Crash n' Survival Course with R

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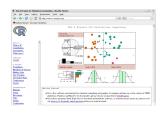
MFPL - CIBIV
Center for Integrative Bioinformatics in Vienna

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Your biggest friend and partner in statistics



Installation of R, starting and additional libraries







- ► R Webpage http://www.r-project.org/
- There is code and precompiled binaries
- Start with the command R
- ► Additional libraries which are installed are loaded with library(name)
- A usefull interface is the R commander which is started with library(Rcmdr), conatins a script and an output window

Using R as calculator: Basic Operations

```
> 5-2 # addition, subtraction
                        [1] 3
                        > 5*2 # multiplication, division
* , /
                        [1] 10
                        > 5**2 # power
^ or **
                        [1] 25
                        > 5\%/\%2 # division
%/%
                        [1] 2
                        > 5%%2 # modulo
%%
                        [1] 1
```

Basic arithmetic functions and values

predefined arithmetic functions:

```
max(), min()
abs()
sqrt()
round, floor(), ceiling()
sum(), prod()
log(), log10(), log2()
exp()
sin(), cos(), tan(), asin(), acos(), atan()
```



Basic arithmetic functions and values

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    max(), min()
    abs()
    sqrt()
    round, floor(), ceiling()
    sum(), prod()
    log(), log10(), log2()
    exp()
    sin(), cos(), tan(), asin(), acos(), atan()
```



Constant Values:

 $\begin{array}{cccc} \text{pi} & \pi & \text{pi} \\ \text{Inf, -Inf} & \text{infinity} & 1/0 \\ \text{NaN} & \text{Not a Number} & 0/0 \\ \text{NA} & \text{Not Available} & 'NA' \\ \text{NULL} & \text{Not defined} \end{array}$

Assigning values to variables

We do not need to declare the datatype of a variable in R.

The simplest datatype in R is a scalar.

variables can be used to represent values:

```
> x <- 15
> x
[1] 15
```

values can also be assigned by:

```
> a = 15
> a
[1] 15
```

Logical values:

```
> a == 15
[1] TRUE
```

A Vector is a array of values

construct a vector with the "concatenate function" c()

Scalar vector:

```
> v <- c(15, 4, 67, 5, 9)
> v
[1] 15 4 67 5 9
```

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Scalar vector:

```
> v <- c(15, 4, 67, 5, 9)
> v
[1] 15 4 67 5 9
```

Logical vector:

```
> v <- c(T, T, F, T)
> v
[1] TRUE TRUE FALSE TRUE
```

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> v <- c(15, 4, 67, 5, 9)
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[1] 15 4 67 5 9
```

Logical vector:

```
> v <- c(T, T, F, T)
> v
[1] TRUE TRUE FALSE TRUE
```

Character vector:

```
> v <- c("Hund", "Katze", "Maus", "Mensch")
> v
[1] "Hund" "Katze" "Maus" "Mensch"
```

► Single value from vector:

```
> x <- c(4, 5, 8, 23, 12)
> x[2]
[1] 5
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► Multiple values from vector:

```
> x[c(2,4)]
[1] 5 23
```

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Multiple values from vector:

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> x[c(2,4)]
```

[1] 5 23

Subsequent values from vector:

```
> x[1:3]
```

[1] 4 5 8

- Single value from vector:
 - > x <- c(4, 5, 8, 23, 12)
 - > x[2]
- Multiple values from vector:
- > x[c(2,4)]
- [1] 5 23
- ► Subsequent values from vector:
 - > x[1:3]
 - [1] 4 5 8
- ▶ Logical selection (boolean vectors of same length):
- > x>7
 - [1] FALSE FALSE TRUE TRUE TRUE
 - > x[x>7]
 - [1] 8 23 12

Operations on vectors

arithmetic operations on vectors:

```
> x <- c(4, 5, 8, 23, 12)
> x*2
[1] 8 10 16 46 24
```

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> x <- c(4, 5, 8, 23, 12)
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```

Vectors need to have the same length!

```
> x + x
[1] 8 10 16 46 24
> x + x[1:4]
[1] 8 10 16 46 16
Warning message: longer object length is not a
multiple of shorter object length in: x + x[1:4]
```

Operations on vectors

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Warning message: longer object length is not a
multiple of shorter object length in: x + x[1:4]
```

Test whether a vector contains an element

```
> c(5,6) %in% x [1] TRUE FALSE
```

Some functions on vectors

► Getting the length of a vector

```
> x <- c(4, 5, 8, 23, 12)
> length(x)
[1] 5
```

Some functions on vectors

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> x <- c(4, 5, 8, 23, 12)
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Getting and changing the names of a vector

```
> names(x)=c("E1","E2","E3","E4","E5")
> x
E1 E2 E3 E4 E5
    4    5    8    23    12
> x["E3"]
E3
    8
```

Some functions on vectors

Getting the length of a vector

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E1 E2 E3 E4 E5
   4   5   8   23   12
> x["E3"]
E3
   8
```

Getting the order of a vector

```
> o=order(x)
> o
[1] 1 2 3 5 4
> sort(x) #same as x[o]
[1] 4 5 8 12 23
```

Matrices are represented as vectors with two dimensions

We can set or change dimensions with the function dim

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Matrices can be created using the matrix function

(here the Matrix will be filled in a rowwise fashion)

Useful functions that operate on matrices

Indexing

> MAT<-matrix(5:12,c(4,2),byrow=T)	> MAT		
> MAT[3]		[,1]	[,2]
[1] 7	[1,]	5	6
> MAT[,1] [1] 5 7 9 11	[2,]	7	8
> MAT[1,]	[3,]	9	10
[1] 5 6 > MAT[3,2]	[4,]	11	12
Γ1 ₁ 10			

Useful functions that operate on matrices Indexing

```
> MAT<-matrix(5:12,c(4,2),byrow=T)
> MAT[3]
[1] 7
> MAT[,1]
[1] 5 7 9 11
> MAT[1,]
[1] 5 6
> MAT[3,2]
[1] 10
assign rownames and colnames
> rownames(M) <- c("X","Y")
> colnames(M) <- c("A","B","C")</pre>
> M
  ABC
X 1 3 5
Y 2 4 6
> M["X",]
ABC
1 3 5
```

```
> MAT
[,1] [,2]
[1,] 5 6
[2,] 7 8
[3,] 9 10
[4,] 11 12
```

Factors: storing categorical variables

A factor consists of a vector of integers and a character vector

```
> pain <- c(0,3,2,2,1)
```

- > fpain <- factor(pain,levels=0:3)</pre>
- > levels(fpain) <- c("none", "mild", "medium", "severe")</pre>

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> pain <- c(0,3,2,2,1)
> fpain <- factor(pain,levels=0:3)
> levels(fpain) <- c("none","mild","medium","severe")</pre>
```

The level argument assures the ordering of our categories (otherwise alphabetical ordering)

```
> fpain
[1] none severe medium medium mild
Levels: none mild medium severe
```

- > as.numeric(fpain)
- [1] 1 4 3 3 2

Factors: storing categorical variables

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```
> pain <- c(0,3,2,2,1)
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The level argument assures the ordering of our categories (otherwise alphabetical ordering)

```
> fpain
[1] none severe medium medium mild
Levels: none mild medium severe
> as.numeric(fpain)
[1] 1 4 3 3 2
```

Count the categories (can also be applied to normal vectors):

```
> table(fpain)
fpain
  none mild medium severe
     1     1     2     1
```

Data frames: Lists of vectors with the same length

Creating a data frame object

4 Schnabeltier

2

Data frames: Selection

▶ get the names of a object

```
> names(dframe)
[1] "Spec" "Num"
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\$ gives the column vectors

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> dframe$Num #equals dframe[,2] and dframe[,"Num"]
[1] 3 5 7 2
```

Data frames: Selection

get the names of a object

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> names(dframe)
[1] "Spec" "Num"
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\$ gives the column vectors

```
> dframe$Num #equals dframe[,2] and dframe[,"Num"]
[1] 3 5 7 2
```

can be indexed like a matrix

```
> dframe$Num[1] # equals dframe[1,2]
[1] 7
```

Lists: composite collection of objects

creating object using list() > lobject < -list(c(1,2,2,2), age=c(98,21,56))> lobject $\lceil \lceil 1 \rceil \rceil$ [1] 1 2 2 2 \$age [1] 98 21 56 selection of particular values (Indexing) > lobject[[1]] #is a vector [1] 1 2 2 2 > lobject\$age #also a vector [1] 98 21 56 > lobject[1] #is still a list $\lceil \lceil 1 \rceil \rceil$ [1] 1 2 2 2

Testing and converting variable types

Test a special type
> is.list(lobject)
[1] TRUE
> is.list(lobject[[1]])
[1] FALSE
> is.vector(lobject[[1]])
[1] TRUE
> is.na(c(1,0/0))

[1] FALSE TRUE

Testing and converting variable types

```
Test a special type
  > is.list(lobject)
  [1] TRUE
  > is.list(lobject[[1]])
  [1] FALSE
  > is.vector(lobject[[1]])
  [1] TRUE
  > is.na(c(1,0/0))
  [1] FALSE TRUE
Convert to a special type
  > a <- as.logical(c(3,2,0,1))
  > a
  [1]
       TRUE TRUE FALSE TRUE
  > as.integer(a)
  [1] 1 1 0 1
```

Simple example how to read a table into a data frame object:

> dataFrame1 <- read.table("data.txt",header=T)</pre>

You can also write a data.frame into a file:

> write.table(data,filename)

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> write.table(data,filename)

Write R-Objects:

- > #save all the variables in the workspace to a file:
- > save(file=filename)
- > #save only the variables MAT and v to a file:
- > save(MAT, v, file=filename2)
- > load(filename) #load the data in another R session
- > #get the names of variables in the workspace:
- > ls()

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Load a script with R-commands:

> source(script)

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Get and set the working directory of R:

- > getwd()
- > setwd(dirstring)

R as a programming language

► Loops: for, while and repeat

```
> v=c(1,2,3,4,5)
> for (i in 1:5){
+ print (sum(v[1:i]))
+ }
[1] 1
[1] 3
[1] 6
[1] 10
[1] 15
```

► There is a much faster way

```
> sapply(1:5,function(i) sum(v[1:i]))
[1] 1 3 6 10 15
```

R as a programming language

▶ Loops: for, while and repeat
▶ If and else

```
> v=c(1,2,3,4,5)
> for (i in 1:5){
+ print (sum(v[1:i]))
+ }
[1] 1
[1] 3
[1] 6
Γ1 10
Γ1 15
```

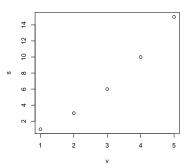
```
> if (v[3]==3){
+ v[2]=3
+ }
> v
[1] 1 3 3 4 5
```

► There is a much faster way

```
> sapply(1:5,function(i) sum(v[1:i]))
[1] 1 3 6 10 15
```

Plotting with R

```
> v = c(1,2,3,4,5)
> s = sapply(1:5,function(i)
+ sum(v[1:i]))
> plot(v,s)
```

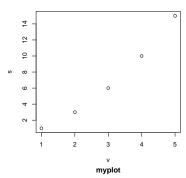


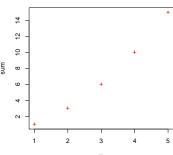
Plotting with R

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There are many parameters to modify the plot, e.g.

```
> plot(v,s,main="myplot",
```





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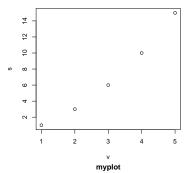
There are many parameters to modify the plot, e.g.

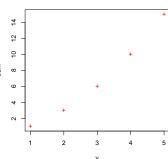
```
> plot(v,s,main="myplot",
```

There are different graphical devices, e.g.

```
x11() (screen, default)
pdf()
```

dev.off() closes a device





Statistics with R

There are many many statistical tests, distributions, ... implemented in R.

```
e.g. Fisher-Test on count data:
```

```
> l=matrix(c(2,11,15,5),2)
> 1
      [,1] [,2]
[1,] 2 15
[2,] 11 5
> fisher.test(1)
```

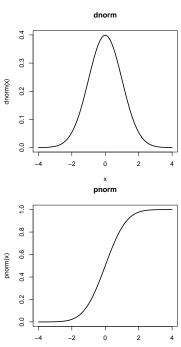
Fisher's Exact Test for Count Data

```
data: 1
p-value = 0.001268
alternative hypothesis: true odds ratio is not equal to 1
95 percent confidence interval:
    0.00547646 0.45530971
sample estimates:
odds ratio
0.06755424
```

Probability and distributions

Distributions can be obtained by:

- probability density d
- cummulative distribution p
- quantiles q
- ▶ random r

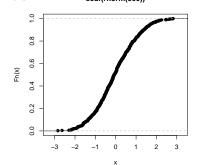


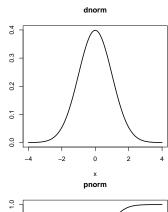
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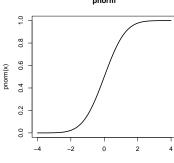
Distributions can be obtained by:

- probability density d
- cummulative distribution p
- quantiles q
- random r

Empirical cummulative distribution: ecdf(rnorm(500))







Literature:

Main source of the information and for further reading:



There are also numerous free documentations of R. In R, you can get help for a function e.g. ?sum