

Advanced R

Day 3

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Course Content - Advanced R (Day 3)

- ▶ Long and wide format

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- ▶ Function in R

Wide and long

Long/Wide format

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Long/Wide format

- ▶ A dataset can be written in two different formats: wide and long
- ▶ A **wide format** has one line for each patient/animal + i.e., column with unique identifier has only unique entries
- ▶ A **long format** can have more more than one line for each patient/animal + i.e., column with unique identifier has recurring entries

Long/Wide format - Example 1

| id | age | weight |
|----|-----|--------|
| P1 | 224 | 67 |
| P2 | 31 | 63 |
| P3 | 50 | 81 |
| P4 | 26 | 88 |

Long/Wide format - Example 1

| id | parameter | value |
|----|-----------|-------|
| P1 | age | 224 |
| P1 | weight | 67 |
| P2 | age | 31 |
| P2 | weight | 63 |
| P3 | age | 50 |
| P3 | weight | 81 |
| P4 | age | 26 |
| P4 | weight | 88 |

Long/Wide format - Example 2

| id | sys1 | sys2 |
|----|------|------|
| P1 | 120 | 125 |
| P2 | 118 | 125 |
| P3 | NA | 110 |

Long/Wide format - Example 2

| id | parameter | value |
|----|-----------|-------|
| P1 | sys1 | 120 |
| P1 | sys2 | 125 |
| P2 | sys1 | 118 |
| P2 | sys2 | 125 |
| P3 | sys1 | NA |
| P3 | sys2 | 110 |

Long/Wide format - why?

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- ▶ long format
 - repeated measurements
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 - can be more efficient in terms of storage space

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- ▶ each format has advantages and is more useful for certain tasks
- ▶ long format
 - repeated measurements
 - ▶ especially if number of repetitions differ per patient/animal
 - ▶ long format needed for plots over repeated measurements
 - can be more efficient in terms of storage space
- ▶ wide format
 - easier to read and look up a patient/animal
 - often easier to make calculations (e.g., BMI)

Long/Wide format - reshape

in R package *tidyr*

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Long/Wide format - reshape

in R package *tidyr*

- ▶ **pivot_wider()** “widens” data, increasing the number of columns and decreasing the number of rows
- ▶ **pivot_longer()** “lengthens” data, increasing the number of rows and decreasing the number of columns

Example - pivot_wider()

```
dt_example <-  
  tibble(id = rep(paste0("P", 1:3), 2),  
         sys = c(120,118,NA, 125,125,110),  
         measurement = c(1, 1, 1, 2, 2, 2))  
  
dt_example %>%  
  pivot_wider(data = ., names_from = measurement, values_from = sys)
```

Example - pivot_longer()

```
dt_example <-  
  tibble(id = paste0("P", 1:4),  
         age = c(224, 31, 50, 26),  
         weight = c(67, 63, 81, 88))  
  
dt_example %>%  
  pivot_longer(data = ., cols = c(age, weight))
```

Functions in R

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```

- ▶ a function returns a 'value' (e.g., value, ggplot, ...)

Example `mean()`

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`mean()` - calculate the mean value of a vector

- ▶ `mean()` has several arguments
- ▶ `x`: numeric/logical vectors (others also possible, e.g. time intervals)
- ▶ `trim`: the fraction (0 to 0.5) of observations to be trimmed from each end of `x` before the mean is computed
- ▶ `na.rm`: a logical evaluating to TRUE or FALSE indicating whether NA values should be stripped before the computation proceeds
- ▶ `...` : further arguments passed to or from other methods

```
mean(x, trim = 0, na.rm = FALSE, ...)
```

Write your own function

```
myFunction <- function() {  
  
}
```

- ▶ object *myFunction* is now a function

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```

- ▶ object *myFunction* is now a function
- ▶ `()` → currently we have no arguments
- ▶ `{ }` is the place for all commands we want

Write your own function

```
myFunction <- function(name) {  
  greetings <- paste0("Hallo, ", name, "!")  
  cat(greetings)  
}
```

- ▶ *name* is our argument
- ▶ {} creates the *greetings* and returns the last line

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```

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```
myFunction(name = "Susi")
```

```
## Hallo, Susi!
```

Write your own function - Example

```
CreateSampleDataset <- function(nrow = 100) {  
  Condition <- rbinom(n = nrow, size = 1, prob = 0.5)  
  IQ <- rnorm(n = nrow, mean = 100, sd = 15)  
  Age <- rnorm(n = nrow, mean = 40, sd = 7.5)  
  Motivation <- runif(n = nrow, min = 1, max = 10)  
  dfSampleData <- tibble(Condition, IQ, Age, Motivation)  
  return(dfSampleData)  
}
```

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```
myFunction2 <- function(x = 0) {  
  x-3  
  return(c(answer = 42))  
}
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  return(c(answer = 42))  
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```

```
myFunction2()
```

```
## answer  
##      42
```

Write your own function - Example

```
myFunction2 <- function(x = 0) {  
  x-3  
  return(c(answer = 42))  
}
```

```
myFunction2()
```

```
## answer  
##      42
```

► *return()*

- variable can be any R object
- a function in R can only ever return one object (use `list()` for more)

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- ▶ Functions are used when the same or similar program code is used in several places in the script
- ▶ Variables declared inside a function are local to that function
- ▶ Function name should make it clear what the function is good for
- ▶ Inputs and outputs should be clear

Links

Links (I)

- ▶ Introduction to R
 - R for Data Science (<https://r4ds.hadley.nz/>)
- ▶ Plots using ggplot
 - Overview with further links to course material: <https://ggplot2.tidyverse.org/>
- ▶ Display tables using flextable
 - flextable bool <https://ardata-fr.github.io/flextable-book/>
 - Function references <https://davidgohel.github.io/flextable/reference/index.html>
- ▶ `knit_child()`
 - link (<https://bookdown.org/yihui/rmarkdown-cookbook/child-document.html>)

Links (II)

- ▶ Download R
 - CRAN (<https://cran.r-project.org/>)
- ▶ Download RStudio
 - RStudio Desktop (<https://posit.co/download/rstudio-desktop/>)