# Introductory Workshop in Bioinformatics - Part 2

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#### Setting up RStudio / Installing required packages

```
install.packages("tidyverse")

### Warning: Paket 'tidyverse' wird gerade benutzt und deshab nicht installiert

install.packages("ggpubr")

### Warning: Paket 'ggpubr' wird gerade benutzt und deshab nicht installiert
```

#### Download files required for this tutorial

- Go to: https://github.com/jentiger82/BioinformaticsWorkshop2022
- · Download all files and save them in an easy to find folder

#### Getting to know R

- Run: library(tidyverse)
- For the first exercise load files SEQ1.txt and SEQ2.txt
- You can replace the web address below with your local file path to the files:
   "D:/Bioinformatics\_Seminar\_2022/SEQ1.txt"

```
seq1 <- read_file(
   "https://raw.githubusercontent.com/jentiger82/BioinformaticsWorkshop2022/main/SEQ1.txt")
seq2 <- read_file(
   "https://raw.githubusercontent.com/jentiger82/BioinformaticsWorkshop2022/main/SEQ2.txt")
seq_1 <- read_file("D:/Bioinformatics_Seminar_2022/SEQ1.txt")</pre>
```

## We want to know if sequence 1 is the same as sequence 2

```
#This is a comment.
#Printing seg1 and seg2
seq1
## [1] "agagacacagccgaaccccaggcagttttggccggtttgccaggttccccgttaccgtcctt"
seq2
## [1] "agagacacagccgaaccccaggcagttttggccgctttgccaggttccccgttaccgtcctt"
#Are they equal?
seq1==seq2
## [1] FALSE
```

#### 5 basic data structures in R

- Character
- Numeric
- Integer
- Logical
- · Complex

#### Character

```
# Holds "string" data. For example:
"Hello World"

## [1] "Hello World"

# You can ask R to tell you what type of data you have:
class("Hello World")

## [1] "character"
```

#### **Numbers**

```
# Numbers can be integers or numerics (i.e., floats) in R. For example:
1
## [1] 1
2.2
## [1] 2.2
3E100
## [1] 3e+100
```

#### **Numbers**

```
# The default for R is to make every number a numeric
class(1)
## [1] "numeric"

# Which means that we have hidden decimal points behind the 1 (1.00 for example)
```

## But what if we wanted specifically an integer?

```
1L
## [1] 1
class(1L)
## [1] "integer"
```

```
# Is this a numeric or a character?
"2"
### [1] "2"
```

```
# Is this a numeric or a character?
"2"

## [1] "2"

class("2")

## [1] "character"
```

## Logicals

```
# Logicals are either True of False. For example
TRUE
## [1] TRUE
FALSE
## [1] FALSE
class(TRUE)
## [1] "logical"
class(FALSE)
## [1] "logical"
```

```
# R is *case sensitive*.
# With that knowledge, what do you think the below would give us?
#class(true)
```

Output: Error: object 'true' not found

```
# R is *case sensitive*.
# With that knowledge, what do you think the below would give us?
#class(true)
```

#### Data Structure Classes cont...

```
# So far we have only been using class()
# But we can also use a function to see the structure of any R object:
str(1)
## num 1
str("Hello World!")
## chr "Hello World!"
str(TRUE)
   logi TRUE
```

```
# R is an object oriented language, and it tends to put everything in some sort of class.
# What would the below tell us?
Str(class)
```

```
# R is an object oriented language, and it tends to put everything in some sort of class.
# What would the below tell us?
str(class)
## function (x)
```

## Simple Data Structures Methods

#### **Arithmetic Methods**

```
# * Addition
1+1
## [1] 2
# * Subtraction
1-1
## [1] 0
# * Multiplication
2*2
## [1] 4
# * Division
10/2
```

## [1] 5

20/41

#### **Equivalence Comparisons**

```
# Equivalence comparisons are a way to check if any two objects are the same
#Does 1 equal 1?
1==1

## [1] TRUE

#Does "Hello" equal "World?"
"Hello" == "World"

## [1] FALSE
```

## **Equivalence Comparisons**

```
#We can also invert an equivalancy comparison to check if two objects are *not* equal #Does 1 not equal 1?

1!=1
```

## [1] FALSE

## **Mathematical Comparisons**

```
#For numeric data types, mathematical comparisons can also be made
#Is 1 Less than 100?
1<100

## [1] TRUE

#Is 2+2 greater than 2^2
2+2>2^2
## [1] FALSE
```

```
# Arithmetic in R follows PEMDAS
4+5*3
## [1] 19
# vs
(4+5)*3
## [1] 27
# What would we get with the following?
4+53>(4+5)3
```

```
# Arithmetic in R follows PEMDAS
4+5*3
## [1] 19
# vs
(4+5)*3
## [1] 27
# What would we get with the following?
4+5*3>(4+5)*3
## [1] FALSE
```

#### **Variables**

```
# Variables hold objects that are assigned to them.
a <- 1
# They can be identical to the object assigned to them
a <- 1
b <- a
#does b equal 1?
b==1
## [1] TRUE</pre>
```

#### Variables enable complex operations on data

```
h <- 2^100
i <- h/3E100
j <- 1E5
k <- j^(-1*i)
#is k greater than 1?
k > 1
## [1] FALSE
```

## **Complex Data Structures**

#### **Vectors**

```
# A Vector is an ordered collection of either numerics, characters, or logicals
c(1, 2, 3)
## [1] 1 2 3
c(TRUE, FALSE, TRUE)
## [1] TRUE FALSE TRUE
c("hello", "world")
## [1] "hello" "world"
```

```
# A vector in R has a limitation: they can only allow one type of class in each vector.

# If you add more than one type of class, it will default to a character.

# With that said, what would the following give you?

class(c("puppies", 2))
```

```
class(c("puppies", 2))
## [1] "character"
```

```
# I can change the following vector in my variable to all numbers
my_num <- as.numeric(c("2", 2))
class(my_num)

## [1] "numeric"

# But what would happen if I tried to do the following?

as.numeric(c("salt", "2")) print(as.numeric(c("salt", "2")))</pre>
```

```
print(as.numeric(c("salt", "2")))

## Warning in print(as.numeric(c("salt", "2"))): NAs durch Umwandlung erzeugt

## [1] NA 2

class(as.numeric(c("salt", "2")))

## Warning: NAs durch Umwandlung erzeugt

## [1] "numeric"
```

#### **Vectors Continued**

95

##

```
# Vectors can also have a vector of names which describe each element
grades <- c(98, 95, 82)
names(grades) <- c("Jimmy", "Alice", "Susan")</pre>
grades
## Jimmy Alice Susan
      98
            95
                  82
##
# Elements from a vector can be accessed using the index of the desired data. Lets say I wanted to know wh
grades[2]
## Alice
##
      95
# But lets say that I didn't know Alice was the second index
grades["Alice"]
## Alice
```

#### **Vectors Continued**

```
# If you want to add a series of numbers to a vector of integers, you can do the following:
my_ints <- 1:10
my_ints
## [1] 1 2 3 4 5 6 7 8 9 10</pre>
```

#### Lists

```
# A list is an ordered collection of any objects. This differs from vectors, because vectors can only be o
my_list <- list(1, "b", TRUE, c(1,2,3))</pre>
# Lists can also have names
names(my list)=c("My #", "My Letter", "My logical", "My num Set")
my list
## $`My #`
## [1] 1
##
## $`My Letter`
## [1] "b"
##
## $`My logical`
## [1] TRUE
##
## $My_num_Set
## [1] 1 2 3
```

#### Lists continued

```
# The elements within a list can be accessed by using numeric indexes or by the element name
my list[[2]]
## [1] "b"
my_list[["My_num_Set"]]
## [1] 1 2 3
my list$My num Set
## [1] 1 2 3
#If I did the following, I would get the whole vector within the list
my_list[2]
## $`My Letter`
## [1] "b"
```

#### **Data Frames**

Jimmy

Alice

Susan

98

95

82

## 1

## 2

## 3

#### **Data Frames**

```
#Data frames can be accessed numerically by expressing the row and column of interest.
#What grade did Susan get?
my_df[3,2] #row, column
## [1] 82
# They can also be accessed with the $ sign
my df$Grades
## [1] 98 95 82
#What grade did Alice get?
my_df$Grades[2]
## [1] 95
```

#### Using Data Frames - Iris data

```
# How does the iris data Look Like?
#View(iris)
# Get row 1
row1 <- iris[1,]
# Get column 1
col1 <- iris[,1]
# Access a column by name
pl <- iris$Petal.Length
# How to get the value in row 1, column 2?
# How would I get the "species" in row 3?</pre>
```

We continue in 15 minutes...

#### Iris data - solutions

```
# How to get the value in row 1, column 2?
iris[1,2]

## [1] 3.5

# How would I get the "species" in row 3?
iris$Species[3]

## [1] setosa
## Levels: setosa versicolor virginica
```