

# Spring 2022 courses

## DONE ds Quiz 4

### Settings

- After the first play, the quiz will be opened for unlimited play
- Let me know if you have any comments or corrections

### Which program executes an R program?

Tip: an R program or script has the file type .R

TRUE:

- Rscript
- R CMD Batch

FALSE:

- Rterm
- Rgui
- R

### In the code example, which variables are local?

```
oddcounr <- function(z=0) {  
  k <- 0  
  for (n in x) {  
    if (n %% 2 == 1) k <- k+1  
  }  
  p << 1  
  return(k)  
}
```

TRUE:

- n
- k

FALSE:

- x
- n in x

Feedback: variables that are defined inside a function are not known outside of it. x is an argument variable that must exist outside for the function to be used. If they were, they'd be global variables. To make a variable global from within a function, you can use the super-assignment operator <<-.

### Be the interpreter!

The function `oddcount` is defined in the code block below. What is the output of the last command, `oddcount()`?

```
oddcount <- function(x=0) {
  k <- 0
  for (n in x) {
    if (n %% 2 == 1) k <- k+1
  }
  return(k)
}
oddcount()
```

TRUE:

- 0

FALSE:

- 1
- ERROR
- <bytecode: 0x00000000051a7cf8>

Feedback: If no argument is given, the default value 0 is returned. The bytecode is the byte-compiled version of the function. These versions run faster than the non-compiled versions. `oddcount` is not byte-compiled, but the built-in functions like `mean()` are. Test it by entering `mean` without an argument. You may know the bytecode concept from Java whose bytecode is executed by the Java virtual machine.

## Argument list with `args()`

`ggplot` is a function in the `ggplot2` package, which is not built-in. When will `args(ggplot)` return the list of arguments?

TRUE:

- When `ggplot2` is installed and loaded

FALSE:

- When `ggplot2` is installed
- When `dev.list() == NULL`
- Never, because `args()` only works for built-in functions like `mean`

## Complete the code ??? to return the output:

```
speed <- c("medium", "slow", "fast", "fast", "fast")
f_speed <- factor(speed,
                  ordered = TRUE,
                  levels = ???)
summary(f_speed)
```

Output:

```
slow medium  fast
  1       1     3
```

TRUE:

- `c("slow", "medium", "fast")`

FALSE:

- `c("fast", "medium", "slow")`
- `c("slow", "medium")`
- `c("fast", "fast", "fast")`

Feedback: you can create ordered factors with `factor()` by setting the `ordered` and `level` args. `summary()` is a generic function. It was introduced in the "Introduction to R" DataCamp course.

## Be the interpreter!

The function call below should print: "Hello, Marcus Birkenkrahe !" But it returns this error instead:

```
: Error: unexpected symbol in "hello2(fname="Marcus" lname"
```

Can you fix the code below?

```
hello2 <- function(fname, lname) {  
  print(paste("Hello, ", fname, lname, "!"))  
}  
hello2(fname="Marcus" lname="Birkenkrahe")
```

```
Error: unexpected symbol in "hello2(fname="Marcus" lname"
```

TRUE:

- The function call is missing a comma between the two arguments

FALSE:

- User-defined functions can only have one argument
- The arguments "Marcus" and "Birkenkrahe" should not be named in the function call
- The function call to `hello2()` contains the unexpected symbol =

## Which command below returns the average of the vector `foo`?

```
foo <- c(1,3,4,100,NA,5,NA)
```

TRUE:

- `mean(foo, na.rm=TRUE)`

FALSE:

- `mean(foo, rm.na=TRUE)`
- `average(foo)`

- `mean(foo)`

## Assign the output to the correct command!

Tip: Nile is a time series of 100 elements.

---

```
Nile[Nile > 1200 & Nile < 1250] 1210 1230 1210 1220
which(Nile > 1200 & Nile < 1250) 4 8 22 26
```

---

## Which plotting command belongs to which plot type?

---

|                                  |                           |
|----------------------------------|---------------------------|
| <code>plot(data)</code>          | Scatter plot              |
| <code>hist(data)</code>          | Histogram                 |
| <code>boxplot(data)</code>       | Boxplot                   |
| <code>plot(density(data))</code> | Density distribution plot |

---

## Extract column vectors

ToothGrowth contains the factor vector `supp`, which indicates if a test subject received Vitamin C ("VC") or Orange Juice ("OJ"). The data frame looks like this:

```
'data.frame':      60 obs. of  3 variables:
 $ len : num  4.2 11.5 7.3 5.8 6.4 10 11.2 11.2 5.2 7 ...
 $ supp: Factor w/ 2 levels "OJ","VC": 2 2 2 2 2 2 2 2 2 2 ...
 $ dose: num  0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 0.5 ...
```

Which command confirms that 30 subjects received Vitamin C?

TRUE:

- `sum(ToothGrowth$supp=="VC")`
- `sum(ToothGrowth[,2]=="VC")`
- `length(ToothGrowth[ToothGrowth=="VC"])`
- `length(which(ToothGrowth=="VC"))`

Author: Marcus Birkenkrahe

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