has a richer scoping structure.

Control statements:

Control statements in R very similar to those of the ALGOL descent family.

Here we will look at loop and if-else statements.

Loops:-

Here we define the for loop function in that function the following line should have been instantly recognized by python programmers.

Syntax: for (Declared variable in Initialized Variable)

for any next it means that there will be one iteration of the loop for each component of vector n with n taking on the values of those components.

In the 1st iteration $n = n[1]$ in the 2nd iteration $n = n[2]$ and so on.

For example we use this structure to output the square of every element in the vector

Eg:- for > $n \leftarrow c(5, 6, 7)$
 > for (n in n)
 > print(n^2)

O/p: 25 36 49

Eg:- > $i \leftarrow 1$
 > while ($i \leq 10$)
 > $i = i + 1$
 > print(i)

O/p:- [1] 13

Eg:- > a ← 1
 > while (a < 10)
 > print(a)
 > a = a + 1
 }

O/p: [1] 1
 [1] 2
 [1] 3
 [1] 4
 [1] 5
 [1] 6
 [1] 7
 [1] 8
 [1] 9

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2 Eg:- > i ← 1
 > while (i <= 10)
 > print(i)
 > i ← i + 1

O/p: 1 5 9

Eg:- > i ← 1
 > while (TRUE)
 > i ← i + 4

O/p: [1] 13

> if (i > 10) break

[while(TRUE) = repeat]

> i

Eg:- > i ← 1
 > repeat
 > i ← i + 4
 > if (i > 10)
 break

O/p: [1] 13

> i

(ans - i) 13

In loopings with while and repeat is also available with this while and repeat functions are complete with break. A statement that causes control to leave loop.
 → In the 1st example the variable i took on the values 1, 5, 9 and 13 as the loop went through its iterations. In that last case the condition $i \leq 10$ fail so, the break took hold and we left the loop.

Note:- repeat has no boolean exist condition. We must use break or something like return.

Next function

The next statement in R programming language is useful when we want to skip the current iteration of a loop without termination it.

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Ques: > $i \leftarrow 1$

> while ($i \leq 10$)

> if ($i == 5$)

> next

> point(1)

> $i = i + 1$

O/P: 1 2 3 4 6 7 8 9 10

Ques: > $x \leftarrow 1; 10$

> for (val in x)

{

 if (val == 3)

 {

 next

 }

 point(val)

}

O/P: 1 2 4 5 6 7 8 9 10

```

3eg!- > v ← LETTERS [1:6]
      for (i in v)
      {
        if (i == "D")
          o/p: A B C. E F
    
```

function $f(x)$ is called convex if for all $x_1, x_2 \in \text{dom } f$ and $\lambda \in [0, 1]$, we have
$$f(\lambda x_1 + (1-\lambda)x_2) \leq \lambda f(x_1) + (1-\lambda)f(x_2)$$

Scan() function: 19/05/2022
Here we use scan() read the no.of items and displays the items. here we can use in any looping function

```

File-1:1
1 > for fn in c ("file 1", "file 2")
2   print (scan (fn))
3
4   Read 6 items
5   [1] 1 2 3 4 5 6
6

File-2:1
1
2   Read 3 items
3   [1] 5 12 13

```

looping over non vector sets:

R does not directly support iteration over non vector sets. But there are a couple of indirect ways to accomplish it.

lapply:

The iterations of the loop are independent of each other, thus allowing to be performed in any order.

```
> file1 <- c(1, 2, 3, 4, 5, 6)  
> file2 <- c(5, 12, 13)  
> lapply(c("file1", "file2"), sum)
```

O/p: 6 14 16 + 5 6

get() function:

As its name implies in R programming language, it is used to return an object with the name specified as argument to the function. This is another method of print the values of the object i.e. Just like a print function this function can also be used to copy one object to another.

Syntax: get("Object")

Object are nothing but vector, array, list etc.

```
eg1: x1 <- c("a", "b", "c")
```

```
x2 <- c(1, 2, 3)
```

```
get("x2") O/P: 1 2 3
```

```
x4 <- get("x1")
```

```
print(x4) O/P: "a" "b" "c"
```

```
list1 <- list(x1, x2)
```

```
(where x1 & x2 are get("list1")) O/P: $x1 $x2
```

```
a b c
```

1 2 3

if-else

Syntax: v <- if(condition)

```
else exp1  
exp2
```

Eg1 > n <- 2

> m <- if (n==2) 1 else 0
 after press > n o/p: 2 → it's digit is 2 in result due to 2
 > else
 > n+1
 > y o/p: TRUE → it's digit is 1 in result due to 1

The result of y depending on weather condition is TRUE or FALSE we can use this fact to compile our code with out taking n value

> y <- if (n==2)
 > y <- n
 > else
 > y <- n+1

here n is not mentioned so output gets error!

Arithmetic and Boolean operators and values: 20/5/2022

Basic R Operators

Operation	Description	x%/%y	Integer division
x+y	Addition	x==y	Test for equality
x-y	Subtraction	x<=y	Test for less than or =
x*y	multiplication	x>=y	Test for greater than or equal to
x/y	division	x&&y	Boolean AND for scalars
x^y	Exponentiation	x y	Boolean OR for scalars
x%/%y	modular arithmetic	x&y	Boolean AND for scalars
			Vectors (vector x,y,result)
			Boolean OR for vectors
			(Vector x,y result)
		!x	Boolean negation

Example:

> n
 (i) TRUE FALSE TRUE
 > y
 (i) TRUE TRUE FALSE

> $x+y$

[1] TRUE FALSE FALSE

> $x \& y$ & & y[1]

[1] TRUE

> $x \& y$ # looks at just the first elements
of each vector.

[1] TRUE

> If($x \& y$ & & y[1]) print ("both TRUE")

[1] "both TRUE"

> if ($x \& y$) print ("both TRUE")

[1] "both TRUE"

warning message:

In If ($x \& y$) print ("both TRUE")

the condition has length > 1 and only the
first element will be used.

In R programming has no scalar types with scalars being
treated as one element vectors

In the above table there are different boolean operators
for the scalar and vector cases.

In the above example the central point is that in evaluating
an If, we need a single boolean not a vector of booleans.

Hence the warning seen in the proceeding example as
well as the need for having both the & and && operators

⇒ The boolean values TRUE and FALSE can be abbreviated
as T and F these values changes to one and zero

in arithmetic expression

Eg: $(1 < 2) & (3 < 4)$ frequent with operators & &

TRUE

> $(1 < 2) * (3 < 4)$

[1] 1

> $(1 < 2) * (3 < 4) * (5 < 1)$

[1] 0

> (1<2) == TRUE

> (1<2) == 1

In the 2nd compilation (statement) for instance the comparison (1<2) returns TRUE and (3<4), yields TRUE as well both values are treated as 1 so the product is 1

R functions look similar to those of C, JAVA, python and so on. Now, ever they have much more of a function programming flavours which has directly implications for the R programmers.

Default Values for Arguments

In R programming function can have default values of its parameters it means while defining a function, its parameters can be set values, it makes function more generic.

Eg1: function-name <- function (a, b = 10)

{

Point (a+b) base of this named arguments

function-name(10) o/p: 20
function-name(30, 40) o/p: 70

The console points on 70 because it overrides the 20. So,

Eg2: >m <- function(a, b, c = "Hai", d = "TKR")

Point (10, 20) base of this named arguments

m(a, b, c, d) base of this named arguments

o/p: 10 20 "Hai" "TKR"

The terminology named arguments that because R uses lazy evaluation. It does not evaluate an expression until or unless it needs. To the named argument may not actually be used

(P>E) & (S>I) <

P [D]

(I>D) & (P>E) & (S>I) <

I [D]

Return Values:

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The return values of a function can be any R object. The return value is often a list, it could even be another function we can transmit a value back to the caller by explicitly calling `return`. The value of the last executed statement will be returned by default.

Eg1:- Return value

odd count

> function (n)

{

> k <- 0

> for (n in n)

{

> if (n %% 2 == 1)

{

> k <- k + 1

}

> return(k) (0) > k

}

This function returns the count of odd numbers in the argument.

- we could simplify the code by eliminating the call to `return()` to do this we evaluate the expression to be returned `k` in the last statement.

Eg2:- odd count

> function (n)

{

> k <- 0

> for (n in n) {

> if (n %% 2 == 1)

> k <- k + 1

}

> k

}

Eg3:- odd count

> function (n)

{

> k <- 0

> for (n in n) {

> if (n %% 2 == 1)

> k <- k + 1

}

> for (n in n) {

> if (n %% 2 == 1)

> k <- k + 1

}

> k

}

In the eg3 the last executed statement here is the call to `for()` which returns the value `NULL` that means there would be no return value at all.

Returning Complex Objects

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The return object can value can be any R object (84)

We can return complex objects along with function name

Eg: g is a function which return a matrix and also

function f also left for writing self-explanatory problems

f ← function (x)

return (x^2)

return (t)

}

g()

function (x)

return (x^2)

order output - 103

using who

using diff. & same

o>t

Concepts - 103

In the above example if function have multiple return values place them in a list or other container.

functions are objects:

R functions are first class objects, that means they can be used for the most part just like other objects

In R a function is an object so the R interpreter is able to pass control to the function along with arguments that may be necessary for the function to accomplish the action.

Syntax of function creation:

f ← function (x)

{

return (x+1)

}

using who - 103

Function is a built-in R function whose job is to create functions

- There are two arguments to function the first is the formal argument list for the function we are creating i.e in the
- Second is the body of the function i.e the single statement return (x+1) the and argument must be of class expression, so the point is that the right hand side creates a function object i.e is "f"

Eg: f ← function (a,b)

{

return (a+b)

> f2 ← function (a,b)

{

return (a-b)

> f ← f1

{

return (a+b)

> f(3,2)

{

$f_1 \leftarrow f_2$

$f_2(3, 2)$

[ij 1] (Nested function, (or) function within a function)

$g \leftarrow \text{function}(h, a, b)$

$h(a, b)$

$g(f_1, 3, 2)$

[ij 5]

$g(f_2, 3, 2)$

[ij 1]

In the above example functions are objects we can also assign them use them as arguments to other functions e.g. h this process we can call it as a function call, nested within another function or function call with in a function.

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Environment Scope Issues:

Environment can be through of a collection of objects "functions, variables". An environment is created when we first sign up the R Interpreter that means RStudio. Any variable we define is now in this environment. A function formally referred to as a closure in the R documentation consist not only of its arguments and body but also of its environment. In this environments works in R is essential for writing effective R functions.

e.g:- $a \leftarrow 2$

$b \leftarrow 5$

$f \leftarrow \text{function}(x)$

$x \leftarrow 0$

$w \leftarrow 12$

$g \leftarrow \text{function}(y)$

$y \leftarrow 0$

$d \leftarrow 8$

$h \leftarrow \text{function}(z)$

$z \leftarrow 0$

$\text{return}(h(w))$

}

Environment (f)

3 of background

variables

O/p: R_GlobalEnv

$a \leftarrow 2$

$b \leftarrow 5$

$f \leftarrow \text{function}$

$n \leftarrow 0$

$g()$

\rightarrow

it gives output for how many

variables are present

O/p: a b f

variables

are present

Here the function `f` is created to at the top level i.e. at the interpreter command prompt and thus has the top level environment which in R output is referred to as `R_GlobalEnv`. But which we refer to in R code as `GlobalEnv`.

In your R program if we run any program as a batch file i.e. is consider top level

`ls()`:

```
a<-2  
b<-5  
f<function(a)  
n<-0  
ls()
```

O/P: "a" "b" "f"

`Eq2: w<-12`

`f<function(y)`

```
d<-8  
h<function()  
return(h)
```

`is.function(f)`
`w: num[12]`
3 Environment(f)
`ls()`
O/P:- "w" "d"

In the above function `ls()` list the object of an environment. If we call it at the top level we get the top level environment.

The top level environment here includes the variable "w". which is actually used in "f" function

> environment()

<environment: R_GlobalEnv>

> GlobalEnv

<environment: R_GlobalEnv>

In the above two commands we can used to know the R environment

Scope Issues :-

If we were working with the C language we would say that, the variable "w" in the previous example is global to function. While "d" is local to f. things are similar in R, but R is more hierarchical compared to C

> outer_fun ← function()

 {

 b ← 20 } global variable.

 inner_fun ← function()

 {

 scope_1 scope_2 { c ← 30 } local variable

 3

 (local) a ← 10 } global variables (in) outer

 }

(87)

so, here a, b are the global variables and c is local variable.

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we have h() being local to f() just like d. In such a situation scope has to be hierarchical in R setup, d which is local to f() is in turn Global to h(). The same is true for y as argument are considered locals in R. the hierarchical nature of scope implies that since w is global to f. It is global to h as well we do use a within h()

Eg:- w ← 12
f ← function (y),

{

 d ← 8

 h ← function()

 {

 return (h())

 y

 return (h())

 3

On :- f(2)
h ← function

 {

 return (d + (w+y))

 9

The R interpreter finds that there was no local variables of that name, so it ascended to the next higher level in this case the top level where it founds a variables 'w' with value 12. Keep in mind that a variable 'w' with value 12 is local to f(). It is invisible at the top level. If h() is local to f(). It displays error: Object 'h' not found as it is not defined.

```

Eg1 - f ← function(y)
      {
        d ← 8
        h ← function()
        {
          return (d * (w+y))
        }
        return (h())
      }
    
```

O/p: 112

```

      > f ← function(y)
        {
          d ← 8
          return (h())
        }
      }

      > h ← function()
        {
          return (d * (w+y))
        }
      
```

O/p:- error: d,w,y values
are not there

In the above examples environments

Created by inheritance that means inner most environment is used first A reference to d which h() would refers to h()'s d not f()'s compare the both Example the example 2 does not work as d is no longer in the environment h because h() is defined at the top level thus an error is generated

Eg2 - f ← function(y)

```

      {
        d ← 8
        return (h(d,y))
      }
    
```

> h ← function (dee;yyy)

```

      {
        return (dee * (w+yyy))
      }
    
```

> f()

⇒ > f ← function (y, ftn)

```

      {
        d ← 8
      }
    
```

point (environment (ftn))

return ftn (d,y)

}

> h ← function (dee,yyy)

```

      {
        return (dee * (w+yyy))
      }
    
```

> w ← 12

> f (3,h)

when f() executed the formal argument ftn was matched by the actual argument h() ftn has h environment

More on Isc()

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Isc() function is form with in a function returns the name of the current local variables with the environment arguments it will point the names of the local of any frame in the calling function

Eg:- f<function (y)

{

 d<=8

 return p(n(d,y))

}

> h<function (dee, yyyy)

 Point(Isc())

Point (is(envir = parent.frame (n=1)))

 return '(dee*(w+yyyy))

{

>f(2)

O/P:- [1] "dee" "yyyy"

so what [1] "d" "y" is variable that got set to as global
or top level

In the above example parent.frame(), the argument 'n' specifies how many frames to go up in the call chain (or) call function. here we were in the executing of h() which had been called from f. So, Specifying n=1 gives us f()'s frame and thus we get its locals

Function have no side effects :-

function programming property is that functions do not change non-local variables that is generally their are no side effects the code in the function has read access to its non-local variables but it does not have write access to them. in the previous examples code than appear to re assign those values but the action with will effect only copies. not the variables themselves. lets see the one example by adding some more go to our previous example.

```

> w ← 13
f ← function (y)
{
    a ← 8
    w ← w + 1
    y ← y - 2
    Point(w)
}
h ← function ()
{
    return (x)
    Point(w)
}
t ← 4
f(t)
0/p t [1] 13
> w
12
> t
4

```

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Here w at the top level did not change even though it appears to change within $f()$. Only a local copy of w within $f()$ changed. Similarly the top level variable t didn't change even though its associated formal argument y did change.

Note: Reference to the local w actually go to the same memory location as the global one until the value of the local changes. In that case a new memory location is used.

Extended Example: A function to display the contents of a call frame

The values of the local variables in our current function we want to know through our board in a debugging settings using Single Step. This means to know the values of the locals in the parent function i.e. the one from which the current function was called here we will develop code to display these values thereby further demonstrating access to the environment hierarchy.

Eg: f <function()

{
 a <- 1

 return (g(a)+a) o/p:- 2

g(1)+1

g(a)

(11)

g <function (aa)

{
 b <- 2

 aa+b <- b(a+a+b) o/p:- 3

 return (aa+b)

g(aa)

h <function (aaa)

{
 c <- 3

 aaa+c <- return (aaa+aa) o/p:- 4

when we calls to f it interns calls g() which then calls h() we want to know the values of the local variables of the current function .the variables aa, aab, b and while we are in g() we also wish to know the values of the locals in f() at the time of the call to g().as well as the values of the global variables.

09/06/2022.

No pointers in R

R does not have variables corresponding to pointers or references like 'c' language this can make programming more difficult in some cases

for example you can't write a function directly changes its arguments like

Eg:- >n= [13, 5, 12] (array)

> n.sort()

>n

O/p:- 5 12 13

The current version of R programming has an experimental feature called reference classes which may reduce the difficulty hence in the above example the value of n, the argument to sort() changed

Eg:- >n <- c(13, 5, 12)

> sort(n)

>n

O/p:- 5 12 13

>n

13 5 12

Eg 8- > n <- c(13, 5, 12)

> m <- sort(n)

> m

5 12 13

(12)

In the example 2 the argument to sort does not change if we do want m to change in this R code the solution is to reassign the arguments.

Writing upstairs :-

In any variable higher in the environment hierarchy than the level at which your writing statement exists.

We can use the super "super assignment operator" or the assign function. because we cannot directly write access to variables at the higher level why are the standard ' \leftarrow ' operator is not possible.

Writing to non-locales with the Super Assignment Operator.

Eg 9 two \leftarrow function (x)

{

u \leftarrow 2 * u

z \leftarrow 2 * z

}

> n \leftarrow 1

> z \leftarrow 3

> u

Error: "Object" "u" not found

> two(1)

> m [1] 1

> z [3] 3

> u [2]

1. m was the actual argument two(). In the above example it retains the value 1 after the call. this is because its value 1 was copied to the formal argument u. which is treated as a local variable with in the function. Thus when u changed, m did not change with it.

1. The two z values are entirely unrelated to each other. One is top level and other is local to f . The change in the local variable has no effect on the global variable. Of course having two variables with the same name is probably not good programming.
2. The u value did not even exist as a top level variable prior to f . Hence the 'not found' error message. It was created as a top level variable by the super assignment operator to function $\rightarrow f$. As confirmed after the call.

Eg: $\text{f} \leftarrow \text{function}()$

u
[1]
 $b <$
[3]

Req: In \leftarrow function (x) it aligns works with all
functions. \leftarrow function (x) is different from \leftarrow function ($x + 1$)
as the \leftarrow $x + 1$ option of a function that reflects
all of y (super assignment) but x is defined
 y $\leftarrow 3$ (super assignment) but x is defined

$\text{f}(1)$

Environment last got

$\text{return}(m)$

Environment last got

y
 $> x$

(P/P) # (Error - no object with previous)

$\text{f}(m)$

(x or y) not

[1] 4

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$> x$ ((1, 4)) object names

Error: Object 'm' is not found

Here m is defined within function f , when \leftarrow (m) is executed when R interpreter sees a 'super assignment' to m it starts going up the hierarchy at the first top level the environment with PN function & it does not find an m and so that x is the one that $\text{return}()$ to not m at the top level writing to non-locals with assigns \leftarrow

Assign a value to a name in environment assigns to write to upper level variables in R current versions

Syntax:-

assign (variable, value)

Programmatic approach used to store data to

Example :-

```

> assign("a", 10)
> print(a)
[1] 10

```

Eg:-

```

> two <- function(x) {
    > assign("a", x, envir = globalEnv)
    > x^2
}
> two(2)
[1]
[2]
[3]
[4]
[5]
[6]
[7]
[8]
[9]
[10]
[11]
[12]
[13]
[14]
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[89]
[90]
[91]
[92]
[93]
[94]

```

In the above example we replaced the super assignment operator with a call to `assign()` function. that call instructs R to assign the value x^2 to a variable. A further up the call stack specifically in the top level environment

Paster() :-

Here we are using `paste()` to create new variables combining the prefix `a-` with the remaining index!

Eg:-

```

for(i in 1:3)

```

```

{
  assign(paste("a-", i))
}
```

```

}
```

O/p:-

```

a-1 [1] 1
a-2 [2] 4
a-3 [3] 9

```

Extended Example:-

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Discrete Event Simulation in R :-

DES is widely used in businesses, industry and government sectors. The term discrete refers to the fact that the state of the system changes only in discrete quantities rather than changing continuously.

eg:- ATM machine

(95)

this example the state of our system at a time t' to be the no. of people in queue at the time. the state changes only by $+1$ when someone arrives, or by -1 when a person finishes a an ATM transaction.

General to DES Operation is maintenance of the event list which is simply a list of scheduled events. this is general DES term so the word list here does not refer to the R Data type. Instead of that we will represent the event list by a data frame.

Eg:- cus 1 arrives at time 23.12

2 arrives at time 25.88

3 arrives at time 25.97

1 finishes services at time 26.02

So, In the above data frame with the 1st row containing the earliest schedule the 2nd containing the 2nd earliest and so on

In this example if the customer arrival occurs when the queue is empty. that customer services will begin. One event triggers setting upto another

The person will be finished which is another event that must be added to event list

One of the oldest approach to writing DES code is the event oriented approach. consider the ATM situation at time '0' (zero) the queue is empty the stimulation code randomly generates the code of the 1st arrival at the 2.3 at this point the event list is simply (2.3,"arrival") this event is pulled of the list. stimulated time is updated to 2.3

The queue for the ATM is empty so we start the service by randomly generating the service time say it is 1.2 time units. then the completion of service will occur at stimulated time $2.3 + 1.2 = 3.5$,

- Library functions using in DFS :-
- Scheduled event \rightarrow SchedEvent(): It algorithm 8/10
Insert a newly created event into the event list
 - getNextEvent(): pulls the earliest event of the event list
 - doStimulation \rightarrow doStim()
 - Include the core loop of the stimulations repeatedly calls get next event. To gets earliest of the pending events, updates the current stimulated time.
 - Sim\$currtime:
to reselect the occurrences the event and calls the application specific function readyEvent() to process this newly occurred event.
 - ReactEvent(): It performs the event when it will occurs
 - InitGbls(): It is used to initialize global variables.
 - PointResults() \rightarrow point results.
Points the application specific results of specific events

When should we use global variables?

14/06/2022

Variables that are created outside of function are known as global variables.

- Global variables can be used by every programmer, But inside and outside of a function.
- The term global variable will be used to include any variable located higher in the environment than the level of the given code of interest in R library super assignment operator is used in many functions. So, threaded code and GPU code which are used for writing fast programs to make heavy use of global variables, which provides the main avenue of communication b/w parallel actors (debugging, maintenance and error detection used)

1. `cat("R Programming")` → En-2 string with all in
my_fun ← function() { cat("R Programming language") }
my_fun()

{
 Paste (txt, "language")
}
my_function()

O/p: R programming language.

So, the above two examples are cleaner code is usually easier to write, debug and maintain for this reason avoiding unanticipated errors or uncleared errors and simplifying the code that we choose to use.

global

Closures :=

R closure consisting of a function's arguments and body together with its environment at the time of the call. The fact that the environment is included in exploited in a type of programming that uses a feature also known as closure.

A closure consisting of a function that steps up a local variable and then creates another function that access the variables

Variables

fgt > counter ← function() > c2 ← c1(c)

9

$\text{ctr} \leftarrow 0$ (x)

47 *Fauvelot*

$\text{ctr} \leftarrow \text{ctr} + 1$

8

$\text{ctr} \leftarrow \text{ctr} + 1$

`Cat("This count currently has value ", ctr, "in")`

Value", cr, "n")

3) $My > 1000 \text{ Nm} \cdot \text{rad}^{-1}$ $\rightarrow 250 \text{ rad} \rightarrow 182^\circ$

Digitized by srujanika@gmail.com

$c_1 \leftarrow \text{counter}()$ $(N2)2P \Rightarrow N2$

~~counter~~ $(\sqrt{2})^2 \rightarrow \infty$

systems (e.g., biological systems).

$\text{ctr} \leftarrow \text{ctr} + 1$

`Car("this count currently has value", cto, "in")`

In the above example, c_1 function and c_2 functions series as completely independent counters where we invoke each of them a few times. (98)

Recursion :-

A recursion function call itself the idea is this -

1. To solve a problem type x by writing a recursive function,
- Break the original problem, of type x into one or more smaller problems of type x .
- Within $f(x)$ call $f(x)$ on each of the smaller problems.
- Within $f(x)$ piece together the result of $f(x)$ point to solve the original problem.

Quick Sort Implementation :-

In quick sort algorithm used to sort a vector of number from smallest to largest

Bn^{o2} 5, 4, 12, 13, 3, 8, 88

- First compare every thing to the first element 5
- To form two sub vectors one consisting of the elements less than 5 and other consisting of the greater than or = to 5
 - i.e. Sub vector 1 is 4, 3 & sub vector 2 is 12, 13, 8, 88
- we then call the function on the sub vector 1 & return 3, 4 and sub vector 2 returning 8, 12, 13, 88 then we string goes together with the final e.g. 3, 4, 5, 8, 12, 13, 88

Source code in Al-

```
qs ← function(x)
    if (length(x) ≤ 1)
        return (x)
    pNot ← x[0]
    therest ← x[1]
    sv1 ← therest (therest < pNot)
    sv2 ← therest (therest ≥ pNot)
    sv1 ← qs(sv1)
    sv2 ← qs(sv2)
    return (c(sv1, pNot, sv2))
```

from the above source code if (length(x) ≤ 1) return (x) without the function would keep calling itself.

repeatedly an empty vector executing forever. So, recursion certainly is an elegant way to solve many problems (99). But recursion has two potential drawbacks.

It's fairly abstract & the recursion is really just the inverse of proof by mathematical induction; but many programmers find it tough.

Recursion is very lavish in its use of memory which may be an issue in R. It applies to large problems.

Binary Search Tree (Extended Example):

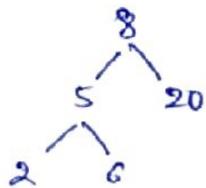
20/6/2022

The library function in R programming to create a new tree uses "TSt" In other programming languages like C, C++, Java, etc., fortune we use Data structures are common in both computer science and statistics.

But in R, part library for a recursive partitioning approach to regression and classification is very popular.

Trees can be implemented in R itself and it performs well. It is not an issue.

Eg:-



We know very well binary search tree principle.

If value is less than the value at leftly if any less than or equal to that of the root (parent) while the value at the rightly if any greater than of the parent (root).

A matrix in R is a time consuming activity. We allocate new space for a trees matrix several rows at a time instead of row by row. The no. of rows allocated each time will be given the variable INC (increment) as it is common with tree travels we implement our algorithm with recursion.

Eg:- $x \leftarrow \text{new}^{\text{tree}}(8, 3)$

with this $x[1] \leftarrow \text{new}^{\text{tree}}(5, 3)$ $x[2] \leftarrow \text{new}^{\text{tree}}(6, 3)$ $x[3] \leftarrow \text{new}^{\text{tree}}(NA, 3)$

\$mat

\$inc (increment)

[1] NA NA 8

[2] NA NA NA

[3] NA NA NA

The session continues that when our initial allocation of 3 rows is full `ps()` allocates three new rows for a total of 6 (101)

22/06/2022

In R programming the replacement function named as "names"

Eg:- 1 > `x <- c(1, 2, 3)`

> `Replace(x)` o/p:- NULL

> `names(x)` o/p:- NULL

Eg2:- > `x <- c(1, 2, 3)`

> `names(x) <- c("a", "b", "c")`

> `names(x)` o/p:-

1 "a" 2 "b" 3 "c"

Eg3:- > `x <- c(1, 2, 3)`

> `names(x) <- c("a", "b", "c")`

> `Replace(x, names(x),`

"x", "y", "z")

> `Replace(y)`

o/p:-

x y z

1 2 3

In the above examples we will use the names and replace functions in the R console

What's consider a Replacement functions :-

Any assignment statement in which the left side is not just a identifier is consider as a replacement function.

Eg:- > `g(u) <- v`

> `u <- "g <- (v, value=v)"`

} same, executes

Note :- the replacement function has one or more arguments than the original function `g()`, a named argument `value` for reasons replaced. same.

Eg2:- > `m <- q(8, 88, 5, 12, 13)`

> `x <- 8, 88, 5, 12, 13`

> `m[3]` o/p:- 5

> `"["(m, 3)` o/p:- 5

In the above example we will use the function "[" if for reading vector elements, the complicated call of a function "`c<-()`" is used to write.

Eg3:- > `p <- "(m, 2:3 value = 99:100)"`

> `m` o/p:- 8 10 100 13

Is simply performing what happen in the console when execute this `m[2:3] <- 99:100`. we can easily verify what occurring like

Eg 4:- > `x <- c(8, 88, 5, 12, 13)`

> `x[2:3] <- 99:100`

> `x` o/p:- 8 99 100 12 13

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