

Do you have a github account?  
Give us a star! :)

# Welcome!

Please download all of the course material:

---

[tinyurl.com/introRData](https://tinyurl.com/introRData) > [Clone or download ▾](#) > [Download ZIP](#)

and store it in a single, accessible folder on your computer. **Don't forget to unzip!**

# Introduction to R & data

---

- Part I: Basics of R
- Lunch (~12:15)
- Part II: Modern R with Tidyverse
- Final remarks (~17:00)

move the samples from the data  
{r}  
select the samples to keep  
keepsamples <- row.names(pheno[  
apply sample selection to  
counts.sub <- counts.sub[,keepsamples]  
pheno.sub <- pheno[keepsamples]  
mt.assay <- mt.assay[,keepsamples]  
rld.sub <- rld[,keepsamples]

# Introduce yourself!

---

- Your name & pronouns
- What is your faculty/background? (Economics, Medicine, Biology...)
- What part of your education are you in? (Bachelor, PhD, prof...)
- What is your motivation for learning R?
- What is your experience with R?

Go to [tinyurl.com/introRDatadoc](https://tinyurl.com/introRDatadoc) and add your name to the roll call!

# Asking for help

---

- Option 1: ask a direct question, in the Zoom chat
- Option 2: indicate that you're stuck, in the Status Chart

## Status chart

**GOAL: Open RStudio and the exercise file**

Name	:-)	HELP!	<i>link to breakout room [will be added by helper]</i>
Chris	X		
Sam		X	<a href="http://tinyurl.com/helpmeJacques">tinyurl.com/helpmeJacques</a>

*NB: Also use the status chart when you are done with an exercise! See GOAL.*

# Introduction to R & data

---

## Part 1: Basics of R



# What is R?

---

- Widely used programming language for data analysis
- Based on statistical programming language **S** (1976)
- Developed by **Ross Ihaka & Robert Gentleman** (1995)
- Very active community, with many (often subject-specific) packages
- Open source, interoperable!



## We will work in RStudio

---

- Integrated Development Environment for R
- Founded by JJ Allaire, available since 2010
- Bloody useful! Let's take a look: please open RStudio!

# The Rstudio interface



The screenshot displays the RStudio interface with four main panels:

- Script Panel (Top Left):** Shows an R script named "programming\_exercise.R" with code for calculating averages from the iris dataset. Overlaid text includes "script", "markdown", and "data preview".
- Environment Panel (Top Right):** Shows the Global Environment tab with the message "Environment is empty". Overlaid text includes "environment" and "history".
- Console Panel (Bottom Left):** Shows the R console output for version 3.5.2. Overlaid text includes "console" and "Natural language support but running in an English locale".
- Files Panel (Bottom Right):** Shows the Files tab with tabs for "Plots", "Packages", "Help", and "Viewer". Overlaid text includes "files", "plots", "packages", and "help".

# Have you downloaded the course material?

---

[tinyurl.com/introRData](https://tinyurl.com/introRData) > [Clone or download ▾](#) > [Download ZIP](#)

Please store it in a single, accessible folder on your computer. **Don't forget to unzip!**

# R syntax & the console

---

- Data types in R
- Using functions
- Combining data
- Indexing data

move the samples from the data  
`{r}  
select the samples to keep  
keepsamples <- row.names(pheno[  
apply sample selection to  
counts.sub <- counts.sub[,keepsamples]  
pheno.sub <- pheno[keepsamples]  
mt.assay <- mt.assay[,keepsamples]  
rlid.sub <- rlid[,keepsamples]

# How to read our slides (console version)

---

you are writing in the console

```
> x <- 1
```

*giving R information*

console output

```
> x  
[1] 1
```

*asking R a question*

index of output

*NB! Focus on the slides when we are presenting! We will have exercises, too.*

# Variable assignment

---

>

# Variable assignment

---

```
> x <- 6
```

# Variable assignment

> x <- 6

> x <- 'apple'

> x <- "hello world"

> x = 6

in cheatsheets/ folder  
(course materials)

## Variable Assignment

```
> a <- 'apple'
> a
[1] 'apple'
```

**Base R Cheat Sheet**

### Getting Help

- Accessing the help files**
  - ?mean
  - Get help of a particular function.
  - help.search('weighted mean')**
  - Search the help files for a word or phrase.
  - help(package = 'dplyr')**
  - Find help for a package.

### Creating Vectors

c(2, 4, 6)	2 4 6	Join elements into a vector
2:6	2 3 4 5 6	An integer sequence
seq(2, 3, by=0.5)	2.0 2.5 3.0	A complex sequence
rep(1:2, times=3)	1 1 2 1 2 1 2	Repeat a vector
rep(1:2, each=3)	1 1 1 2 2 2	Repeat elements of a vector

### Vector Functions

sort(x)	rev(x)
Return x sorted.	Return x reversed.
table(x)	unique(x)
See counts of values.	See unique values.

### Selecting Vector Elements

By Position	
x[4]	The fourth element.
x[-4]	All but the fourth.
x[2:4]	Elements two to four.
x[-(2:4)]	All elements except two to four.
x[c(1, 5)]	Elements one and five.
By Value	
x[x == 10]	Elements which are equal to 10.
x[x < 0]	All elements less than zero.
x[x %in% c(1, 2, 5)]	Elements in the set 1, 2, 5.
Named Vectors	
x['apple']	Element with name 'apple'.

### Conditions

a == b	Are equal	a > b	Greater than	a >= b	Greater than or equal to	is.na(a)	Is missing
a != b	Not equal	a < b	Less than	a <= b	Less than or equal to	is.null(a)	Is null

### Types

Converting between common data types in R. Can always go from a higher value in the table to a lower value.

as.logical	TRUE, FALSE, TRUE	Boolean values (TRUE or FALSE).
as.numeric	1, 0, 1	Integers or floating point numbers.
as.character	'1', '0', '1'	Character strings. Generally preferred to factors.
as.factor	'1', '0', '1', levels: '1', '0'	Character strings with preset levels. Needed for some statistical models.

### Maths Functions

log(x)	Natural log.	sum(x)	Sum.
exp(x)	Exponential.	mean(x)	Mean.
max(x)	Largest element.	median(x)	Median.
min(x)	Smallest element.	quantile(x)	Percentage quantiles.
round(x, n)	Round to n decimal places.	rank(x)	Rank of elements.
signif(x, n)	Round to n significant figures.	var(x)	The variance.
cor(x, y)	Correlation.	sd(x)	The standard deviation.

### Matrices

m <- matrix(x, nrow = 3, ncol = 3)	Create a matrix from x.
m[2, ]	Select a row
m[, 1]	Select a column
m[2, 3]	Select an element
t(m)	Transpose
m %*% n	Matrix Multiplication
solve(m, n)	solve(m, n)
find(x in m: x = n)	Find x in m: x = n

### Lists

l <- list(x = 1:5, y = c('a', 'b'))	A list is a collection of elements which can be of different types.
l[[2]]	Second element of l.
l[1]	New list with only the first element.
l\$x	Element named x.
l['y']	New list with only element named y.

### Data Frames

df <- data.frame(x = 1:3, y = c('a', 'b', 'c'))	A special case of a list where all elements are the same length.
df	df\$x
df[2]	df[[2]]

### Matrix subsetting

df[, 2]	Understanding a data frame
View(df)	See the full data frame.
head(df)	See the first 6 rows.
nrow(df)	Number of rows.
ncol(df)	Number of columns.
dim(df)	Number of columns and rows.
cbind	- Bind columns.
rbind	- Bind rows.

### Variables

plot(x, y)	Values of x in order.
plot(x, y)	Values of x against y.
hist(x)	Histogram of x.

### Dates

See the lubridate package.



# Executing R code

---

In a script/Rmarkdown document:

- Place your cursor in the line of code you want to execute
- Press  or **ctrl + enter**
- When running multiple lines: select all lines, then press ‘Run’ or **ctrl+enter**

In an Rmarkdown document: execute a *chunk* with the green triangle

```
9 ## EXERCISE 0
10
11 Try to run this code.
12
13 ``{r}
14 x <- 6
15
16 apple <- "apple"
17 ````
```



# Exercise 0: open the exercise document

---

1. File > Open Project > introduction-to-R-and-data-github.Rproj
2. From the ‘files’ menu (bottom right), select **baseR\_exercises.Rmd**
3. Run the R code in Exercise 0
4. When you are done: place an X in the green column of the status chart

## Exercise 1

---

1. Do the following calculation in R:

$$\frac{1 + 5}{9}$$

2. Assign the result of the calculation to a variable.
3. Bonus: Round off the result to 1 decimal. *Tip: Use the **Maths Functions** section of your cheat sheet!*

# Exercise 1

---

1. Do the following calculation in R:

$$\frac{1 + 5}{9}$$

```
> (1+5)/9
```

2. Assign the result of the calculation to a variable.

```
> MyCalc <- (1+5)/9
```

3. Bonus: Round off the result to 1 decimal. *Tip: Use the **Maths Functions** section of your cheat sheet!*

```
> round(MyCalc, 1)
```

# Another data type: logical

---

T	TRUE
F	FALSE

```
> TRUE
```

```
[1] TRUE
```

```
> F
```

```
[1] FALSE
```

```
> x==6
```

```
[1] TRUE
```

```
> 2>4
```

```
[1] FALSE
```

`==` is equal to

`!=` is not

`>=` larger than or equal to

`<` smaller than

# Combining data: creating vectors

```
> c(1,2,3) [1] 1 2 3
```

a numeric vector

```
> c("a", "b", "c")  
[1] "a"  "b"  "c"  ← a character vector
```

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

a logical vector

```
> c(TRUE, "a", 3)          ??????  
[1] TRUE    "a"      "3"      ←
```

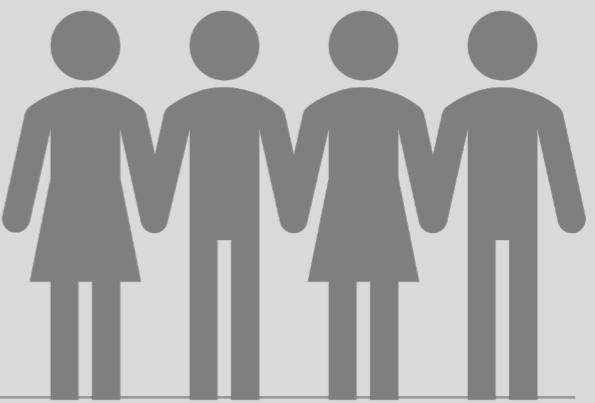
# Vector functions

---

```
> p <- 1:5  
> p  
[1] 1 2 3 4 5  
  
> mean(p)  
[1] 3  
  
> p * 2  
[1] 2 4 6 8 10  
  
> q <- 5:1  
> q  
[1] 5 4 3 2 1  
  
> p * q  
[1] 5 8 9 8 5
```

p * 2		
p	2	
1	2	2
2	2	4
3	2	6
4	2	8
5	2	10

p * q		
p	q	
1	5	5
2	4	8
3	3	9
4	2	8
5	1	5

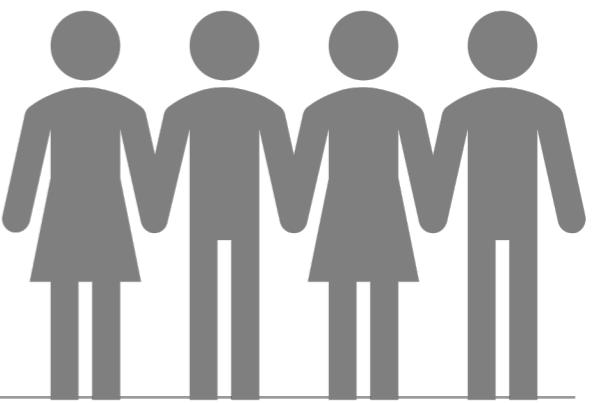


## Exercise 2: create vectors

---

Meet Ann, Bob, Chloe, and Dan.

1. Make a character vector with their names, using the function `c()`. Save the vector as `name`.
2. How old are Ann, Bob, Chloe, and Dan? Design a numeric vector with their respective ages. Save it as `age`.
3. Bonus: What is their average age? Use a function in R to calculate this.



## Exercise 2: create vectors

---

Meet Ann, Bob, Chloe, and Dan.

1. Make a character vector with their names, using the function `c()`. Save the vector as `name`.

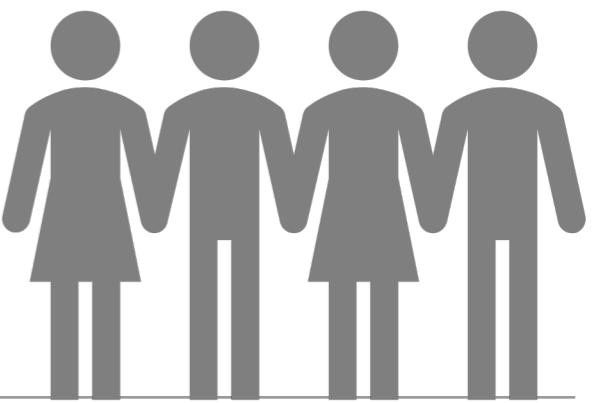
```
> name <- c("Ann", "Bob", "Chloe", "Dan")
```

2. How old are Ann, Bob, Chloe, and Dan? Design a numeric vector with their respective ages. Save it as `age`.

```
> age <- c(35, 22, 50, 51)
```

3. Bonus: What is their average age? Use a function in R to calculate this.

```
> mean(age)  
[1] 39.5
```



# Data structures

```
> name  
[1] "Ann"     "Bob"      "Chloe"   "Dan"  
> age  
[1] 35 22 50 51  
  
> c(name,age)  
[1] "Ann"     "Bob"      "Chloe"   "Dan"      "35"       "22"       "50"       "51"  
  
> list(name,age)  
[[1]]  
[1] "Ann"     "Bob"      "Chloe"   "Dan"  
  
[[2]]  
[1] 35 22 50 51  
  
> data.frame(name,age)  
  name  age  
1  Ann   35  
2  Bob   22  
3 Chloe  50  
4  Dan   51
```

# Data structures

---

	how many dimensions?	function
vector	1	<code>c()</code>
list	any number	<code>list()</code>
data frame	2	<code>data.frame()</code>



# Vectors and factors

---

```
> country <- c("UK", "USA", "USA", "UK")
```

```
> country
```

```
[1] "UK"   "USA"  "USA"  "UK"
```

```
> as.factor(country)
```

```
[1] UK   USA  USA  UK
```

```
Levels: UK USA
```

A factor is defined by its **levels**  
(most useful for a category)

## Exercise 3: combining data

---

1. Create a vector **country** containing four countries (use at least one duplicate!).
2. Create a data frame combining **name**, **age**, and **country**, and save it as **df**.
3. Bonus: create a list combining **name**, **age**, and **country**.

## Exercise: combining data

---

1. Create a vector **country** containing four countries (use at least one duplicate!).

```
> country <- c("UK", "USA", "USA", "UK")
```

2. Create a data frame combining **name**, **age**, and **country**, and save it as **df**.

```
> df <- data.frame(name, age, country)
```

3. Bonus: create a list combining **name**, **age**, and **country**.

```
> myList <- list(name, age, country)
```

# Questions: data types inside a data frame

---

What happened to the character vectors when you put them in a data frame?

```
> df$name  
[1] Ann     Bob      Chloe   Dan  
Levels: Ann Bob Chloe Dan
```

```
> df$country  
[1] UK     USA    USA    UK  
Levels: UK USA
```

Is this useful? Where? Where not?

Did the same happen when you put the vectors in a list? *Hint: you can select vector elements with double brackets: myList[[1]]*

```
> myList[[1]]  
[1] "Ann"    "Bob"    "Chloe"  "Dan"
```

# adding info to an existing data frame

Data without data: NA

```
> df$pet <- c("cat", "none", "fish", NA)
```

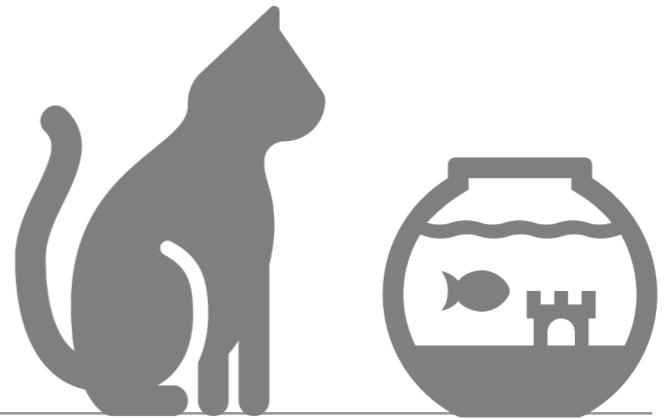
```
> df
```

	name	age	country	pet
1	Ann	35	UK	cat
2	Bob	22	USA	none
3	Chloe	50	USA	fish
4	Dan	51	UK	<NA>

```
> as.factor(df$pet)  
[1] cat none fish <NA>
```

Levels: cat fish none

NA = Not Available



name	age	country	pet
Ann	35	UK	cat
Bob	22	USA	none
Chloe	50	USA	fish
Dan	51	UK	NA

In other words:

We **know** that Bob has **no pets**.

We **do not know** if Dan has pets.

## Exercise 4: predict the answer

---

Predict the results. Does the (real) answer make sense to you?

```
> 5 == 5
```

```
> 5 == NA
```

```
> NA == NA
```

## Exercise 4: predict the answer

---

Predict the results. Does the (real) answer make sense to you?

```
> 5 == 5
```

```
[1] TRUE
```

```
> 5 == NA
```

```
[1] NA
```

```
> NA == NA
```

```
[1] NA
```

```
> is.na(NA)
```

```
[1] TRUE
```

# Selecting vector elements by position

> name

```
[1] "Ann"    "Bob"    "Chloe" "Dan"
```

> name[2]

```
[1] "Bob"
```

> name[1:3]

```
[1] "Ann"    "Bob"    "Chloe"
```

## Base R Cheat Sheet

### Getting Help

#### Accessing the help files

?mean  
Get help of a particular function.  
help.search('weighted mean')  
Search the help files for a word or phrase.  
help(package = 'dplyr')  
Find help for a package.

#### More about an object

str(iris)  
Get a summary of an object's structure.  
class(iris)  
Find the class an object belongs to.

### Using Packages

install.packages('dplyr')  
Download and install a package from CRAN.

library(dplyr)  
Load the package into the session, making all its functions available to use.

dplyr::select  
Use a particular function from a package.

data(iris)  
Load a built-in dataset into the environment.

### Working Directory

getwd()  
Find the current working directory (where inputs are found and outputs are sent).

setwd('C://file/path')  
Change the current working directory.

Use projects in RStudio to set the working directory to the folder you are working in.

Vectors		
Creating Vectors		
c(2, 4, 6)	2 4 6	Join elements into a vector
2:6	2 3 4 5 6	An integer sequence
seq(2, 3, by=0.5)	2.0 2.5 3.0	A complex sequence
rep(1:2, times=3)	1 2 1 2 1 2	Repeat a vector
rep(1:2, each=3)	1 1 1 2 2 2	Repeat elements of a vector

Vector Functions		
sort(x)	rev(x)	Return x sorted. Return x reversed.
table(x)	unique(x)	See counts of values. See unique values.

Selecting Vector Elements		
By Position		
x[4]	The fourth element.	
x[-4]	All but the fourth.	
x[2:4]	Elements two to four.	
x[-(2:4)]	All elements except two to four.	
x[c(1, 5)]	Elements one and five.	
By Value		
x[x == 10]	Elements which are equal to 10.	
x[x < 0]	All elements less than zero.	
x[x %in% c(1, 2, 5)]	Elements in the set 1, 2, 5.	
Named Vectors		
x['apple']	Element with name 'apple'.	

Conditions	a == b	Are equal	a > b	Greater than	a >= b	Greater than or equal to	is.na(a)	Is missing
	a != b	Not equal	a < b	Less than	a <= b	Less than or equal to	is.null(a)	Is null

Programming				
For Loop				
for (variable in sequence){ Do something }				
Example				
for (i in 1:4){ j <- i + 10 print(j) }				
While Loop				
while (condition){ Do something }				
Example				
while (i < 5){ print(i) i <- i + 1 }				
Functions				
function_name <- function(var){ Do something return(new_variable)				
Example				
square <- function(x){ squared <- x*x return(squared)				
Reading and Writing Data				
Also see the <a href="#">readr</a> package.				
Input				
df <- read.table('file.txt')	write.table(df, 'file.txt')	Read and write a delimited text file.		
Output				
df <- read.csv('file.csv')	write.csv(df, 'file.csv')	Read and write a comma separated value file. This is a special case of read.table/write.table.		
Description				
load('file.RData')	save(df, file = 'file.Rdata')	Read and write an R data file, a file type special for R.		
Types				
Converting between common data types in R. Can always go from a higher value in the table to a lower value.				
as.logical	TRUE, FALSE, TRUE	Boolean values (TRUE or FALSE).		
as.numeric	1, 0, 1	Integers or floating point numbers.		
as.character	'1', '0', '1'	Character strings. Generally preferred to factors.		
as.factor	'1', '0', '1', levels: '1', '0'	Character strings with preset levels. Needed for some statistical models.		
Matrices				
m <- matrix(x, nrow = 3, ncol = 3) Create a matrix from x.				
m[2, ]	Select a row	t(m)		
m[, 1]	Select a column	Transpose m %*% n		
m[2, 3]	Select an element	Matrix Multiplication solve(m, n) Find x in: m * x = n		
Lists				
l <- list(x = 1:5, y = c('a', 'b')) A list is a collection of elements which can be of different types.				
l[2]	Second element of l.	l[1]		
	New list with only the first element.	l\$x		
	Element named x.	l['y']		
	New list with only element named y.			
Data Frames				
df <- data.frame(x = 1:3, y = c('a', 'b', 'c')) A special case of a list where all elements are the same length.				
x   y				
df\$x	df[2]			
Variables Assignment				
> a <- 'apple' > a [1] 'apple'				
The Environment				
ls() List all variables in the environment.				
rm(x)	Remove x from the environment.			
rm(list = ls())	Remove all variables from the environment.			
You can use the environment panel in RStudio to browse variables in your environment.				
Strings				
Also see the <a href="#">stringr</a> package.				
paste(x, y, sep = ' ')	Paste multiple vectors together.			
paste(x, collapse = ' ')	Join elements of a vector together.			
grep(pattern, x)	Find regular expression matches in x.			
gsub(pattern, replace, x)	Replace matches in x with a string.			
toupper(x)	Convert to uppercase.			
tolower(x)	Convert to lowercase.			
nchar(x)	Number of characters in a string.			
Factors				
factor(x)				
Turn a vector into a factor. Can set the levels of the factor and the order.				
Statistics				
Also see the <a href="#">stats</a> package.				
lm(y ~ x, data=df)	Linear model.			
glm(y ~ x, data=df)	Generalised linear model.			
summary	Get more detailed information out a model.			
pairwise.t.test	Perform a t-test for paired data.			
avov	Analysis of variance.			
Distributions				
Random Variates				
Normal	rnorm	dnorm	pnorm	qnorm
Poisson	rpois	dpois	ppois	qpois
Binomial	rbinom	dbinom	pbinom	qbinom
Uniform	runif	dunif	punif	qunif
Plotting				
Also see the <a href="#">ggplot2</a> package.				
plot(x, y)	Values of x in order.	plot(x, y)		
hist(x)	Values of x against y.	hist(x)		
Dates				
See the <a href="#">lubridate</a> package.				

### Selecting Vector Elements

#### By Position

**x[4]** The fourth element.

**x[-4]** All but the fourth.

**x[2:4]** Elements two to four.

**x[-(2:4)]** All elements except two to four.

**x[c(1, 5)]** Elements one and five.

# Selecting vector elements by value

```
> age  
[1] 35 22 50 51
```

```
> age[age>40]
```

```
[1] 50 51
```

```
> age>40
```

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

```
> name[name %in% c("Chloe", "Ann", "Evie")]
```

```
[1] "Ann" "Chloe"
```

age	age>40	result
35	FALSE	
22	FALSE	
50	TRUE	50
51	TRUE	51

## Selecting Vector Elements

### By Value

**x[x == 10]** Elements which are equal to 10.

**x[x < 0]** All elements less than zero.

**x[x %in% c(1, 2, 5)]** Elements in the set 1, 2, 5.

### Named Vectors

**x['apple']** Element with name 'apple'.

## Exercise 5: selecting vector elements

---

1. Return only the first number in your vector `age`.
2. Return the 2<sup>nd</sup> and 4<sup>th</sup> name in your vector `name`.
3. Return only ages under 30 from your vector `age`.

## Exercise 5: selecting vector elements

---

1. Return only the first number in your vector `age`.

```
> age[1]  
[1] 35
```

2. Return the 2<sup>nd</sup> and 4<sup>th</sup> name in your vector `name`.

```
> name[c(2,4)]  
[1] "Bob" "Dan"
```

3. Return only ages under 30 from your vector `age`.

```
> age[age<30]  
[1] 22
```

# Indexing lists

---

```
> myList <- list(name,age)
```

```
> myList  
[[1]]  
[1] "Ann"     "Bob"      "Chloe"   "Dan"
```

```
[[2]]  
[1] 35 22 50 51
```

```
> myList[1] ← select the first list element
```

```
[[1]]  
[1] "Ann"     "Bob"      "Chloe"   "Dan"
```

```
> myList[[1]] ← select the content of the first list element  
[1] "Ann"     "Bob"      "Chloe"   "Dan"
```

## Indexing lists (continued)

---

```
> myList[[1]][2]
```

```
[1] "Bob"
```

```
> myList[1][2]
```

```
[[1]]
```

```
NULL
```

**NA != NULL**

---

**NA** Information is **Not Available**

**NULL** Information **does not exist**

**“None” or 0** Data entry specifying **content of 0**

## Exercise 6: predict the answer

---

```
> is.na(NA)
```

```
[1] TRUE
```

```
> is.null(NULL)
```

```
[1] TRUE
```

Predict the results. Does the (real) answer make sense to you?

```
> is.null(NA)
```

```
> is.na(NULL)
```

## Exercise: predict the answer

---

```
> is.na(NA)
```

```
[1] TRUE
```

```
> is.null(NULL)
```

```
[1] TRUE
```

Predict the results. Does the (real) answer make sense to you?

```
> is.null(NA)
```

```
[1] FALSE
```

```
> is.na(NULL)
```

```
[1] logical(0)
```

# Indexing a data frame

---

## Matrix subsetting

`df[ , 2]`



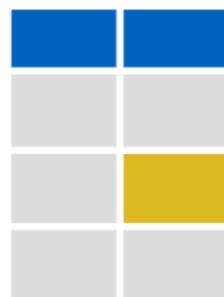
`df[2, ]`



row

column

`df[2, 2]`



name	age	country
Ann	35	UK
Bob	22	USA
Chloe	50	USA
Dan	51	UK

# Indexing a data frame

```
> df[,2]  
[1] 35 22 50 51  
  
> df[, "age"]  
[1] 35 22 50 51  
  
> df$age  
[1] 35 22 50 51  
  
> df[2,]  
  name age country  
2 Bob 22 USA  
  
> df[df$name=="Bob",]  
  name age country  
2 Bob 22 USA  
  
> df[df$name=="Bob", "age"]  
[1] 22
```

**columns**  
df[,2]  
df[, "age"]  
df\$age



name	age	country
Ann	35	UK
Bob	22	USA
Chloe	50	USA
Dan	51	UK



**rows**  
df[2,]  
df[df\$name=="Bob", ]

# Remove variables

---

Before we start with the exercise, please **remove the vectors** that were the basis of your data frame from your environment, like this:

```
> rm(name,age,country)
```

*(We do this so they cannot confuse you during the following exercise!)*

## Exercise 7: selecting from dataframe

---

1. From your dataframe `df`, return the entries for everyone living in a country of your choice.
2. Return **only the names** of everyone in your data frame `df` under 40.  
*(Hint: what information should you use for row indexing?  
What information should you use for column indexing?)*
3. Bonus: can you use vector indexing on a column to achieve the same result?

## Exercise 7: selecting from dataframe

---

1. From your dataframe df, return the entries for everyone living in a country of your choice.

```
> df[df$country=="USA", ]
```

	name	age	country
2	Bob	22	USA
3	Chloe	50	USA

2. Return only the names of everyone in your data frame df under 40.

(*Hint: what information should you use for row indexing?*

*What information should you use for column indexing?*)

```
> df[df$age<40, "name"]
```

```
> df[df$age<40, 1]
```



indexing the dataframe df

```
[1] Ann Bob
```

```
Levels: Ann Bob Chloe Dan
```

3. Bonus: can you use vector indexing on a column to achieve the same result?

```
> df$name[df$age<40]
```



indexing the vector df\$name

# Which bracket does what?

---

- [ ] **Indexing** vectors, lists, dataframes...
- ( ) Passing **arguments** to functions
- { } **Defining content** of loops, functions, etc.

# Let's breathe and recap!

---

What data types have you encountered so far?

`logical`

`numeric`

`character`

How can data be absent?

`NA` (not available)

`NULL` (non-existent)

And what data structures have you encountered?

`vector` (one dimension)

`factor` (one dimension, level-based)

`data frame` (two dimensions)

`list` (++ dimensions)

# Functions

---

What functions have you encountered so far?

```
> c()  
> as.factor()  
> data.frame()  
> is.na()  
> mean()  
...
```

And do you still know what they mean? And how to use them? **No?**

```
> ?mean()  
> help.search("standard deviation")
```

(or use the Help window to the right of your console)

# Help!

table {base} ← function & package names

R Documentation

## Cross Tabulation and Table Creation

### Description

table uses the cross-classifying factors to build a contingency table of the counts at each combination of factor levels.

### Usage

table(..., ← how would you use the function?  
  exclude = if (useNA == "no") c(NA, NaN),  
  useNA = c("no", "ifany", "always"),  
  dnn = list.names(...), deparse.level = 1)}

wait what? These are extra arguments. If you see argument = "default" then the setting is already specified, so you won't have to.

as.table(x, ...) ← functions so related they share a help page  
is.table(x)

```
## S3 method for class 'table'  
as.data.frame(x, row.names = NULL, ...)
```

# Help? Scroll down!

---

## Examples

```
require(stats) # for rpois and xtabs
## Simple frequency distribution
table(rpois(100, 5))
## Check the design:
with(warpbreaks, table(wool, tension))
table(state.division, state.region)

# simple two-way contingency table
with(airquality, table(cut(Temp, quantile(Temp)), Month))

a <- letters[1:3]
table(a, sample(a))                      # dnn is c("a", "")
table(a, sample(a), deparse.level = 0) # dnn is c("", "")
table(a, sample(a), deparse.level = 2) # dnn is c("a", "sample(a)")

## xtabs() <-> as.data.frame.table() :
UCBAdmissions ## already a contingency table
DF <- as.data.frame(UCBAdmissions)
class(tab <- xtabs(Freq ~ ., DF)) # xtabs & table
## tab *is* "the same" as the original table:
all(tab == UCBAdmissions)
```

# Take a break!

---



# Programming

---

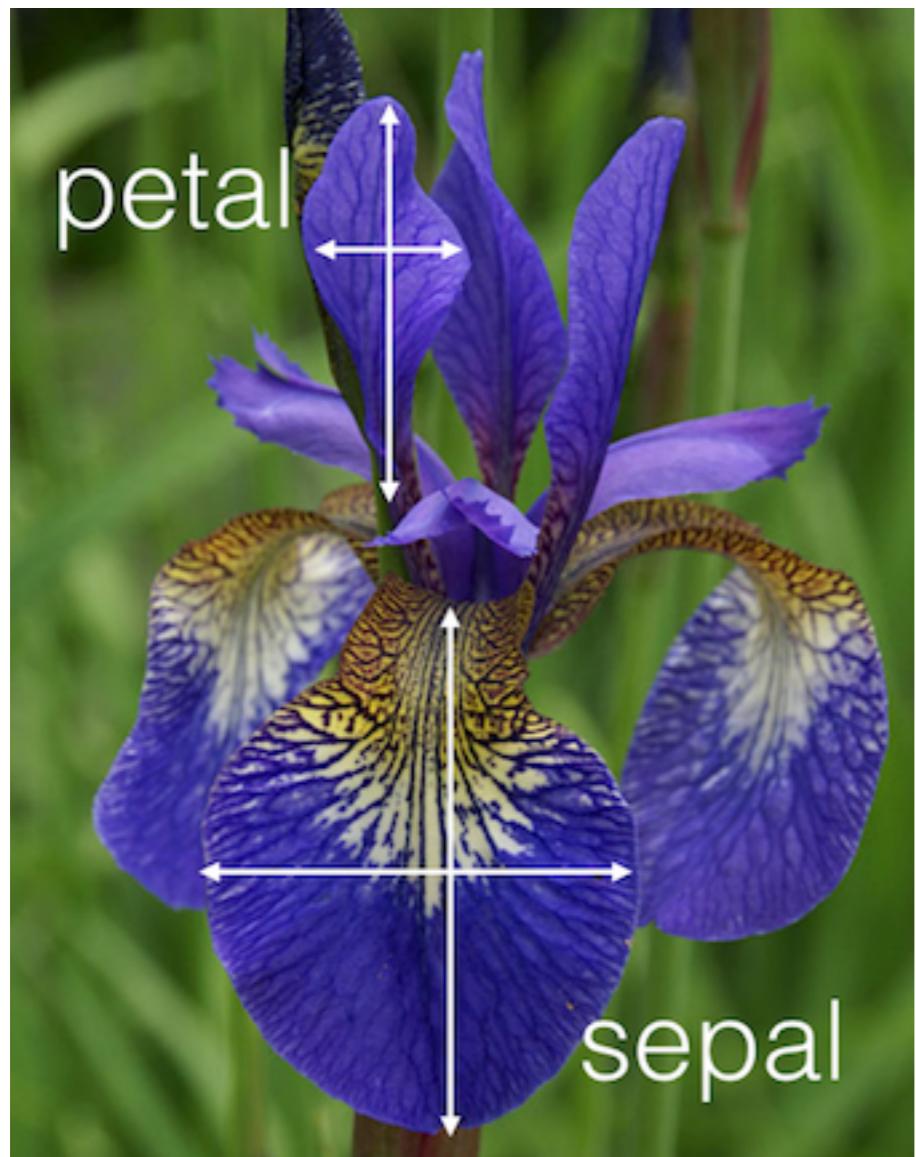
- Writing a multiline script
- Programming with loops and functions
- Handling and processing a dataset

```
lines(density(glnorm[,sname],na.rm=TRUE))  
move the samples from the data  
* {r}  
select the samples to keep  
keepsamples <- row.names(pheno)  
apply sample selection to counts  
counts.sub <- counts.sub[,keepsamples]  
pheno.sub <- pheno[keepsamples]  
mt.assay <- mt.assay[,keepsamples]  
rlid.sub <- rlid[,keepsamples]
```

# Introducing a sample dataset

---

- Dataset: ‘iris’
- Standard dataset in R, measurements on 3 species of iris flowers



# Introducing a sample dataset

---

- Dataset: ‘iris’
- Standard dataset in R, measurements on 3 species of iris flowers

```
> head(iris)
```

	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
1	5.1	3.5	1.4	0.2	setosa
2	4.9	3.0	1.4	0.2	setosa
3	4.7	3.2	1.3	0.2	setosa
4	4.6	3.1	1.5	0.2	setosa
5	5.0	3.6	1.4	0.2	setosa
6	5.4	3.9	1.7	0.4	setosa

```
> summary(iris)
```

Sepal.Length	Sepal.Width	Petal.Length	Petal.Width	Species
Min. :4.300	Min. :2.000	Min. :1.000	Min. :0.100	setosa :50
1st Qu.:5.100	1st Qu.:2.800	1st Qu.:1.600	1st Qu.:0.300	versicolor:50
Median :5.800	Median :3.000	Median :4.350	Median :1.300	virginica :50
Mean :5.843	Mean :3.057	Mean :3.758	Mean :1.199	
3rd Qu.:6.400	3rd Qu.:3.300	3rd Qu.:5.100	3rd Qu.:1.800	
Max. :7.900	Max. :4.400	Max. :6.900	Max. :2.500	

## Exercise 8: iris

---

Explore the `iris` dataframe, using some of the following functions:

`head()`

`tail()`

`names()`

`summary()`

`dim()`

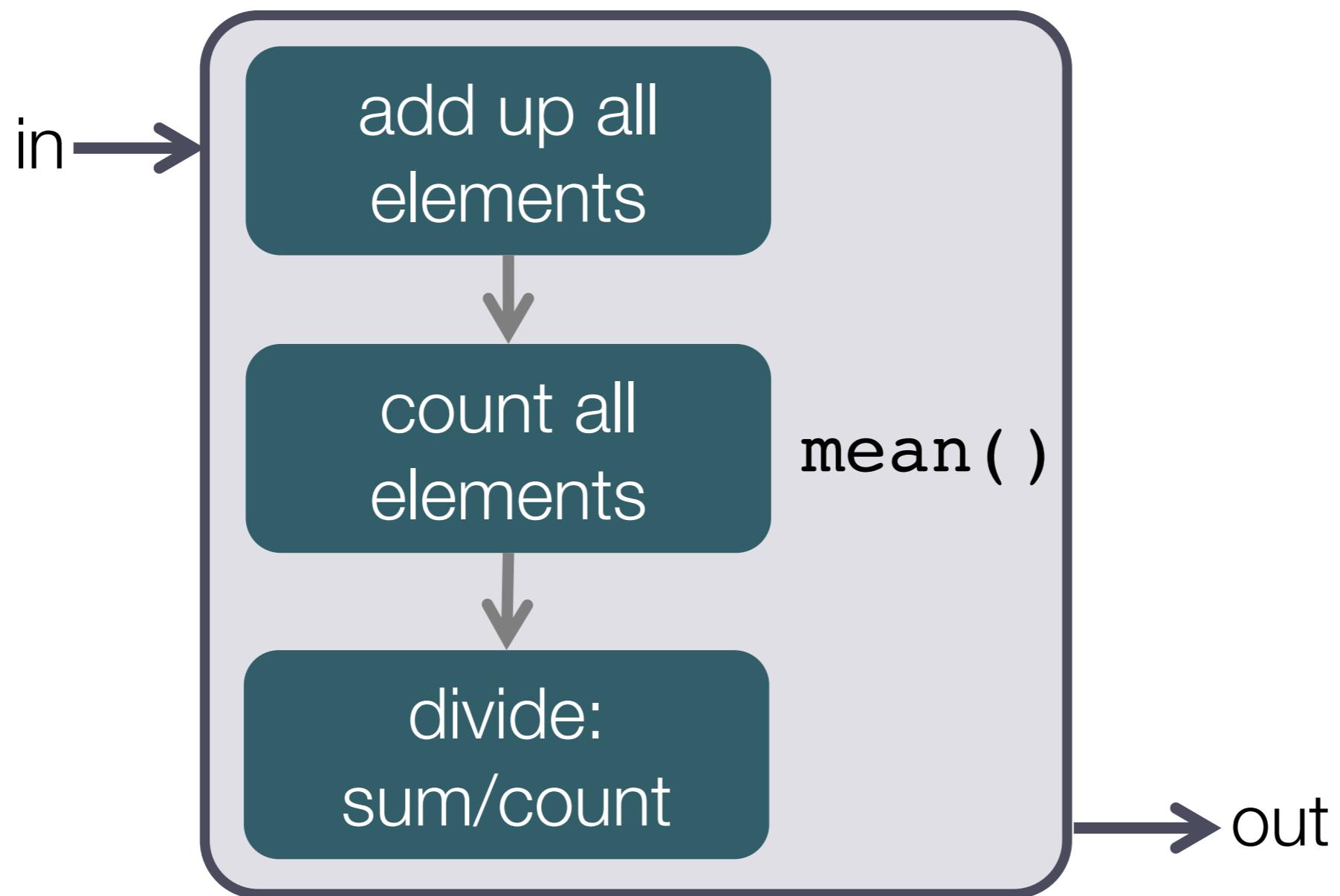
`str()`

Can you figure out what these functions do? What do they teach you about the kind of data in `iris`?

# Programming: functions

---

- Multiple instructions that form a cohesive unit
- Should be able to be repeated on different inputs



# Programming: functions

---

```
myFun <- function(x,y){  
  z <- x*y  
  return(z)  
}
```

```
> myFun(2,4)  
[1] 8
```



Step 1: run the function. This saves the function to memory under the *name* you assigned it.

Step 2: use the function. You can now execute the function by calling its *name*.

## Exercise 9: function

---

1. Write a function that takes a vector as input, and returns the mean of this vector (you can use the existing function `mean()` inside your function).

```
apply_calc <- function(...){  
  ...  
  return(...)  
}
```

2. Add further options to your function:
  - a. for example, the standard deviation (`sd()`), the minimum (`min()`), and the maximum (`max()`) of your input vector.
  - b. Put all of these calculations in a vector using the function `c()`, and return this result vector.

```
apply_calc <- function(...){  
  ...  
  
  allres <- c(...)  
  return(...)  
}
```

## Exercise 9: function

---

1. Write a function that takes a vector as input, and returns the mean of this vector (you can use the existing function `mean()` inside your function).

```
apply_calc <- function(x) {  
  m <- mean(x)  
  return(m)  
}
```

2. Add further options to your function:
  - a. for example, the standard deviation (`sd()`), the minimum (`min()`), and the maximum (`max()`) of your input vector.
  - b. Put all of these calculations in a vector using the function `c()`, and return this result vector.

```
apply_calc <- function(x) {  
  m <- mean(x)  
  s <- sd(x)  
  mi <- min(x)  
  ma <- max(x)  
  allres <- c(m,s,mi,ma)  
  return(allres)  
}
```

# Applying apply\_calc

---

```
apply_calc <- function(x){  
  m <- mean(x)  
  return(m)  
}
```

```
> apply_calc(iris$Sepal.Length)  
[1] 5.8433333
```

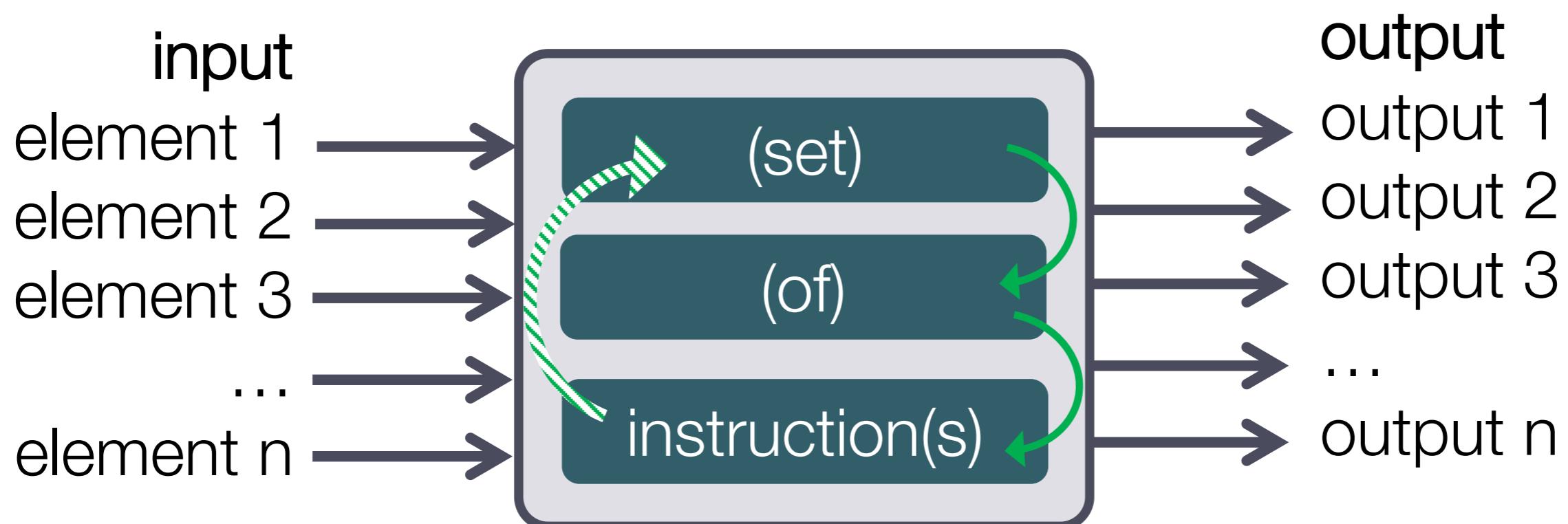
```
apply_calc <- function(x){  
  m <- mean(x)  
  s <- sd(x)  
  mi <- min(x)  
  ma <- max(x)  
  allres <- c(m,s,mi,ma)  
  return(allres)  
}
```

```
> apply_calc(iris$Sepal.Length)  
[1] 5.8433333 0.8280661 4.3000000 7.9000000
```

# Programming: for-loops

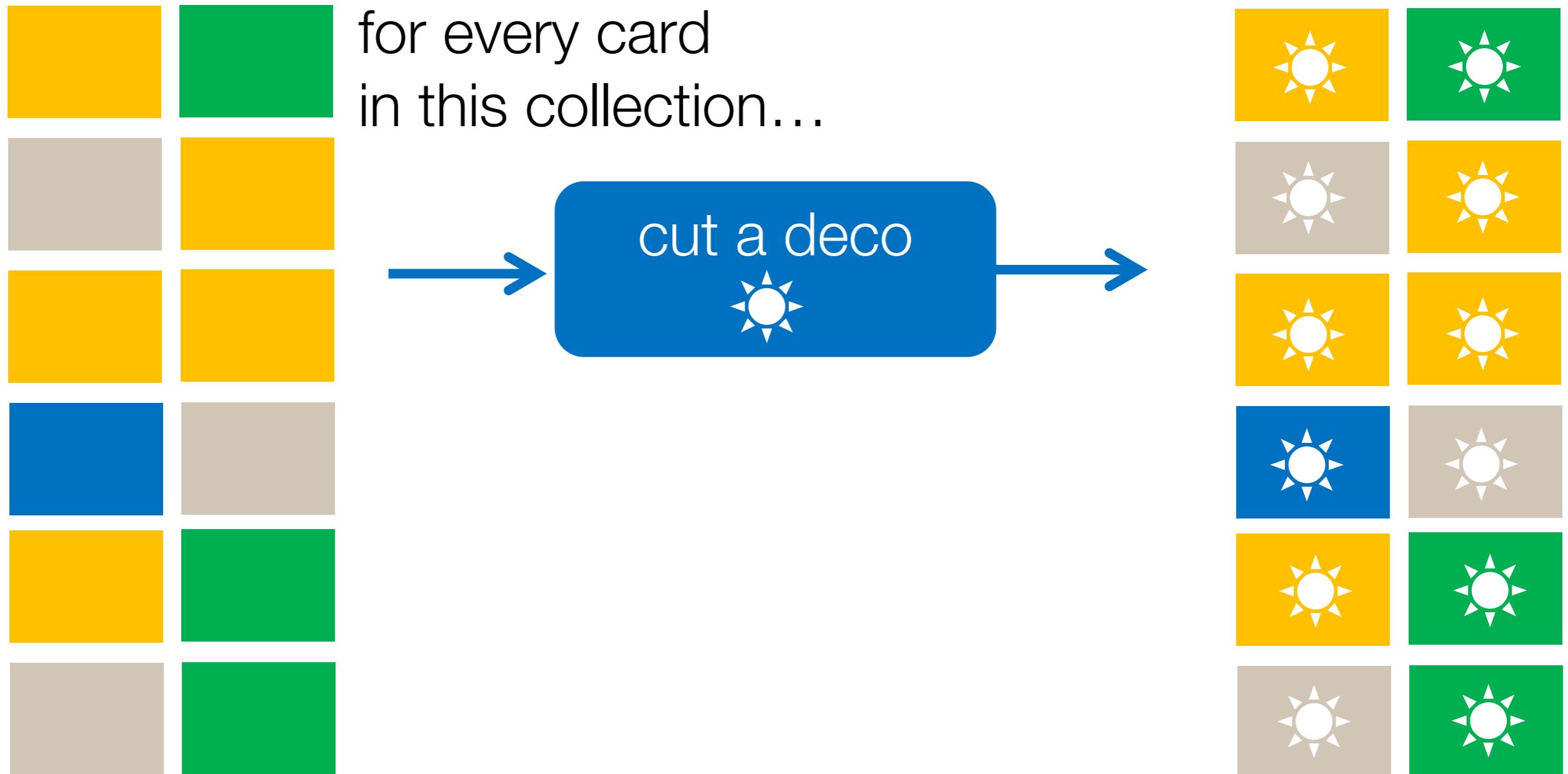
---

- Instruction(s) that need(s) to be applied multiple times
- Input is an iterable object (it consists of multiple similar elements)



# Imagine: the repeated action in a for-loop

---



# How would this look in R?

---

```
for(■ in [■■■■■]) {  
  cut_deco(■)  
}
```

```
[1] ☀  
[1] ☀  
[1] ☀  
[1] ☀  
[1] ☀  
[1] ☀
```

```
for(i in 1:6){  
  print(i)  
}
```

```
[1] 1  
[1] 2  
[1] 3  
[1] 4  
[1] 5  
[1] 6
```

# Exercise 10: for loop

---

1. Make a vector with all the column names in `iris`.

*Hint: use the function `colnames()`.*

```
iriscols <- ...
```

2. Make a for-loop that iterates over all the column names in `iris`, and prints these column names.

```
for(...){  
  ...  
}
```

3. Elaborate on this for-loop: select the corresponding column in `iris`, and print the mean. *Hint: yes, you should get a warning! Do you understand why?*

```
for(...){  
  ...  
}
```

# Exercise

---

1. Make a vector with all the column names in `iris`.

*Hint: use the function `colnames()`.*

```
iriscols <- colnames(iris)
```

2. Make a for-loop that iterates over all the column names in `iris`, and prints these column names.

```
for(i in iriscols){  
  print(i)  
}
```

3. Elaborate on this for-loop: select the corresponding column in `iris`, and print the mean. *Hint: yes, you should get a warning! Do you understand why?*

```
for(i in iriscols){  
  column <- iris[,i]  
  stat <- mean(column)  
  print(stat)  
}
```

# What's wrong? And how do we fix it?

---

```
for(i in iriscols){  
  # select the appropriate column  
  column <- iris[,i]  
  # calculate the mean  
  stats <- mean(column)  
  # print the mean  
  print(stats)  
}
```

```
[1] 5.843333  
[1] 3.057333  
[1] 3.758  
[1] 1.199333  
[1] NA
```

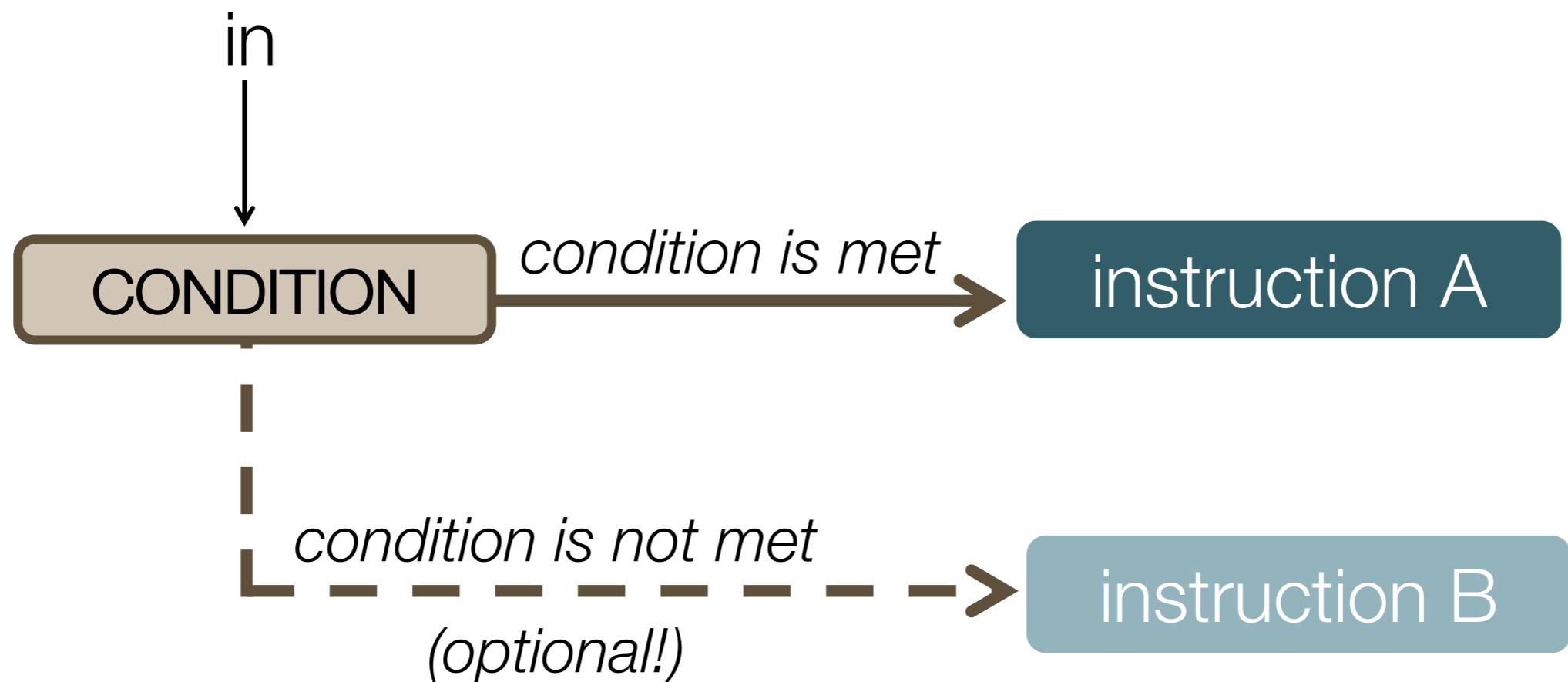
Warning message:

In mean.default(c) : argument is not numeric or logical:  
returning NA

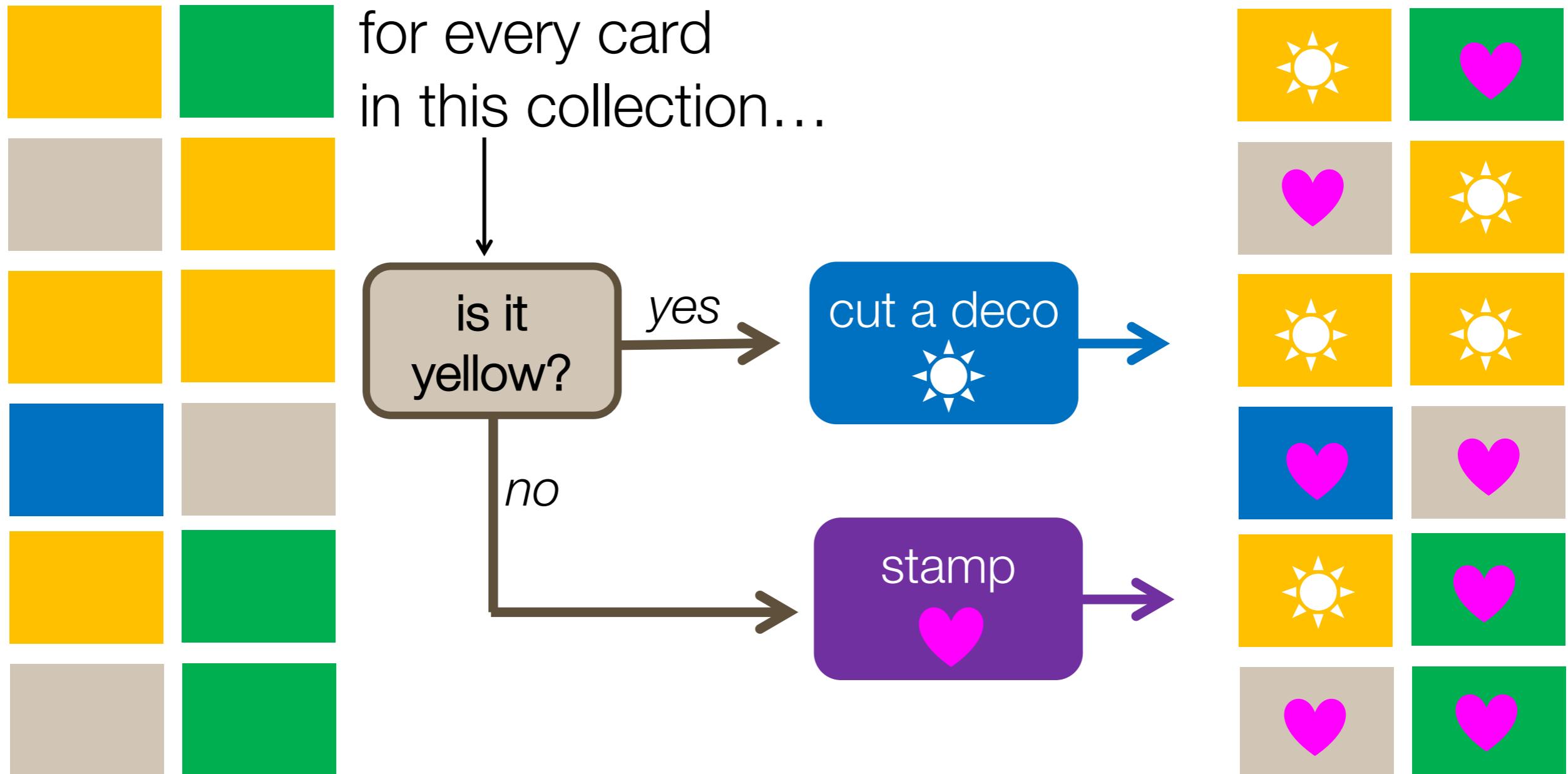
# Programming: if-statement

---

- Test to conditionally apply an instruction
- Possibility for an alternative instruction (if the test is negative)



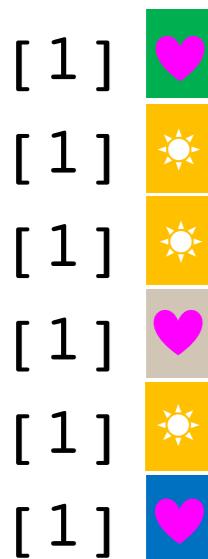
# Imagine: the selectivity of an if-statement



PS: note how this is still a for-loop? That's not a requirement!

# How would this look in R?

```
for(■ in [■■■■■]) {  
  if(■ == ■){  
    cut_deco(■)  
  } else{  
    stamp(■)  
  }  
}
```



```
for(i in 1:6){  
  if(i > 3){  
    print("Large!")  
  } else{  
    print("small...")  
  }  
}
```

```
[1] "small..."  
[1] "small..."  
[1] "small..."  
[1] "Large!"  
[1] "Large!"  
[1] "Large!"
```

# Programming: if-statement

---

```
d <- 7

if(d >= 10){
  print("d is 10 or larger")
} else if(d > 5){
  print("d is between 5 and 10")
} else{
  print("d is 5 or lower")
}
```

```
[1] "d is between 5 and 10"
```

## Exercise 11: if statement

---

Copy-paste the for-loop you made in the previous exercise. Inside this for-loop, add an if-statement, so that `mean()` is only performed on numeric vectors. *Hint: check the function `is.numeric()`.*

```
for(i in iriscols){  
  column <- iris[,i]  
  ...  
}
```

## Exercise 11: if statement

---

Copy-paste the for-loop you made in the previous exercise. Inside this for-loop, add an if-statement, so that `mean()` is only performed on numeric vectors. *Hint: check the function `is.numeric()`.*

```
for(i in iriscols){  
  column <- iris[,i]  
  if(is.numeric(column)){  
    stat <- mean(column)  
    print(stat)  
  }  
}
```

# To check before lunch: please load tidyverse

---

```
> library(tidyverse)
— Attaching packages ————— tidyverse 1.2.1 —
  ggplot2 3.1.0      purrr   0.3.0
  tibble   2.0.1      dplyr    0.7.8
  tidyrr   0.8.2      stringr 1.4.0
  readr    1.3.1     forcats  0.3.0
— Conflicts ————— tidyverse_conflicts() —
  dplyr::filter() masks stats::filter()
  dplyr::lag()    masks stats::lag()
```

Enjoy your lunch!

---

