# Interactive debugging with Environment Browser

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## Overview of base::browser()

- ► This is a brief overview of base::browser() and how to use it for debugging in R.
- browser() is default function found in R.
- Useful tool to "hack into" R functions (that you or other people wrote)

# Typical day in lab

#### head(iris)

```
Sepal.Length Sepal.Width Petal.Length Petal.Width Species
##
## 1
             5.1
                         3.5
                                       1.4
                                                  0.2 setosa
## 2
             4.9
                         3.0
                                       1.4
                                                  0.2 setosa
## 3
             4.7
                         3.2
                                       1.3
                                                  0.2 setosa
## 4
             4.6
                        3.1
                                      1.5
                                                  0.2 setosa
## 5
             5.0
                         3.6
                                      1.4
                                                  0.2 setosa
## 6
             5.4
                         3.9
                                       1.7
                                                  0.4
                                                        setosa
```

Task: Researcher's ruler was off by 5 inches. Let's adjust it!

## Oops

```
# Loop through each column and add 5
df <- iris
for(j in 1:ncol(df)){
   df[,j] <- df[,j] + 5
}</pre>
```

## Warning in Ops.factor(df[, j], 5): '+' not meaningful for factors

```
unlist(lapply(iris, class))
```

```
## Sepal.Length Sepal.Width Petal.Length Petal.Width Species
## "numeric" "numeric" "numeric" "factor"
```

▶ Forgot that Species column is not numeric!

## Quick and dirty solution

```
# Loop through each column EXCEPT the last one
df <- iris
for(j in 1:(ncol(df)-1)){
  df[,j] <- df[,j] + 5
}
head(df)</pre>
```

```
##
    Sepal.Length Sepal.Width Petal.Length Petal.Width Species
            10.1
                        8.5
                                     6.4
                                                5.2 setosa
## 1
## 2
             9.9
                        8.0
                                     6.4
                                                5.2 setosa
             9.7
                        8.2
                                     6.3
                                                5.2 setosa
## 3
             9.6
                       8.1
                                     6.5
                                                5.2 setosa
## 4
## 5
            10.0
                       8.6
                                    6.4
                                                5.2 setosa
                                     6.7
                                                5.4 setosa
## 6
            10.4
                        8.9
```

# We got lucky

- ▶ Not all error messages are obvious
- ▶ Not all bugs come from such easy code
- Not all bugs are this easy to fix

# More likely scenario

```
foo_1 <- function(x) { ... }
foo_2 <- function(x) { ... }
# Code that produces some error
for(j in 1:ncol(df)){
  foo_1(df[,j]); foo_2(df[,j])
}</pre>
```

```
foos <- function(x) {</pre>
  foo 1(foo 2(x))
super_foo <- function(df) {</pre>
  # Code that produces some error
  for(j in 1:ncol(df)){
    foos(df[,j])
super_foo(iris)
```

#### browser() to the rescue

```
for(j in 1:ncol(df)){
  browser() <-----
foo_1(df[,j])
  foo_2(df[,j])
}</pre>
```

#### Example

```
add five <- function(x) \{x + 5\}
df <- iris
outside var <- "I'm outside the loop"
for(j in 1:ncol(df)){
  print(paste("start of the iteration:", j))
  inside_var <- "I'm inside the loop"
  browser()
  df[,j] <- add_five(df[,j])</pre>
  print(paste("end of the interation:", j))
```

#### Trigger browser() mode

#### Run your code

```
> add_five <- function(x) {x + 5}
> df <- iris
> outside_var <- "I'm outside the loop"
> for(j in l:ncol(df)){
        print(paste("start of the iteration:", j))
        inside_var <- "I'm inside the loop"
        browser()
        df[,j] <- add_five(df[,j])
        print(paste("end of the interation:", j))
        + j
        | [1] "start of the iteration: 1"
        called from: top level
        Browsef1)>
```

#### ▶ Run expressions within browser() mode

```
Browse[1]> outside_var: i: inside_var
[1] "I'm outside the loop"
[1] 1
[1] "I'm inside the loop"
Browse[1]> i+1
[1] 2
Browse[1]> head(df)
  Sepal.Length Sepal.width Petal.Length Petal.width Species
                                  1.4
           5.1
                       3.5
                     3.0
3.2
3.1 1.5
2.6 1.4
1.7
                                                0.2 setosa
           4.7
                                                0.2 setosa
           4.6
                                                0.2 setosa
           5.0
                                               0.2 setosa
           5.4
                                               0.4 setosa
Browse[1]>
```

#### What can I do in browser() mode?

► Type help for list of avaiable commands

```
Browse[1]> help next sepinto finish c or cont quit where show stack help sews later the prowse[1]> evaluate expression Browse[1]> expression Browse[1]> expression Browse[1]> expression Browse[1]> help next step sepinton Browse[1]> help sepinton B
```

#### 'c': execute all of the function

Called from: top level Browse[1]>

Browse[1]>

Executes everything left (until the next browser() call that is)
srowse[1] c
[1] "end of the interation: 1"
[1] "start of the iteration: 2"

```
Sanity check: has the first column of df been changed?
   Browse[1]> j; df[1:5, 1:2]; head(iris[1:5, 1:2])
     Sepal.Length Sepal.Width
            10.1
             9.9
                       3.2
            10.0
     Sepal.Length Sepal.Width
            4.9
                       3.0
             4.7
                       3.2
             4.6
                       3.1
             5.0
                       3 6
```

#### 'n': Go to the next line

Moves to beginning of the next line

```
Browse[1]> n
debug at #5: df[, i] <- add_five(df[, i])
Browse[1]> j; df[1:5, 1:2]; head(iris[1:5, 1:2])
  Sepal.Length Sepal.Width
          10.1
           9.9
           9.7
           9.6
          10.0
  Sepal.Length Sepal.Width
           5.1
                       3.0
                       3.2
           4.6
                       3.1
           5.0
```

Also executes your current line (before moving to next ine)

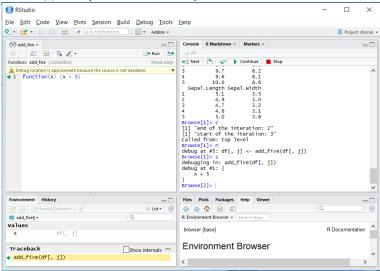
```
Browse[1]> n
debug at #6: print(paste("end of the interation:", j))
Browse[1]> j; df[1:5, 1:2]; head(iris[1:5, 1:2])
  Sepal.Length Sepal.Width
          10.1
                        8.5
           9.9
                        8.0
                        8.2
           9.6
          10.0
  Sepal.Length Sepal.Width
2
           4.9
                        3.0
3
           4.7
                        3.2
           4.6
                        3.1
           5.0
                        3.6
Browse[1]>
```

#### 's': R-ception

► Go into the function call

#### In RStudio

GUI supports (for function calls)



#### Access everything inside the function environment

Access variables as the function sees it

```
debug at #5: df[, i] <- add_five(df[, i])
Browse[1]> s
debugging in: add_five(df[, j])
debug at #1: {
   \bar{x} + 5
Browse[2]> x
 [1] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 1.5 1.6 1.4 1.1 1.2 1.5 1.3 1.4 1.7 1.5 1.7 1.5 1.0 1.7 1.9 1.6 1.6 1.5
[29] 1.4 1.6 1.6 1.5 1.5 1.4 1.5 1.2 1.3 1.4 1.3 1.5 1.3 1.3 1.6 1.9 1.4 1.6 1.4 1.5 1.4 4.7 4.5 4.9 4.0 4.6 4.5
[57] 4.7 3.3 4.6 3.9 3.5 4.2 4.0 4.7 3.6 4.4 4.5 4.1 4.5 3.9 4.8 4.0 4.9 4.7 4.3 4.4 4.8 5.0 4.5 3.5 3.8 3.7 3.9 5.1
[85] 4.5 4.5 4.7 4.4 4.1 4.0 4.4 4.6 4.0 3.3 4.2 4.2 4.2 4.3 3.0 4.1 6.0 5.1 5.9 5.6 5.8 6.6 4.5 6.3 5.8 6.1 5.1 5.3
[113] 5.5 5.0 5.1 5.3 5.5 6.7 6.9 5.0 5.7 4.9 6.7 4.9 5.7 6.0 4.8 4.9 5.6 5.8 6.1 6.4 5.6 5.1 5.6 6.1 5.6 5.5 4.8 5.4
[141] 5.6 5.1 5.1 5.9 5.7 5.2 5.0 5.2 5.4 5.1
Browse[2]> df[,3]
 [1] 1.4 1.4 1.3 1.5 1.4 1.7 1.4 1.5 1.4 1.5 1.5 1.6 1.4 1.1 1.2 1.5 1.3 1.4 1.7 1.5 1.7 1.5 1.0 1.7 1.9 1.6 1.6 1.5
[29] 1.4 1.6 1.6 1.5 1.5 1.4 1.5 1.2 1.3 1.4 1.3 1.5 1.3 1.3 1.3 1.6 1.9 1.4 1.6 1.4 1.5 1.4 4.7 4.5 4.9 4.0 4.6 4.5
 [57] 4.7 3.3 4.6 3.9 3.5 4.2 4.0 4.7 3.6 4.4 4.5 4.1 4.5 3.9 4.8 4.0 4.9 4.7 4.3 4.4 4.8 5.0 4.5 3.5 3.8 3.7 3.9 5.1
[85] 4.5 4.5 4.7 4.4 4.1 4.0 4.4 4.6 4.0 3.3 4.2 4.2 4.2 4.3 3.0 4.1 6.0 5.1 5.9 5.6 5.8 6.6 4.5 6.3 5.8 6.1 5.1 5.3
[113] 5.5 5.0 5.1 5.3 5.5 6.7 6.9 5.0 5.7 4.9 6.7 4.9 5.7 6.0 4.8 4.9 5.6 5.8 6.1 6.4 5.6 5.1 5.6 6.1 5.6 5.5 4.8 5.4
[141] 5.6 5.1 5.1 5.9 5.7 5.2 5.0 5.2 5.4 5.1
Browse[2]>
```

# Effectively use browser(): #1

```
add_five <- function(x) {x + 5}

df <- iris

for(j in 1:ncol(df)){
   if(j == 5) {
      browser()
   }
   df[,j] <- add_five(df[,j])
}</pre>
```

# Effectively use browser(): #2

#### Look at individual group

#### Pass data to browser()

```
> library(dplyr)
> df <- iris
> df %>%
+ group by(Species) %>%
         mutate(avg_sepal_length = list(Species = Species, Sepal.Length = Sepal.Length) %>% browser())
called from: function_list[[k]](value)
Browse[1]> .
Species
 [i] setosa setosa
[15] setosa setosa
[29] setosa setosa
[43] setosa setosa setosa setosa setosa setosa setosa setosa
Levels: setosa versicolor virginica
$Sepal.Length
  [1] 5.1 4.9 4.7 4.6 5.0 5.4 4.6 5.0 4.4 4.9 5.4 4.8 4.8 4.3 5.8 5.7 5.4 5.1 5.7 5.1 5.4 5.1 4.6 5.1 4.8
[26] 5.0 5.0 5.2 5.2 4.7 4.8 5.4 5.2 5.5 4.9 5.0 5.5 4.9 4.4 5.1 5.0 4.5 4.4 5.0 5.1 4.8 5.1 4.6 5.3 5.0
Browse[1]> n
Called from: function_list[[k]](value)
Browse[1]> .
$Species
  [1] versicolor versico
[10] versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor
[19] versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor
[28] versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor
[37] versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor versicolor
[46] versicolor versicolor versicolor versicolor versicolor
Levels: setosa versicolor virginica
```

#### \$Sepal.Length

[1] 7.0 6.4 6.9 5.5 6.5 5.7 6.3 4.9 6.6 5.2 5.0 5.9 6.0 6.1 5.6 6.7 5.6 5.8 6.2 5.6 5.9 6.1 6.3 6.1 6.4 [26] 6.6 6.8 6.7 6.0 5.7 5.5 5.5 5.8 6.0 5.4 6.0 6.7 6.3 5.6 5.5 5.5 6.1 5.8 5.0 5.6 5.7 5.7 6.2 5.1 5.7

#### tl;dr

 browser() allows you to interactively code in any environment scope during R execution