

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

Source on Save Run

```
1 # recursive functions
2
3 recur_fibo <- function(n) {
4   if(n <= 1) {
5     return(n)
6   } else {
7     return(recur_fibo(n-1) + recur_fibo(n-2))
8   }
9 }
10
11 # take input from the user
12 nterms = as.integer(readline(prompt="How many terms? "))
13
14 # check if the number of terms is valid
15 if(nterms <= 0) {
16   print("Please enter a positive integer")
17 } else {
18   print("Fibonacci sequence:")
19   for(i in 0:(nterms-1)) {
20     print(recur_fibo(i))
21   }
22 }
23
```

23:1 (Top Level) R Script

Console Terminal Background Jobs

```
R422 ~/%
> # check if the number of terms is valid
> if(nterms <= 0) {
+   print("Please enter a positive integer")
+ } else {
+   print("Fibonacci sequence:")
+   for(i in 0:(nterms-1)) {
+     print(recur_fibo(i))
+   }
+ }
Error in if (nterms <= 0) {: missing value where TRUE/FALSE needed
>
```

Environment History Connections Tutorial

R Global Environment 122 MiB

Data

Name	Description	Version
a	1 obs. of 1 variable	
A	num [1:2, 1:2] 2 0 1 3	
b	196 obs. of 16 variables	
B	num [1:2, 1:2] 5 2 4 -1	
data	1000 obs. of 9 variables	
data1	1000 obs. of 9 variables	
df	10 obs. of 4 variables	
input	32 obs. of 2 variables	
list1	List of 3	
m	int [1:2, 1:4] 1 2 3 4 5 6 7 8	
M	num [1:2, 1:3] 2 1 6 10 5 4	

Files Plots Packages Help Viewer Presentation

Install Update

Name	Description	Version
askpass	Safe Password Entry for R, Git, and SSH	1.1
assertthat	Easy Pre and Post Assertions	0.2.1
backports	Reimplementations of Functions Introduced Since R-3.0.0	1.4.1
base64enc	Tools for base64 encoding	0.1-3
bit	Classes and Methods for Fast Memory-Efficient Boolean Selections	4.0.5
bit64	A S3 Class for Vectors of 64bit Integers	4.0.5
blob	A Simple S3 Class for Representing Vectors of Binary Data ('BLOBs')	1.2.3
broom	Convert Statistical Objects into Tidy Tibbles	1.0.2
bslib	Custom 'Bootstrap' 'Sass' Themes for 'shiny' and 'markdown'	0.4.2
cachem	Cache R Objects with Automatic Pruning	1.0.6
callr	Call R from R	3.7.3
cellranger	Translate Spreadsheet Cell Ranges to	1.1.0

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

```
1 #1
2 num = as.integer(readline(prompt = "Enter a number: "))
3 if(num < 0) {
4   print("Enter a positive number")
5 } else {
6   sum = 0
7   # use while loop to iterate until zero
8   while(num > 0) {
9     sum = sum + num
10    num = num - 1
11  }
12  print(paste("The sum is", sum))
13 }
```

13.2 (Top Level) R Script

Console Terminal Background Jobs

```
R 4.2.2 ~/> num = as.integer(readline(prompt = "Enter a number: "))
Enter a number: if(num < 0) {
Warning message:
NAS introduced by coercion
>   print("Enter a positive number")
[1] "Enter a positive number"
> } else {
Error: unexpected '}' in "}"
>
```

Environment History Connections Tutorial

Global Environment

Variable	Value
T1	num [1:3, 1:3] -0.2963 -0.0741 0.40...

Values

a1	num [1:3] 3 2 5
a2	num [1:3] 2 3 2
a3	num [1:3] 5 2 4
apple	chr [1:3] "red" "green" "yellow"
apple_colors	chr [1:7] "green" "green" "yellow" "r..."
attempts	num [1:10] 1 3 2 3 2 3 1 1 2 1
geeks	num [1:4] 23 56 20 63
labels	chr [1:4] "Mumbai" "Pune" "Chennai" "..."
name	chr [1:10] "Anastasia" "Dima" "Kather..."
nterms	NA_integer_
num	NA_integer_

Files Plots Packages Help Viewer Presentation

Install Update

Name	Description	Version
<input type="checkbox"/> askpass	Safe Password Entry for R, Git, and SSH	1.1
<input type="checkbox"/> assertthat	Easy Pre and Post Assertions	0.2.1
<input type="checkbox"/> backports	Reimplementations of Functions Introduced Since R-3.0.0	1.4.1
<input type="checkbox"/> base64enc	Tools for base64 encoding	0.1-3
<input type="checkbox"/> bit	Classes and Methods for Fast Memory-Efficient Boolean Selections	4.0.5
<input type="checkbox"/> bit64	A S3 Class for Vectors of 64bit Integers	4.0.5
<input type="checkbox"/> blob	A Simple S3 Class for Representing Vectors of Binary Data ("BLOBs")	1.2.3
<input type="checkbox"/> broom	Convert Statistical Objects into Tidy Tibbles	1.0.2
<input type="checkbox"/> bslib	Custom 'Bootstrap' 'Sass' Themes for 'shiny' and 'rmarkdown'	0.4.2
<input type="checkbox"/> cachem	Cache R Objects with Automatic Pruning	1.0.6
<input type="checkbox"/> callr	Call R from R	3.7.3
<input type="checkbox"/> cellranger	Translate Spreadsheet Cell Ranges to	1.1.0

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

```
1 my_vec <- 1:10
2 my_vec
3
4 my_vec^2
5
```

Run Source

Console Terminal Background Jobs

```
R 4.2.2 ~ /
> %>%
Error: unexpected SPECIAL in "%>%"
> my_vec <- 1:10
> my_vec
[1] 1 2 3 4 5 6 7 8 9 10
> my_vec^2
[1] 1 4 9 16 25 36 49 64 81 100
>
```

Environment History Connections Tutorial

Global Environment

Values

a1	num [1:3]	3 2 5
a2	num [1:3]	2 3 2
a3	num [1:3]	5 2 4
apple	chr [1:3]	"red" "green" "yellow"
apple_colors	chr [1:7]	"green" "green" "yellow" "r..."
attempts	num [1:10]	1 3 2 3 2 3 1 1 2 1
geeks	num [1:4]	23 56 20 63
labels	chr [1:4]	"Mumbai" "Pune" "Chennai" "..."
my_vec	int [1:10]	1 2 3 4 5 6 7 8 9 10
name	chr [1:10]	"Anastasia" "Dima" "Kather..."
nterms	NA_integer_	
num	NA_integer_	

Files Plots Packages Help Viewer Presentation

Install Update

Name	Description	Version
<input type="checkbox"/> askpass	Safe Password Entry for R, Git, and SSH	1.1
<input type="checkbox"/> assertthat	Easy Pre and Post Assertions	0.2.1
<input type="checkbox"/> backports	Reimplementations of Functions Introduced Since R-3.0.0	1.4.1
<input type="checkbox"/> base64enc	Tools for base64 encoding	0.1-3
<input type="checkbox"/> bit	Classes and Methods for Fast Memory-Efficient Boolean Selections	4.0.5
<input type="checkbox"/> bit64	A S3 Class for Vectors of 64bit Integers	4.0.5
<input type="checkbox"/> blob	A Simple S3 Class for Representing Vectors of Binary Data ("BLOBs")	1.2.3
<input type="checkbox"/> broom	Convert Statistical Objects into Tidy Tibbles	1.0.2
<input type="checkbox"/> bslib	Custom 'Bootstrap' 'Sass' Themes for 'shiny' and 'rmarkdown'	0.4.2
<input type="checkbox"/> cachem	Cache R Objects with Automatic Pruning	1.0.6
<input type="checkbox"/> callr	Call R from R	3.7.3
<input type="checkbox"/> cellranger	Translate Spreadsheet Cell Ranges to	1.1.0

RStudio

File Edit Code View Plots Session Build Debug Profile Tools Help

Go to file/function Addins

```
1 # Get the input values.
2 input <- mtcars[,c('wt','disp')]
3
4 # Give the chart file a name.
5 png(file = "scatterplot.png")
6
7 # Plot the chart for cars with weight between 2.5 to 5 and mileage between 15 and 30.
8 plot(x = input$wt, y = input$disp,
9      xlab = "Weight",
10     ylab = "Displacement",
11     xlim = c(2.5, 5),
12     ylim = c(15, 30),
13     main = "Weight vs Displacement")
14 }
```

Environment History Connections Tutorial

R Global Environment

input	32 obs. of 2 variables
list1	List of 3
m	int [1:2, 1:4] 1 2 3 4 5 6 7 8
M	num [1:2, 1:3] 2 1 6 10 5 4
mtcars	32 obs. of 11 variables
n	int [1:2, 1:4] 8 9 10 11 12 13 14 15
new	1000 obs. of 9 variables
relation	List of 12
t	num [1:2, 1:2] 65 82 82 117
T1	num [1:3, 1:3] -0.2963 -0.0741 0.40...
v	32 obs. of 11 variables

Values

a1 num [1:3] 3 2 5

Files Plots Packages Help Viewer Presentation

Console Terminal Background Jobs

```
R 4.2.2 > input <- mtcars[,c('wt','disp')]
>
> # Give the chart file a name.
> png(file = "scatterplot.png")
>
> # Plot the chart for cars with weight between 2.5 to 5 and mileage between 15 and 30.
> plot(x = input$wt, y = input$disp,
+      xlab = "Weight",
+      ylab = "Displacement",
+      xlim = c(2.5, 5),
+      ylim = c(15, 30),
+      main = "Weight vs Displacement")
+ )
>
```

```
R Console
Error in -c("A", "BA", "Bt1", "Bt2", "Cr", "R") :
  invalid argument to unary operator
>
> # REGEX rules
> p <- c("A", "BA|AB", "Bt|Bw", "Bt3|Bt4|2B|C",
+       "Cr", "R")
Error: object 'p' not found
>
> # Compute genhz labels and
> # add to loafercreek dataset
> loafercreek$genhz <- generalize.hz(
+   loafercreek$hname,
+   n, p)
Error: object 'loafercreek' not found
>
> # Extract the horizon table
> h <- horizons(loafercreek)
Error: object 'h' not found
>
> # Examine the matching of pairing of
> # the genhz label to the hnames
> table(h$genhz, h$hname)
Error in table(h$genhz, h$hname) : object 'h' not found
>
> |
```

```
Untitled - R Editor
library(aqp)
library(soilDB)

# Load from the loafercreek dataset
data("loafercreek")

# Construct generalized horizon designations
n <- c("A", "BA", "Bt1", "Bt2", "Cr", "R")

# REGEX rules
p <- c("A", "BA|AB", "Bt|Bw", "Bt3|Bt4|2B|C",
      "Cr", "R")

# Compute genhz labels and
# add to loafercreek dataset
loafercreek$genhz <- generalize.hz(
  loafercreek$hname,
  n, p)

# Extract the horizon table
h <- horizons(loafercreek)

# Examine the matching of pairing of
# the genhz label to the hnames
table(h$genhz, h$hname)
```

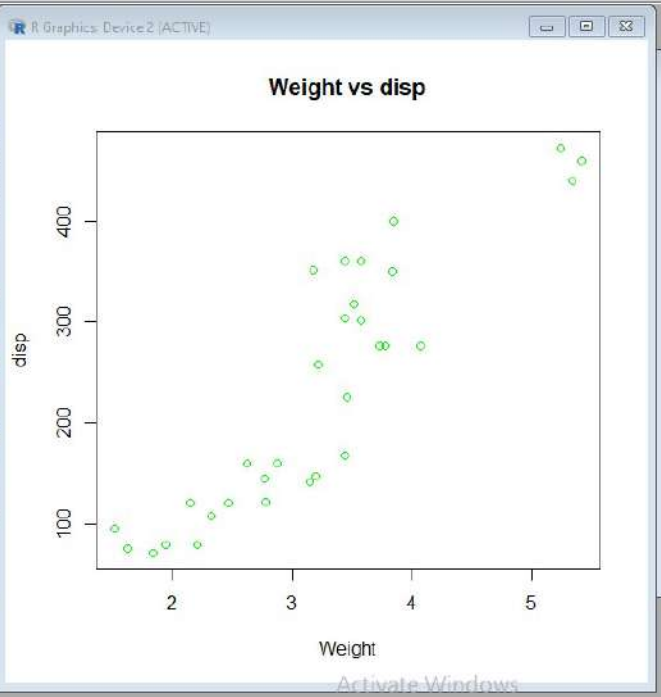
```
#4
a$cyl <- factor(a$cyl)
mylm <- lm(mpg ~ cyl, data = a)
summary(mylm)$coef

with(a, tapply(mpg, cyl, mean))

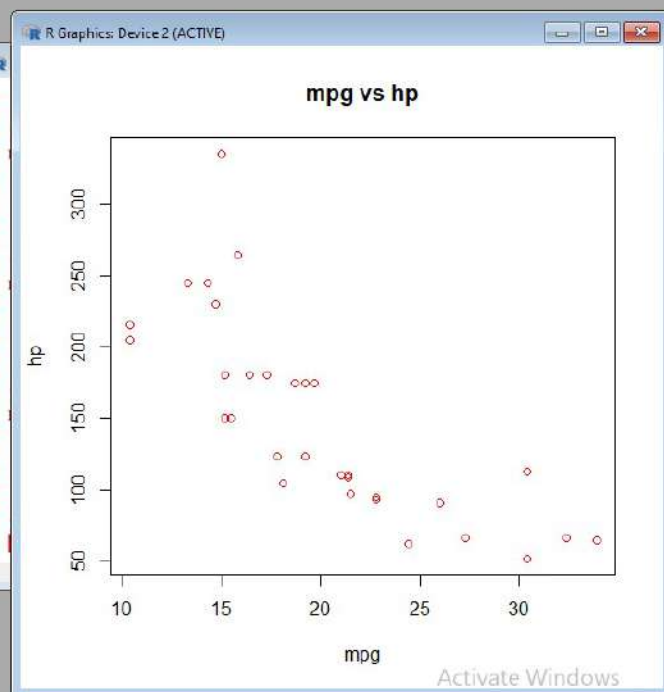
#5
min(a$wt)
max(a$wt)

##3rd
#1
plot(x = a$wt, y = a$displ,
     xlab = "Weight",
     ylab = "displ",
     main = "Weight vs displ",
     col = "green")
```

```
Warning message:
1: In plot.1:
2: In plot.1:
3: In axis(1:
4: In axis(1:
5: In box(1:
6: In title:
>
> plot(x = a$wt, y = a$displ,
+       xlab = "Weight",
+       ylab = "displ",
+       main = "Weight vs displ",
+       col = "green")
+ )
+ )
Error: unexpected
> plot(x = a$wt, y = a$displ,
+       xlab = "Weight",
+       ylab = "displ",
+       main = "Weight vs displ",
+       col = "green")
+ )
+ )
```



```
Untitled - R Editor  
#5  
min(a$wt)  
max(a$wt)  
  
##3rd  
#1  
plot(x = a$wt, y = a$disp,  
      xlab = "Weight",  
      ylab = "disp",  
      main = "Weight vs disp",  
      col="green")  
}  
#2  
plot(x = a$mpg, y = a$hp,  
      xlab = "mpg",  
      ylab = "hp",  
      main = "mpg vs hp",  
      col="red")  
}
```





```
##3rd
#1
plot(x = a$wt, y = a$displ,
     xlab = "Weight",
     ylab = "displ",
     main = "Weight vs displ",
     col="green")
#2
plot(x = a$mpg, y = a$hp,
     xlab = "mpg",
     ylab = "hp",
     main = "mpg vs hp",
     col="red")
#5
library(ggplot2)
ggplot(a, aes(hp, fill=factor(vs))) +
  geom_histogram(aes(y=..density..), alpha=0.6, position="identity")
```

