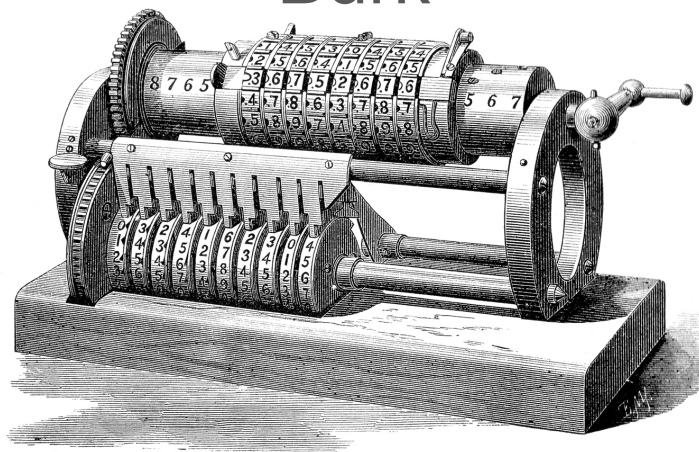
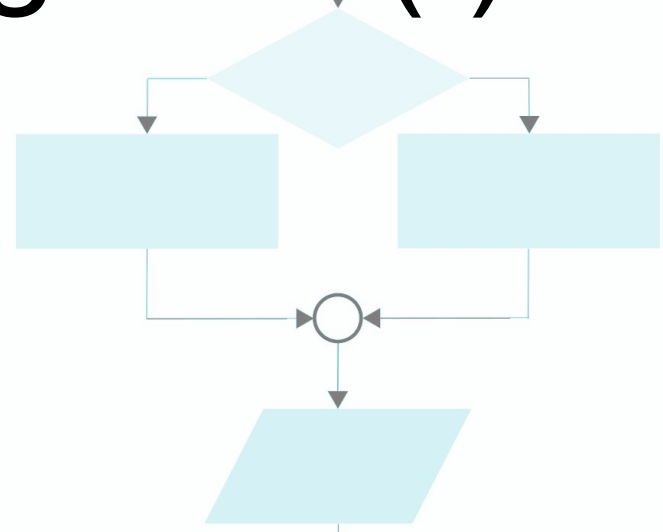


# Programming in R

Binder, Bender,  
Burk



# Unit 2: Structured Programming in R (I)



# Structured Programming in R

Remember, this course has multiple goals:

- Learn things about the R language: "R"
- Get to know nice tools to use: "Tools"
- Learn things about software development in general: "Dev"

This unit:

- "R" Track: Data Types and Control Structures
- "Dev" Track: Programming Style

R Track

Data Types and Control Structures

# "Structured Programming"

- Using conditionals ("if..then"), loops ("for"), and functions to solve a problem
- ... as opposed to, say, copy-pasting your code to execute the same action multiple times<sup>1</sup>
  - people actually do that and it makes for entertaining stories. Dont. Do. It.
- This course unit is about basic structured programming in R and covers a large part of the content of last year's course.
  - Datatypes (atomics, lists, matrices, data.frame)
  - Operators
  - Control structures (conditionals, loops, functions)
- We will go a bit beyond last year to get you a solid foundational knowledge of the "nuts and bolts" in R!

<sup>1</sup>In some situations there is a legitimate technique called "loop unrolling" for performance reasons. Don't do it in R; if you need performance in R you use Rcpp or similar.

# Reminder: Data Types

- "atomic" data types: `logical`, `numeric`, `character`
  - (also less importantly: `integer`, `complex`, `raw`)
  - "vectors"
- "recursive" data types: `list` (and some other more special things)

# Reminder: Data Types

- **atomic** data types: logical, numeric, character
  - (also less importantly: integer, complex, raw)
  - "vectors"
- "atomic": always a vector of zero or more values *of the same type*
- no nesting, no recursive structure

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

```
> c(1, "a")
[1] "1" "a"
```

- **recursive** data types: list
  - (and some other things we omit for now)
- may contain any other values

```
> list(1, "a", list())
[[1]]
[1] 1

[[2]]
[1] "a"

[[3]]
list()
```

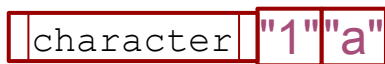
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[1] 1 3
```



```
> c(1, "a")
[1] "1" "a"
```

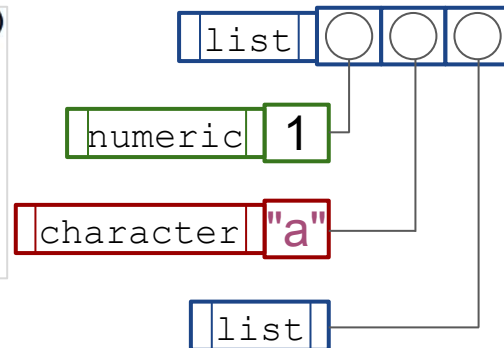


- **recursive** data types: list
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- may contain any other values

```
> list(1, "a", list())
[[1]]
[1] 1

[[2]]
[1] "a"

[[3]]
list()
```





# Reminder: Data Types

- **atomic** data types: logical, numeric, character
  - (also less importantly: integer, complex, raw)
  - "vectors"
- "atomic": always a vector of zero or more values *of the same type*
- no nesting, no recursive structure

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

numeric 1 3

```
> c(1, "a")
[1] "1" "a"
```

character "1" "a"

- **recursive** data types: list
  - (and some other things we omit for now)
- may contain any other values

```
> list(c(1, 3), c(1, "a"), list(2, "b"))
[[1]]
[1] 1 3

[[2]]
[1] "1" "a"

[[3]]
[[3]][[1]]
[1] 2

[[3]][[2]]
[1] "b"
```

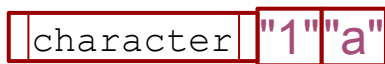
# Reminder: Data Types

- **atomic** data types: logical, numeric, character
  - (also less importantly: integer, complex, raw)
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- "atomic": always a vector of zero or more values *of the same type*
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```
> c(1, 3)
[1] 1 3
```



```
> c(1, "a")
[1] "1" "a"
```



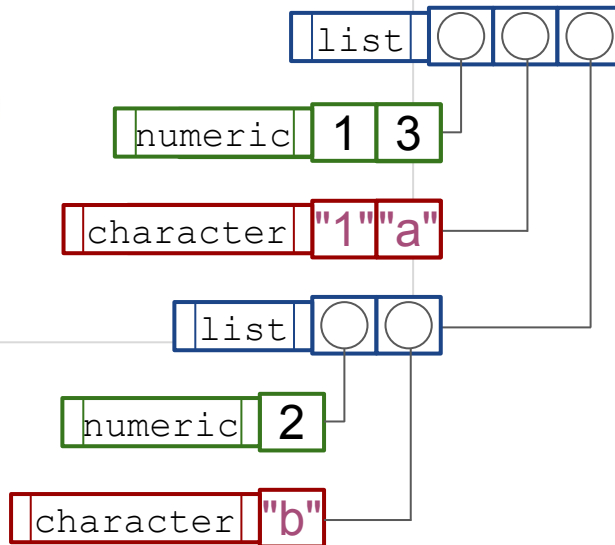
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  - (and some other things we omit for now)
- may contain any other values

```
> list(c(1, 3), c(1, "a"), list(2, "b"))
[[1]]
[1] 1 3
```

```
[[2]]
[1] "1" "a"
```

```
[[3]]
[[3]][[1]]
[1] 2
```

```
[[3]][[2]]
[1] "b"
```



# Reminder: Data Types

- atomic types: indexing `> x <- c(1, 3)`

numeric	1	3
---------	---	---

# Reminder: Data Types

- atomic types: indexing `> x <- c(1, 3)`

numeric	1	3
---------	---	---

`> x[2]`  
`[1] 3`

numeric	index	2
1		
3		

→ 

numeric	3
---------	---

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

numeric	index	1	2	1
1				
3				

→ 

numeric	1	3	1
---------	---	---	---

# Reminder: Data Types

- atomic types: indexing `> x <- c(1, 3)`

numeric	1	3
---------	---	---

`> x[2]`  
`[1] 3`

numeric	index	2
1		
3		

→ 

numeric	3
---------	---

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

numeric	index	1	2	1
1				
3				

→ 

numeric	1	3	1
---------	---	---	---

`> x[c(TRUE, FALSE)]`  
`[1] 1`

index	T	F
numeric	1	3

→ 

numeric	1
---------	---

# Reminder: Data Types

- atomic types: indexing
  - single brackets [ ]: index with integers (or character names) or logicals
  - double brackets [[ ]]: index with single integer (or single character name) only get single entry. **This is preferred if you know you just need a single entry.**

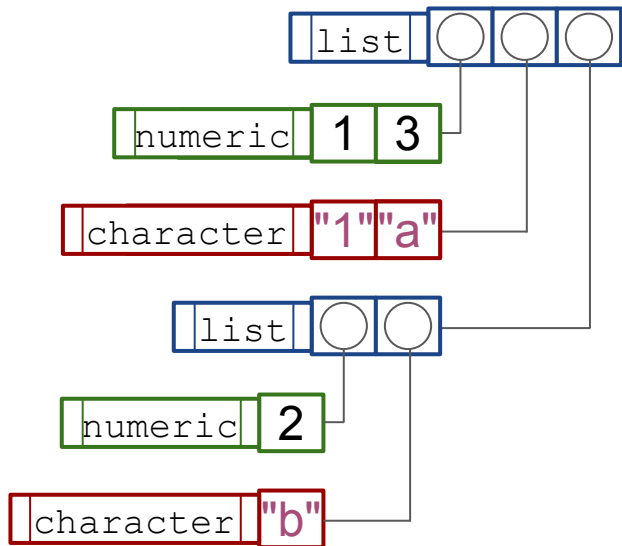
Named indexing of vectors:

```
> y <- c(a=1, b=2)
> y['a']
a
1
> y[['a']]
[1] 1
```

# Reminder: Data Types

- list indexing

```
> x <- list(c(1, 3), c(1, "a"), list(2, "b"))
```

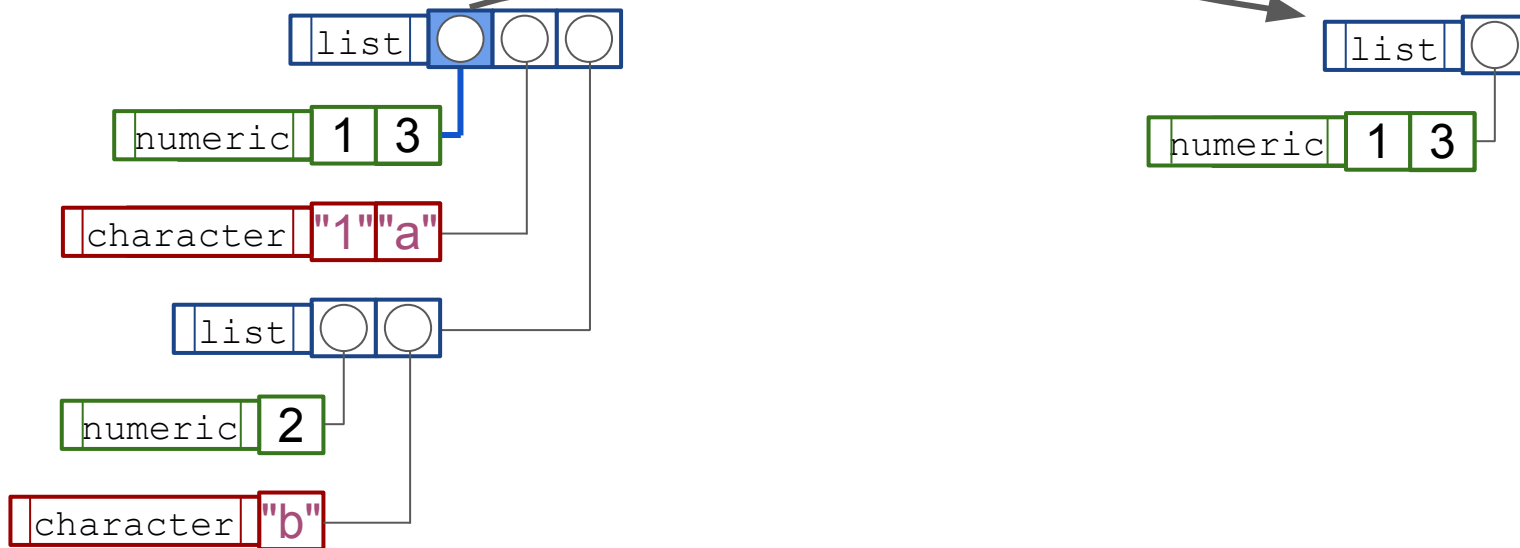


# Reminder: Data Types

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

- list indexing: [ ] just gives a smaller list

```
> x <- list(c(1, 3), c(1, "a"), list(2, "b"))
```

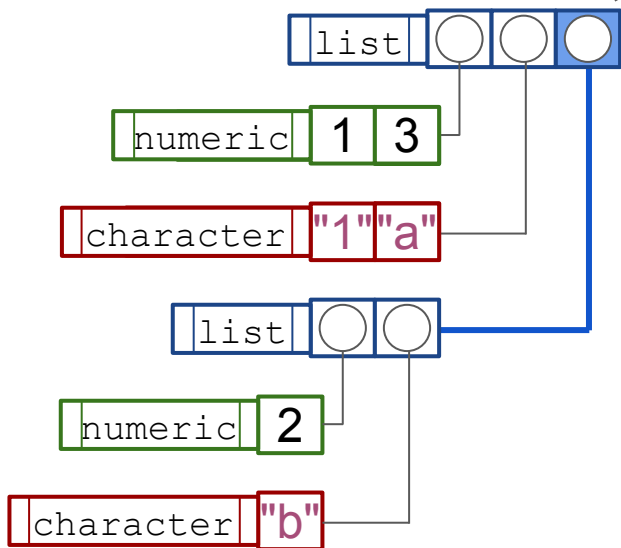




# Reminder: Data Types

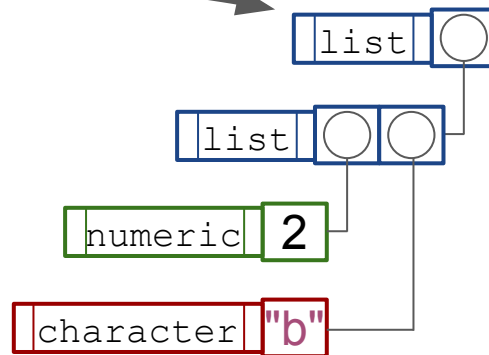
- list indexing: [ ] just gives a smaller list

```
> x <- list(c(1, 3), c(1, "a"), list(2, "b"))
```



```
> x[3]
[[1]]
[[1]][[1]]
[1] 2

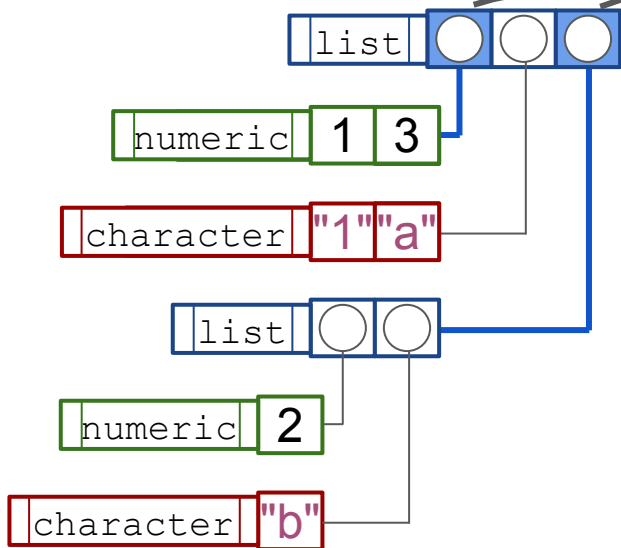
[[1]][[2]]
[1] "b"
```



# Reminder: Data Types

- list indexing: [ ] just gives a smaller list

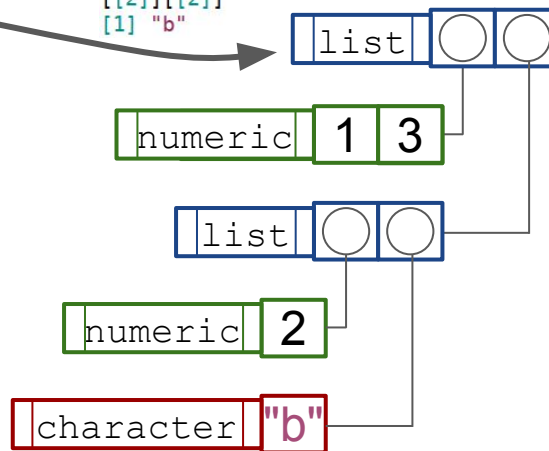
```
> x <- list(c(1, 3), c(1, "a"), list(2, "b"))
```



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

```
[[2]]  
[[2]][[1]]  
[1] 2
```

```
[[2]][[2]]  
[1] "b"
```

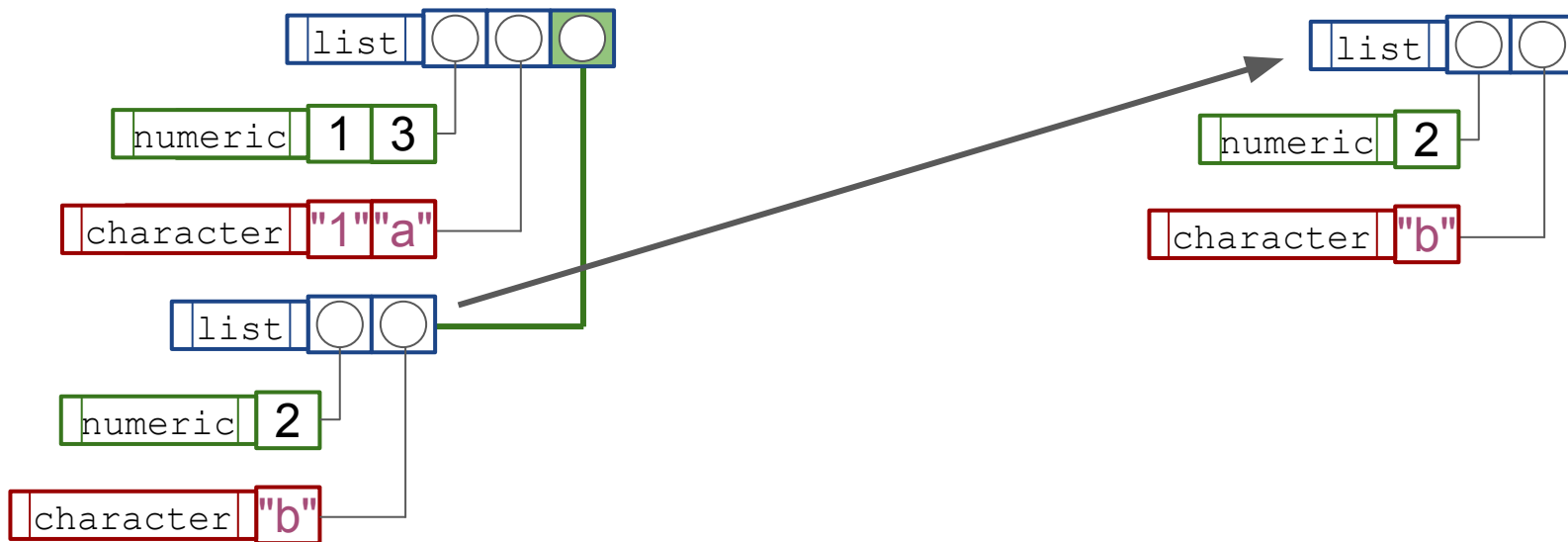


# Reminder: Data Types

- list indexing: `[[ ]]` gives *content* of list element

```
> x <- list(c(1, 3), c(1, "a"), list(2, "b"))
```

```
> x[[3]]  
[[1]]  
[1] 2  
  
[[2]]  
[1] "b"
```

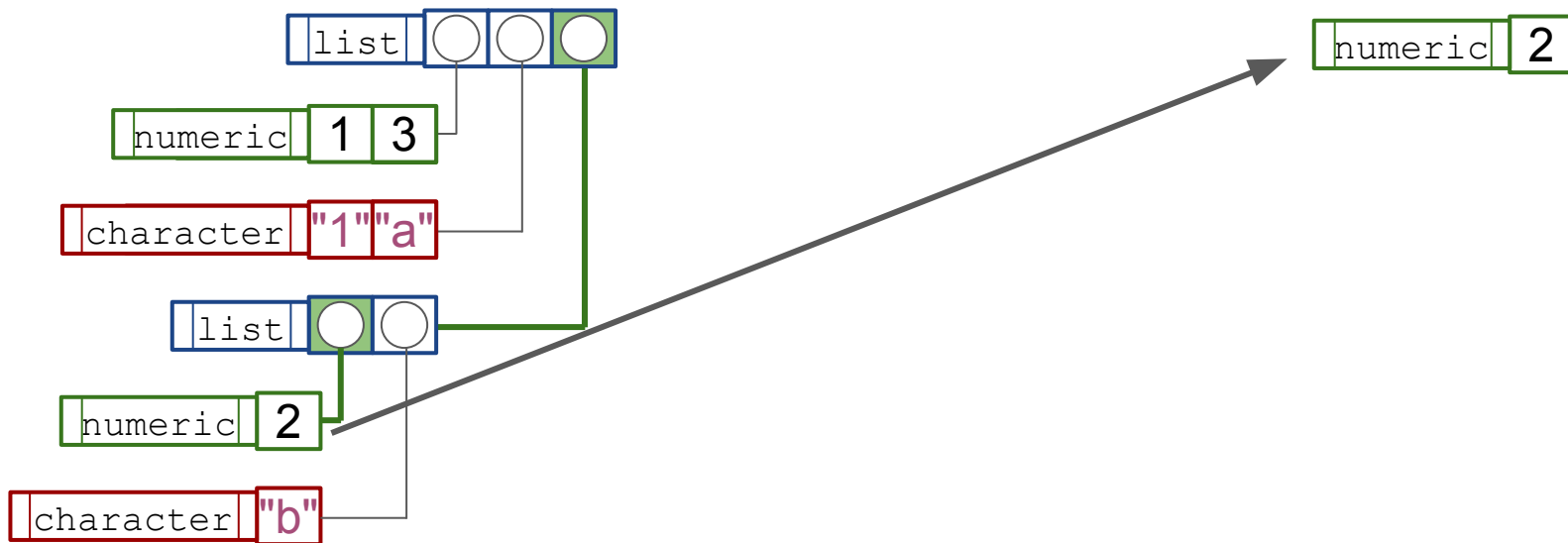


# Reminder: Data Types

```
> x[[3]][[1]]  
[1] 2
```

- list indexing: `[[ ]]` gives *content* of list element

```
> x <- list(c(1, 3), c(1, "a"), list(2, "b"))
```



# Reminder: Data Types

- list indexing:
  - `[]` just gives a smaller list
  - `[[ ]]` gives *content* of list element
  - *both* of these work with names as well
  - the `$`-operator is also common, but note the difference to `[[ ]]`:

```
> x <- list(abc = 1)
> x[["abc"]]
[1] 1
> x[["ab"]]
```

```
> x[["abc"]]
[1] 1
> x[["ab"]]
NULL
```

```
> x$abc
[1] 1
> x$ab
[1] 1
```

This is "partial matching", avoid this if possible!

# Reminder: Data Types

- "atomic" data types: `NULL`, `logical`, `numeric`, `character`
  - (also less importantly: `integer`, `complex`, `raw`)
- "recursive" data types: `list` (and some other more special things)
- many other types are just these with extra attributes!
  - `factor`, `ordered`: `integer` with "levels" attribute and class "factor" or class `c("ordered", "factor")`
  - `matrix`: `numeric` with "dim" attribute
  - `data.frame`: named `list` with "row.names" attribute and class "data.frame"
  - These have their own ways of handling access with `[ ]`, `[, ]`, `[[ ]]`, and `[[, ]]`

# Reminder: Matrix / data.frame access with [, ]

```
> x
      [,1] [,2] [,3]
[1,]     1     4     7
[2,]     2     5     8
[3,]     3     6     9
> x[c(1, 3), ] # select rows 1 and 3
      [,1] [,2] [,3]
[1,]     1     4     7
[2,]     3     6     9
> x[, c(1, 3)] # select columns 1 and 3
      [,1] [,2]
[1,]     1     7
[2,]     2     8
[3,]     3     9
```

## Reminder: Matrix / data.frame access with [, ]

```
> x <- matrix(1:9, nrow = 3)
```

```
> x
```

```
      [,1] [,2] [,3]  
[1,]    1    4    7  
[2,]    2    5    8  
[3,]    3    6    9
```

```
> x[c(1, 3), c(1, 3)]    --> Create a "slice" of columns 1 and 3, and of these slice rows 1 and 3
```

```
      [,1] [,2]  
[1,]    1    7  
[2,]    3    9
```



## Reminder: Matrix / data.frame access with [, ]

```
> x
      [,1] [,2] [,3]
[1,]    1    4    7
[2,]    2    5    8
[3,]    3    6    9

> y
      [,1] [,2]
[1,]    1    1
[2,]    1    3
[3,]    3    3

> x[y] # select elements (1,1), (1,3), (3,3) in turn
[1] 1 7 9
```

# Reminder: A matrix is just a vector

```
> x <- diag(3)
> x
      [,1] [,2] [,3]
[1,]    1    0    0
[2,]    0    1    0
[3,]    0    0    1
> x[4] <- 100
> x[8] <- 200
> x
      [,1] [,2] [,3]
[1,]    1  100    0
[2,]    0    1  200
[3,]    0    0    1
```

## Reminder: A data.frame is a list of vectors

```
> df <- data.frame(x = 1:3, y = letters[1:3], stringsAsFactors = FALSE)
> df
  x y
1 1 a
2 2 b
3 3 c
> df[[2]][1] <- "Z"
> df[[1]][2] <- 9
> df
  x y
1 1 Z
2 9 b
3 3 c
```

# Reminder: Subset Assignment

- `[ ]` and `[ , ]`, make it possible to assign multiple places in a vector at once:

```
> x <- c(1, 2, 3, 4)
> x[c(2, 3)] <- c(10, 11)
> x
[1] 1 10 11 4
```

# Reminder: Subset Assignment

- `[ ]` and `[, ]`, make it possible to assign multiple places in a vector, **matrix**, **data.frame** at once

```
> x <- diag(3)
```

```
> x
```

	[,1]	[,2]	[,3]
[1,]	1	0	0
[2,]	0	1	0
[3,]	0	0	1

```
> x[c(1, 3), c(1, 3)] <- c(100, 200, 300, 400)
```

```
> x
```

	[,1]	[,2]	[,3]
[1,]	100	0	300
[2,]	0	1	0
[3,]	200	0	400

Assign to the "slice" of rows 1, 3 and columns 1, 3

# Reminder: Subset Assignment

- `[ ]` and `[, ]`, make it possible to assign multiple places in a vector at once:

```
> x
      [,1] [,2] [,3]
[1,]    1    0    0
[2,]    0    1    0
[3,]    0    0    1
> y
      [,1] [,2]
[1,]    1    1
[2,]    1    3
[3,]    3    3
> x[y] <- c(100, 200, 300)
> x
      [,1] [,2] [,3]
[1,]  100    0  200
[2,]    0    1    0
[3,]    0    0  300
```

# Reminder: Loops

- "for"-loop: `for (index in <vector/list>) { ... }`

# Reminder: Loops

- "for"-loop: `for (index in <vector/list>) { ... }`

DO NOT use:

```
for (index in 1:number) { ... }
```

this has bad behaviour when `number` is 0!

DO use:

```
for (index in seq_len(number)) { ... }
```



# Reminder: Loops

- "for"-loop: `for (index in <vector/list>) { ... }`

DO NOT use:

```
for (index in 1:len(vector)) { ... }
```

this has bad behaviour when `vector` is empty!

DO use:

```
for (index in seq_along(vector)) { ... }
```

# Reminder: Loops

- "for"-loop: `for (index in <vector/list>) { ... }`
- "while"-loop: `while (condition) { ... }`
- "repeat"-loop: infinite loop. `repeat { ... }`

Control flow within loops:

- break out of loop: `break`
- continue to next round of the loop: `next`

# Reminder: Conditionals

- "if"-conditional: `if (condition) { ... } else { ... }`

Similar:

- ifelse-function: `ifelse(conditions, ..., ...)`

The difference: `ifelse` *vectorizes*:

```
> ifelse(c(TRUE, FALSE, TRUE), letters, LETTERS)
[1] "a" "B" "c"
```

=> use `ifelse` *only* if you have a *vector* of logicals.

=> use `if` otherwise. The following does *not* work with `ifelse`:

```
> condition <- TRUE
> x <- if (condition) c(1, 2, 3) else c(3, 2, 1)
> x
[1] 1 2 3
```

# Reminder: Conditionals

## Vectorization of Logicals

- Be aware: sometimes you want to work with *scalar truth values*, sometimes with *vector truth values*
  - Scalar truth values: e.g. `while`, `if`
  - Vector truth values: e.g. `ifelse`, vector indexing `x[c(TRUE, FALSE, TRUE)]`
- Different operators for them!
  - Operators for *vector truth values*: `|`, `&`.
  - Operators for *scalar truth values* `||`, `&&`.
- Be aware that `==` and similar gives *vector truth values* if comparing more than one value. use `all(x == y)`, `identical(x, y)` or `isTRUE(all.equal(x, y))` instead!

# What We Expect You to Know

## Structured Programming I

Know the material from the last lecture! In particular

- What the different types of vectors are, and what differentiates them from lists
- That matrices and data.frames are internally vectors and lists with special attributes
- Different ways of accessing and setting elements in vectors, lists, matrices, and data.frames
- Vectorization and recycling by operators
- Loops: for, while, repeat, next, break
- Conditionals: if, else, switch(), ifelse()
- Logical operators, differences between &,&| and &&,&&|| and when each is needed

# What We Expect You to Know

## Structured Programming I

Learn about these useful functions included in R that you may not have known before. `help()` is always a good start for this!

- Sequences (similar to colon operator): `seq_len`, `seq_along`, `seq`
- Sets: `setdiff`, `union`, `unique`, `setequal`, `duplicated`, `anyDuplicated`, `table`
- Indexing & finding things: `which`, `which.max`, `which.min`, `max`, `min`, `pmax`, `pmin`, `match`, `row`, `col`
- Logic operations: `all`, `any`, `identical`, `all.equal`, `isTRUE`, `isFALSE`, `xor`
- Value transformations: `cut`, `hist`, `diff`, `floor`, `ceiling`, `round`, `trunc`
- Vector reordering: `head`, `tail`, `append`, `rep`, `rep_len`, `rev`, `sort`, `order`, `sample`
- Matrix creation and manipulation: `rbind`, `cbind`, `diag`, `expand.grid`
- Type and type conversion: `is.<TYPE>`, `as.<TYPE>`, `anyNA`, `is.na`, `is.finite`, `mode`, `type`, `typeof`, `ordered`, `factor`, `unlist`, `class`
- Matrix / Dim info: `dim`, `colnames`, `rownames`, `dimnames`, `nrow`, `ncol`, `NROW`, `NCOL`, `length`, `names`
- String operations: `regexpr`, `gregexpr`, `grep`, `grepl`, `sub`, `gsub`, `regexec`, `nchar`, `substr`, `strsplit`, `sprintf`, `toupper`, `tolower`, `paste`, `paste0`
- Functional operations (more on these in the next unit) `do.call`, `replicate`, `lapply`, `sapply`, `tapply`, `vapply`, `mapply`, `rapply`, `Map`, `Filter`, `apply`

# Dev Track Programming Style

# Programming Style

- As was already said in the first lecture, we not only want to teach you R itself, but also how to write software well.
- **Programming Style:** Suppose you know what sequence of commands you want R to execute.
  - How should you write these commands down?
  - What formatting do you use?
  - How do you break your code up into functions?
  - How do you name these functions?

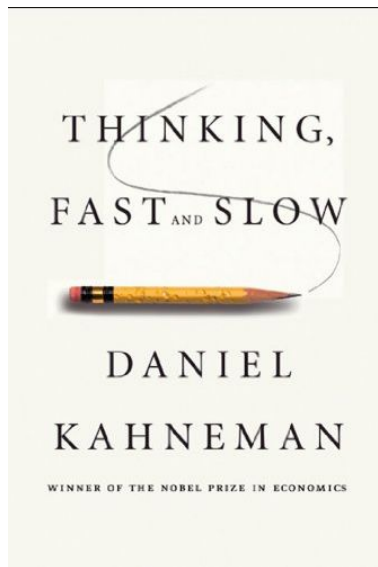


# Programming Style

- The program you write will not only be read by your computer, but also by humans:
  - Other people. Even if you are not (planning to) collaborate with others, eventually you will.
    - Unlike some physical tasks, programming naturally lends itself to division of labour. When you build a chair for yourself and your friend also wants a chair, you have to put in the same effort again. When you write a program to do your data analysis for your thesis, and your friend wants to do the same analysis, then he can just copy your code... and you better hope you wrote the program in an understandable way!
  - The future "you".
    - Look at your calendar from a year ago and try to remember what you meant with all the different one-word entries you wrote down in a hurry.
    - Future you in a long enough time is essentially a stranger who will hate you if they can't read your code.
- Adapt your code to be readable and easily adaptable!

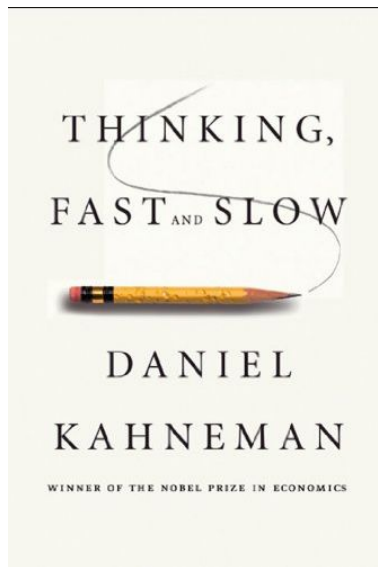
# A Short Detour to Human Psychology

- Kahneman: Thinking, Fast and Slow
  - "System 1": emotional, stereotypic, unconscious, going by first appearances, fast, automatic
  - "System 2": logical, calculating, conscious effort, slow, effortful



# A Short Detour to Human Psychology

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  - "System 1": emotional, stereotypic, unconscious, going by first appearances, fast, automatic
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- Computers do "System 2"!
  - does not care about what your code "looks like"
  - there is no ambiguity, just does the calculation

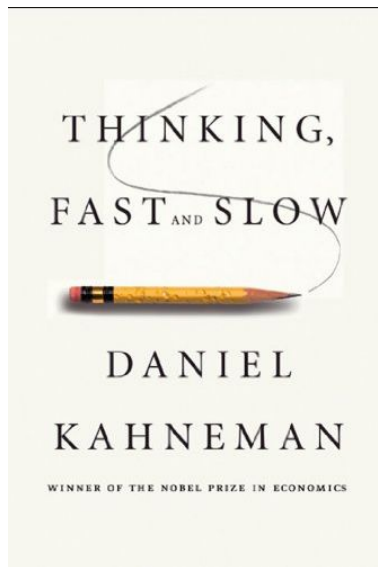


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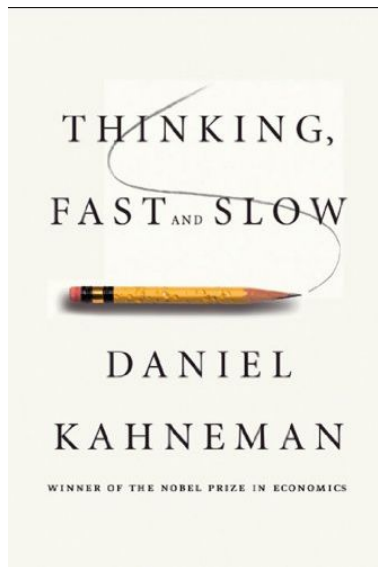
Computers have no problems with this:

```
f <- (function(f) (function(x) f(x(x)))(function(x) f(x(x))))(  
  function(x) function(n) if (n < 1) 1 else n * x(n - 1))
```



# A Short Detour to Human Psychology

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  - "System 2": logical, calculating, conscious effort, slow, effortful
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  - does not care about what your code "looks like"
  - there is no ambiguity, just does the calculation
- Humans prefer "System 1!"
  - judge by appearances and familiarity

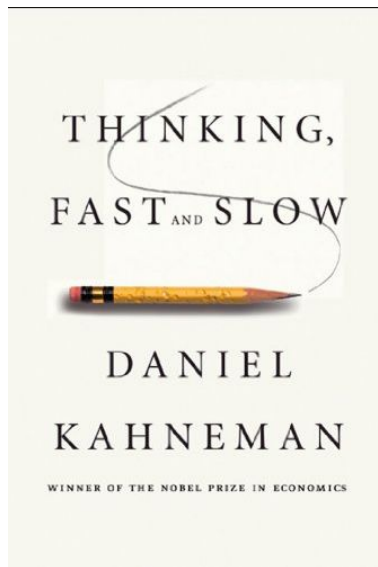


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  - does not care about what your code "looks like"
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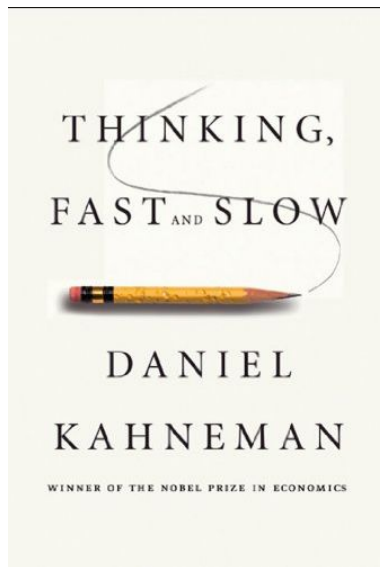
equivalent to  
the function  
two slides  
before:

```
factorial <- function(n) {  
  if (n < 1) {  
    return(1)  
  } else {  
    return(n * factorial(n - 1))  
  }  
}
```



# A Short Detour to Human Psychology

- Kahneman: Thinking, Fast and Slow
  - "System 1": emotional, stereotypic, unconscious, going by first appearances, fast, automatic
  - "System 2": logical, calculating, conscious effort, **slow, effortful**
- Computers do "System 2"!
  - does not care about what your code "looks like"
  - there is no ambiguity, just does the calculation
- Humans prefer "System 1!"
  - judge by appearances and familiarity
- Your program must satisfy both: System 2 (or the computer will not do what you want) and System 1 (Or it will be slow and effortful to understand what is happening)



# Programming Style (I): Avoid Unnecessary Noise & Variation



# Programming Style

## Avoid unnecessary Noise & Variation

```
for (i in seq_len(rows-1)){  
  vertex <- sampleVertex( 3 )  
  point=  
mat[ i , ]; next.point = stepToVertex(point,vertex,0.5)  
  mat[i + 1,]<-next.point}
```

- Come on, it's just ugly. Have some pride in your work!

# Programming Style

## Avoid unnecessary Noise & Variation

```
for (i in seq_len(rows - 1)) {  
  vertex <- sampleVertex(3)  
  point <- mat[i, ]  
  next.point <- stepToVertex(point, vertex, 0.5)  
  mat[i + 1, ] <- next.point  
}
```

- Much more regular!

# Programming Style

## Avoid unnecessary Noise & Variation

```
for (i in seq_len(rows - 1)) {  
  vertex <- sampleVertex(3)  
  point <- mat[i, ]  
  next.point <- stepToVertex(point, vertex, 0.5)  
  mat[i + 1, ] <- next.point  
}
```

- don't write `a+b` at one point and `a + b` at another etc.
- this is sometimes referred to as "**code style**" in the narrower sense

# Programming Style: Code Style

- where to put your parentheses (), braces {}, brackets []
- where to put your spaces, and how many of them
- how to name your variables and functions

# Programming Style: Code Style

- where to put your parentheses (), braces {}, brackets []
- where to put your spaces, and how many of them
- how to name your variables and functions

e.g.

```
if (x == 1) {  
    y <- 2  
}
```

vs.

```
if (x==1)  
    {  
        y=2  
    }
```

# Programming Style: Code Style

- where to put your parentheses (), braces {}, brackets []
- where to put your spaces, and how many of them
- how to name your variables and functions

e.g.

```
if (x == 1) {  
    y <- 2  
}
```

vs.

```
if (x==1)  
    {  
        y=2  
    }
```

- could both be fine as long as you are consistent
- should be chosen for easy readability, but to some degree a matter of aesthetic preferences

# Programming Style: Code Style

- In this course, your code style will be **checked automatically**.
  - Your code deviates from what we dictate --> no points
  - But don't worry, the checker will give you informative feedback!

# Programming Style: Code Style

- In this course, your code style will be **checked automatically**.
  - Your code deviates from what we dictate --> no points
  - But don't worry, the checker will give you informative feedback!
- We **mostly** use [Google's \(old\) style guide](#):
  - Well written, short and to the point without being ambiguous
  - We add the **following alterations**:
    - Function naming: use `lowerCamel` for *functions* and `UpperCamel` for *classes* (you will see classes in a few weeks)
    - Line length: limit to 120 characters instead of 80. Welcome to the future!
    - Indentation: indent one level deeper per `{`, `(` or `[` one level up per `}`, `)` or `]`. Lines starting with closing parens get the reduced indentation level.
    - Don't use `return()` at the end of functions if you don't need to
    - Put comments describing the actions of a function *before* the function, to be consistent with roxygen2 which we will see later in the course.
    - Plus some good practices that you will find out about when checking your answers ;-)



# Programming Style: Code Style

- Notable other style guides:
  - ["tidyverse"](#) style guide.
    - Mostly similar, but much more to read. Largely concerned with "tidyverse"-internal extensions to R that are a bit niche.
    - Notable difference to what we do: they use `snake_case`, while we avoid underscores in variable and function names
  - [Style of the "mlr"-package](#)
    - Not that notable in the wide scheme of things, but the Bischl group at LMU uses it, so you will probably see it at some point.
    - Prominent differences to what we (and most other people) do: uses "=" instead of "<-" (which has some drawbacks). Make integers explicit with the "L" suffix.
- If this interests you, there is a [study on prevalence of different styles in the R community by Chia-Yi Yen et al.](#)

Programming Style (II):  
Use All Communication Channels  
Available to You

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names

Not so nice: single letters

```
f <- function(l) {  
  p <- rep(TRUE, l)  
  p[1] <- FALSE  
  for (m in seq_len(sqrt(l))) {  
    if (!p[m]) next  
    p[seq(m * 2, l, m)] <- FALSE  
  }  
  seq_len(l)[p]  
}
```

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names

```
getPrimes <- function(upper.bound) {  
  is.prime <- rep(TRUE, upper.bound)  
  is.prime[1] <- FALSE  
  for (current.prime in seq_len(sqrt(upper.bound))) {  
    if (!is.prime[current.prime]) next  
    prime.multiples <- seq(  
      from = current.prime * 2,  
      to = upper.bound,  
      by = current.prime)  
    is.prime[prime.multiples] <- FALSE  
  }  
  seq_len(upper.bound)[is.prime]  
}
```

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names

Descriptive  
function name

English. Always.  
No exceptions.

Introduce temporary  
variable we didn't  
have before for  
clarity

call less-well-known  
positional function  
arguments by name

Descriptive  
argument name

```
getPrimes <- function(upper.bound) {  
  is.prime <- rep(TRUE, upper.bound)  
  is.prime[1] <- FALSE  
  for (current.prime in seq_len(sqrt(upper.bound))) {  
    if (!is.prime[current.prime]) next  
    prime.multiples <- seq(  
      from = current.prime * 2,  
      to = upper.bound,  
      by = current.prime)  
    is.prime[prime.multiples] <- FALSE  
  }  
  seq_len(upper.bound)[is.prime]  
}
```

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
  - Some further notes:
    - short variable names are fine if they are established practice. E.g. `"i"` as iterator in for-loops, `"tmp"` for temporary values.
    - avoid collisions with functions in base R or common packages. Don't call your function `"mean"` or `"ggplot"`. Don't call your variable `"c"` because of collision with `c(1, 2)!`
    - avoid names that look similar, e.g. only differentiated through `1` (one) vs. `l` (lowercase L) vs. `I` (uppercase i). Even if *your* system has a font that distinguishes those, someone else's might not.
    - Keep with a pattern. The pendant to `"coord.x"` should *not* be `"y.coordinate"`
    - Consider tab-completion when naming things: names should be `<general>.<special>`. `"coord.x"` is preferred, because if someone else is editing the program and wants to get a coordinate, they enter `coord.<tab>` and can see what coordinates are available.

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
- Use comments!

```
# Get prime numbers
# Uses the "Sieve of Eratosthenes" algorithm
# upper.bound: `numeric(1)` get primes to this number (inclusive)
# returns a `numeric` containing all primes from 2 to `upper.bound`.
getPrimes <- function(upper.bound) {
  is.prime <- rep(TRUE, upper.bound)
  is.prime[1] <- FALSE
  # numbers below `upper.bound` that are not primes
  # must have a divisor less than `sqrt(upper.bound)`, so
  # we don't loop further.
  for (current.prime in seq_len(sqrt(upper.bound))) {
    if (!is.prime[current.prime]) next
    prime.multiples <- seq(
```

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
- Use comments!

English. Always.  
No exceptions.

What does the  
function do?

What are its  
arguments?

What is the  
return value?

What are the  
argument /  
return **types**?

```
# Get prime numbers
# Uses the "Sieve of Eratosthenes" algorithm
# upper.bound: `numeric(1)` get primes to this number (inclusive)
# returns a `numeric` containing all primes from 2 to `upper.bound`.
getPrimes <- function(upper.bound) {
  is.prime <- rep(TRUE, upper.bound)
  is.prime[1] <- FALSE
  # numbers below `upper.bound` that are not primes
  # must have a divisor less than `sqrt(upper.bound)`, so
  # we don't loop further.
  for (current.prime in seq_len(sqrt(upper.bound))) {
    if (!is.prime[current.prime]) next
    prime.multiples <- seq(

```

Parts of your  
code that are not  
obvious?



# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
- Use comments! (**but don't overdo it**)

```
# we get a vector of `TRUE` with length `upper.bound`  
is.prime <- rep(TRUE, upper.bound)  
# `1` is not a prime  
is.prime[1] <- FALSE
```

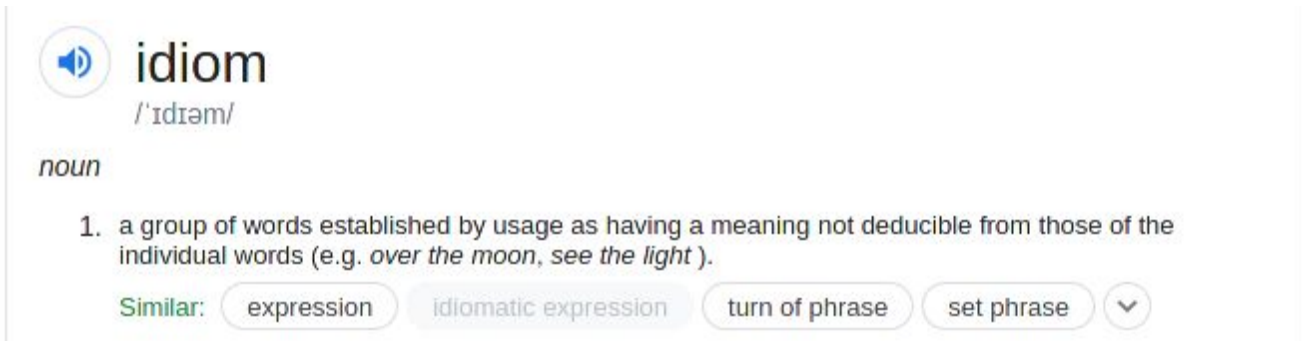
don't just translate the code to english

what happens here is obvious from good naming already


# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
- Use comments! (but don't overdo it)
- Make your code **idiomatic**



The screenshot shows the Cambridge Dictionary entry for the word "idiom". It includes a speaker icon for pronunciation, the phonetic transcription /'ɪdɪəm/, and the part of speech "noun". The first definition is: "a group of words established by usage as having a meaning not deducible from those of the individual words (e.g. *over the moon*, *see the light* )". Below the definition, there is a "Similar:" section with four buttons: "expression", "idiomatic expression", "turn of phrase", and "set phrase", followed by a dropdown arrow.

 **idiom**  
/'ɪdɪəm/  
*noun*

1. a group of words established by usage as having a meaning not deducible from those of the individual words (e.g. *over the moon*, *see the light* ).

**Similar:** expression idiomatic expression turn of phrase set phrase ▼

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
- Use comments! (but don't overdo it)
- Make your code **idiomatic**
  - express it in a natural and commonly accepted way specific to the language

~~`seq_len(upper_bound)[is.prime]`~~

"what kind of  
trick is this?"

"I've seen this  
before!"

`which(is.prime)`



# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
- Use comments! (but don't overdo it)
- Make your code **idiomatic**
  - express it in a natural and commonly accepted way specific to the language
  - Don't be "too clever"
  - Use constructs that the reader of your code is used to from other code
  - Solve problems the way someone would expect them to be solved in R

# Programming Style

## Use All Communication Channels Available to You

- Use descriptive variable / function names
- Use comments! (but don't overdo it)
- Make your code **idiomatic**
  - express it in a natural and commonly accepted way specific to the language
  - Don't be "too clever"
  - Use constructs that the reader of your code is used to from other code
  - Solve problems the way someone would expect them to be solved in R
  - **This takes experience**, you will learn common idioms by reading other code

# Programming Style (III): Your Audience has Tunnel Vision

# Programming Style

## Your Audience has Tunnel Vision

When you spent all day working on a function of course you know what is happening, but someone else should not have to spend a day to understand it

# Programming Style

## Your Audience has Tunnel Vision

Your code should be understandable

- without needing to see many lines at once



## Your code should be understandable

this is *half* of the  
randomForest() function



```

randomForest::randomForest.default
## Create a new random forest object
## @param x a matrix, or a data frame, or a list, or a file, or a URL, or a
## @param y a vector, or a factor, or a list, or a file, or a URL, or a
## @param ntree the number of trees to grow. If NULL, the default is 500.
## @param mtry the number of variables to sample at each node. If NULL, the
## @param oob a logical value. If TRUE, the out-of-bag (OOB) error is
## @param importance a logical value. If TRUE, the importance of the
## @param keep.forest a logical value. If TRUE, the forest is kept.
## @param verbose a logical value. If TRUE, the progress is printed.
## @param seed a random seed.
## @return a random forest object.
## @examples
## data(USArrests)
## rf <- randomForest(x = USArrests[, 1:4], y = USArrests[, 5], ntree = 100)
## rf
## rf$importance
## rf$varImp
## rf$varImp2
## rf$varImp3
## rf$varImp4
## rf$varImp5
## rf$varImp6
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## rf$varImp434
## rf$varImp435
## rf$varImp436
## rf$varImp437
## rf$var
```

## Your code should be understandable

- without needing to see many lines at once
  - chunk your code into reasonably sized functions
    - should ideally fit on a screen
  - and reasonably sized files

[illegible]

# Programming Style

## Your Audience has Tunnel Vision

Your code should be understandable

- without needing to see many lines at once
  - chunk your code into reasonably sized functions
    - should ideally fit on a screen
  - and reasonably sized files
- without having to keep many things in mind at once

# Programming Style

## Your Audience has Tunnel Vision

- without needing to see many lines at once
  - chunk your code into reasonably sized functions
    - should ideally fit on a screen
  - and reasonably sized files
- without having to keep many things in mind at once
  - limit your "nesting depth": avoid for-loops inside if-clauses inside for-loops inside ...
  - limit the number of args of functions. The user shouldn't have to check the docs all the time.
  - limit the role fulfilled by a single function. If you can't summarize its effect in one sentence, consider splitting it up.
  - related but more general: strive for [loose coupling](#), i.e. limit the interdependencies between parts of your code and the degree to which one part depends on specific implementation details of another part.

# What We Expect You to Know

## Programming Style

Make your program easy to read by:

1. Avoiding unnecessary noise and keeping the appearance of your code uniform.
2. Using all communication channels available to you, like variable / function names, comments, and idioms that your reader is familiar with.
3. Assuming your audience has tunnel vision and should be able to understand small blocks of your code at a time

(And remember, the code you hand in is checked for style automatically)