Lazy Evaluation Data Analysis with R and Python

K 7

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• Consider the following function

```
choose1 <- function(u, a, b) {
   if (u < 0.5) a else b
}</pre>
```

• Chooses and returns one of two arguments depending on a third argument

- Chooses and returns one of two arguments depending on a third argument
- Could be used for treatment randomization

```
choose1(rbinom(1, size = 1, prob = 0.3), "placebo", "treatment")
[1] "placebo"
 choose1(rbinom(1, size = 1, prob = 0.3), "placebo", "treatment")
[1] "placebo"
 choose1(rbinom(1, size = 1, prob = 0.3), "placebo", "treatment")
[1] "placebo"
 choose1(rbinom(1, size = 1, prob = 0.3), "placebo", "treatment")
[1] "treatment"
```

• How does this function work?

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 - o R matches and evaluates the arguments u, a, and b
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- How does this function work?
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 - The function is called with some arguments
 - o R matches and evaluates the arguments u, a, and b
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- Does this sound reasonable?
- How can we check? Code demo
- What happens in Python?

- Suppose we have the following problem:
 - Find all subsets of size 2 from a set of size 5

• Easy to create all *ordered* pairs using the expand.grid() function

```
g <- expand.grid(a = 1:5, b = 1:5)
g
```

```
a b
   1 1
   2 1
  3 1
  4 1
   5 1
  1 2
   2 2
  3 2
  4 2
10 5 2
11 1 3
12 2 3
13 3 3
14 4 3
```

- But we want to retain only the $\binom{5}{2}$ distinct combinations of size 2
- Also not difficult:

```
g$b > g$a  # find the rows for which $a > b$

[1] FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE TRUE TRUE FALSE FALSE
[15] FALSE TRUE TRUE TRUE FALSE FALSE TRUE TRUE TRUE FALSE
```

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```
g$b > g$a
           # find the rows for which $a > b$
 [1] FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE FALSE TRUE TRUE FALSE FALSE
[15] FALSE TRUE TRUE TRUE FALSE FALSE TRUE TRUE
 g[g$b > g$a, ]
                # Then use this as a row index
  a b
 1 2
11 1 3
12 2 3
16 1 4
17 2 4
18 3 4
21 1 5
22 2 5
23 3 5
24 4 5
```

- As noted before, referring to g multiple times not ideal
- We can avoid doing this using the eval() function

• We can replace

```
g$b > g$a
```

```
[1] FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE TRUE TRUE TRUE FALSE FALSE
[15] FALSE TRUE TRUE TRUE FALSE FALSE TRUE TRUE TRUE FALSE
```

• by

```
eval(quote(b > a), g)
```

```
[1] FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE TRUE TRUE TRUE FALSE FALSE
[15] FALSE TRUE TRUE TRUE FALSE FALSE TRUE TRUE TRUE FALSE
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[1] FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE TRUE TRUE TRUE FALSE FALSE
[15] FALSE TRUE TRUE TRUE FALSE FALSE TRUE TRUE TRUE FALSE
```

• A general function that makes this even simpler is the with() function

```
with(g, b > a)
```

```
[1] FALSE FALSE FALSE FALSE TRUE FALSE FALSE FALSE TRUE TRUE TRUE FALSE FALSE
[15] FALSE TRUE TRUE TRUE FALSE FALSE TRUE TRUE TRUE TRUE FALSE
```

• We might guess that this is done using substitute()

- We might guess that this is done using substitute()
- We can verify this by looking at the with function by typing its name

with

```
function (data, expr, ...)
UseMethod("with")
<bytecode: 0x7f8779759448>
<environment: namespace:base>
```

- Actually this just tells us that with() is a *generic* function
- The *method* we want is the default method

```
getS3method("with", "default")

function (data, expr, ...)
eval(substitute(expr), data, enclos = parent.frame())
<bytecode: 0x7f8779758aa8>
<environment: namespace:base>
```

• All of this can now be combined to get all the $\binom{5}{2}$ distinct 2-subsets

```
g[with(g, b > a), ]
   a b
6 1 2
11 1 3
12 2 3
16 1 4
17 2 4
18 3 4
21 1 5
22 2 5
23 3 5
24 4 5
```

- This is a very common use case: obtaining a subset a dataset
- There is an even simpler way using the subset() function

```
subset(g, b > a)
   a b
  1 2
11 1 3
12 2 3
16 1 4
17 2 4
18 3 4
21 1 5
22 2 5
23 3 5
24 4 5
```

• A popular add-on package **dplyr** has a similar function called **filter()**

```
dplyr::filter(g, b > a)
  a b
  1 2
  1 3
  2 3
  2 4
  3 4
  1 5
  2 5
  3 5
10 4 5
```

• A popular add-on package **dplyr** has a similar function called filter()

```
dplyr::filter(g, b > a)
   a b
  1 2
  1 3
  2 3
  2 4
  3 4
  1 5
  2 5
  3 5
10 4 5
```

- Here dplyr::filter is the namespace notation package::function
- Allows us to use the function without attaching the whole package
- Similar to the . accessor in Python

- The **dplyr** package is very useful for routine data manipulation
- We will discuss it again later
- The idea of quoted expressions and *non-standard evaluation* is essential in many other contexts