

# Teste com condigo em R

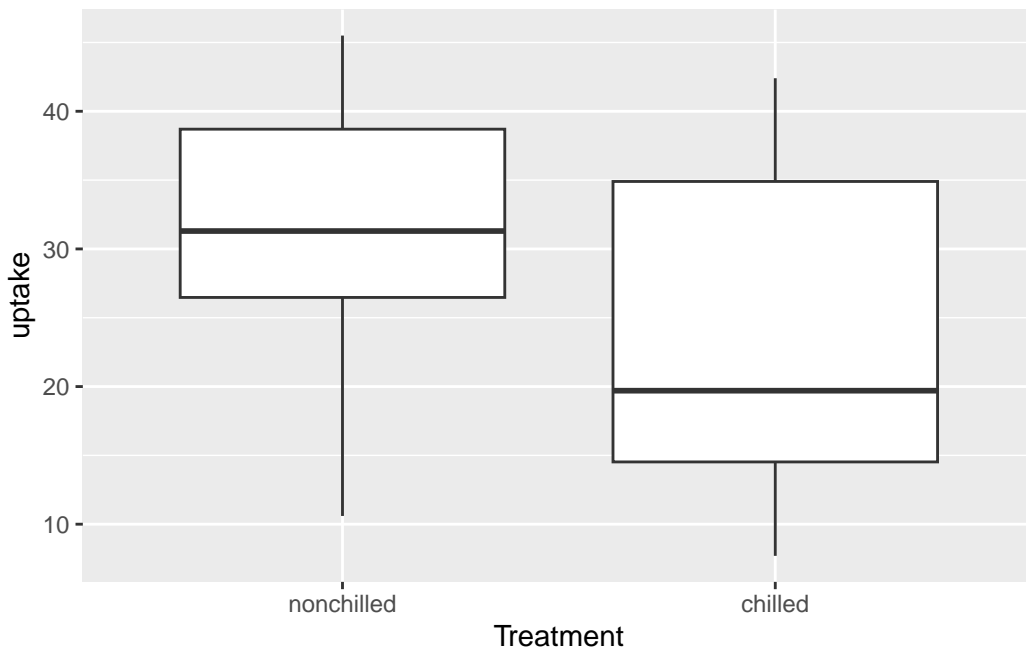
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**Este é um codigo simples em R para testar a executão do quarto**

```
library(ggplot2)

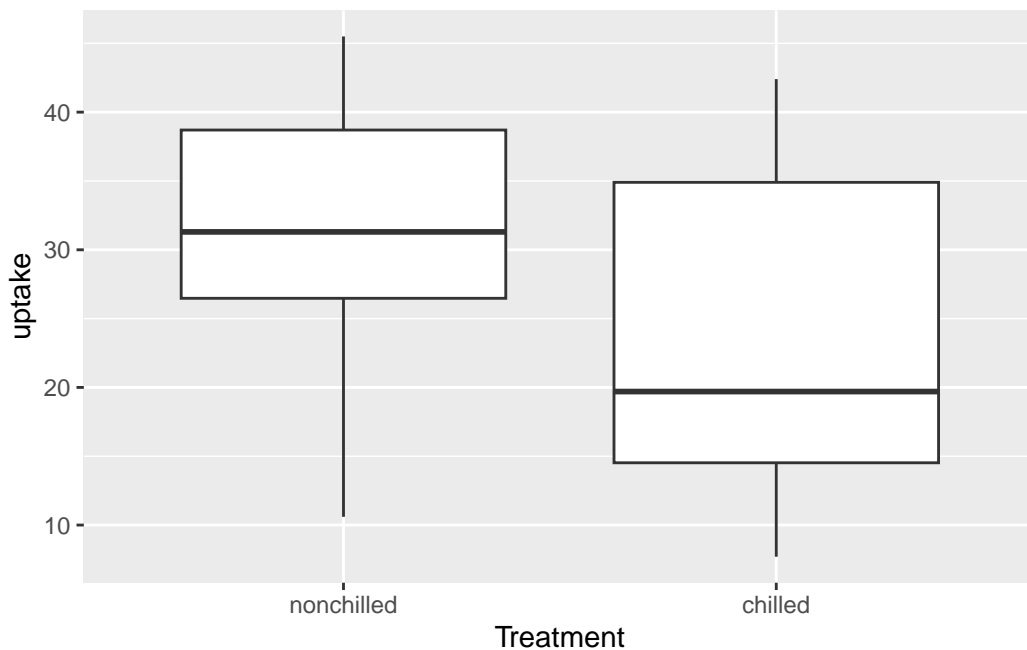
ggplot(C02, aes(Treatment, uptake)) +
  geom_boxplot()
```



```
#library(ggplot2)
summary(CO2)
```

Plant	Type	Treatment	conc	uptake
Qn1 : 7	Quebec :42	nonchilled:42	Min. : 95	Min. : 7.70
Qn2 : 7	Mississippi:42	chilled :42	1st Qu.: 175	1st Qu.:17.90
Qn3 : 7			Median : 350	Median :28.30
Qc1 : 7			Mean : 435	Mean :27.21
Qc3 : 7			3rd Qu.: 675	3rd Qu.:37.12
Qc2 : 7			Max. :1000	Max. :45.50
(Other):42				

```
ggplot(CO2, aes(Treatment, uptake)) +
  geom_boxplot()
```



por algum motivo infelizmente eu não consigo rodar o ggplot2 apenas chamando a biblioteca mãe, como o tidyverse. acontece um erro na execução

```
show("hello world")
```

```
library(tidyverse)
```

```
summary(CO2)
```

```
ggplot(CO2, aes(Treatment, uptake))  
  geom_boxplot()
```

info\_2024/posts/post-with-code/index.qmd -no-browser -no-watch-inputs ERROR: Error reading metadata from /home/daniel/ATA\_bioinfo\_2024/posts/post-with-code/index.qmd.

can not read a block mapping entry; a multiline key may not be an implicit key at line 15, column 1:

ERROR: YAMLError: can not read a block mapping entry; a multiline key may not be an implicit key at line 15, column 1:

Stack trace:

```
at generateError (file:///opt/quarto/bin/quarto.js:10480:12)  
at throwError (file:///opt/quarto/bin/quarto.js:10483:11)  
at readBlockMapping (file:///opt/quarto/bin/quarto.js:11107:24)  
at composeNode (file:///opt/quarto/bin/quarto.js:11299:84)  
at readDocument (file:///opt/quarto/bin/quarto.js:11413:5)  
at loadDocuments (file:///opt/quarto/bin/quarto.js:11448:9)  
at load (file:///opt/quarto/bin/quarto.js:11453:23)  
at parse2 (file:///opt/quarto/bin/quarto.js:11463:12)  
at parseWithNiceErrors (file:///opt/quarto/bin/quarto.js:19695:16)  
at readYamlFromMarkdown (file:///opt/quarto/bin/quarto.js:19635:26)
```