

# PROGRAMMING WITH R

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# INTRODUCTION

- A computer program is a sequence of commands and instructions to effectively solve a given problem.
- Each computer program is based on an underlying procedure called algorithm.
- An algorithm may be implemented in different ways, leading to different programs using the same procedure.

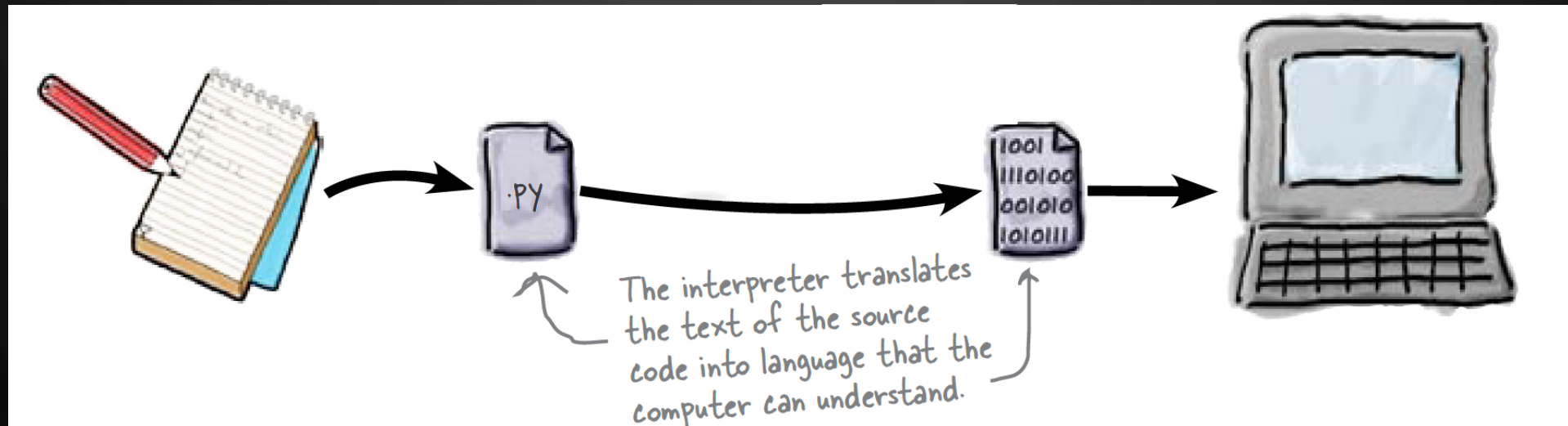
# WHY DO PROGRAMMING?

- If you use other people's software, you will always be limited by what other people think you want to do.
- Write your own programs and the only limit will be your own imagination.
- Even with all those programs on your computer, you still need to do something different, something specific to you.

# COMMON PROPERTIES

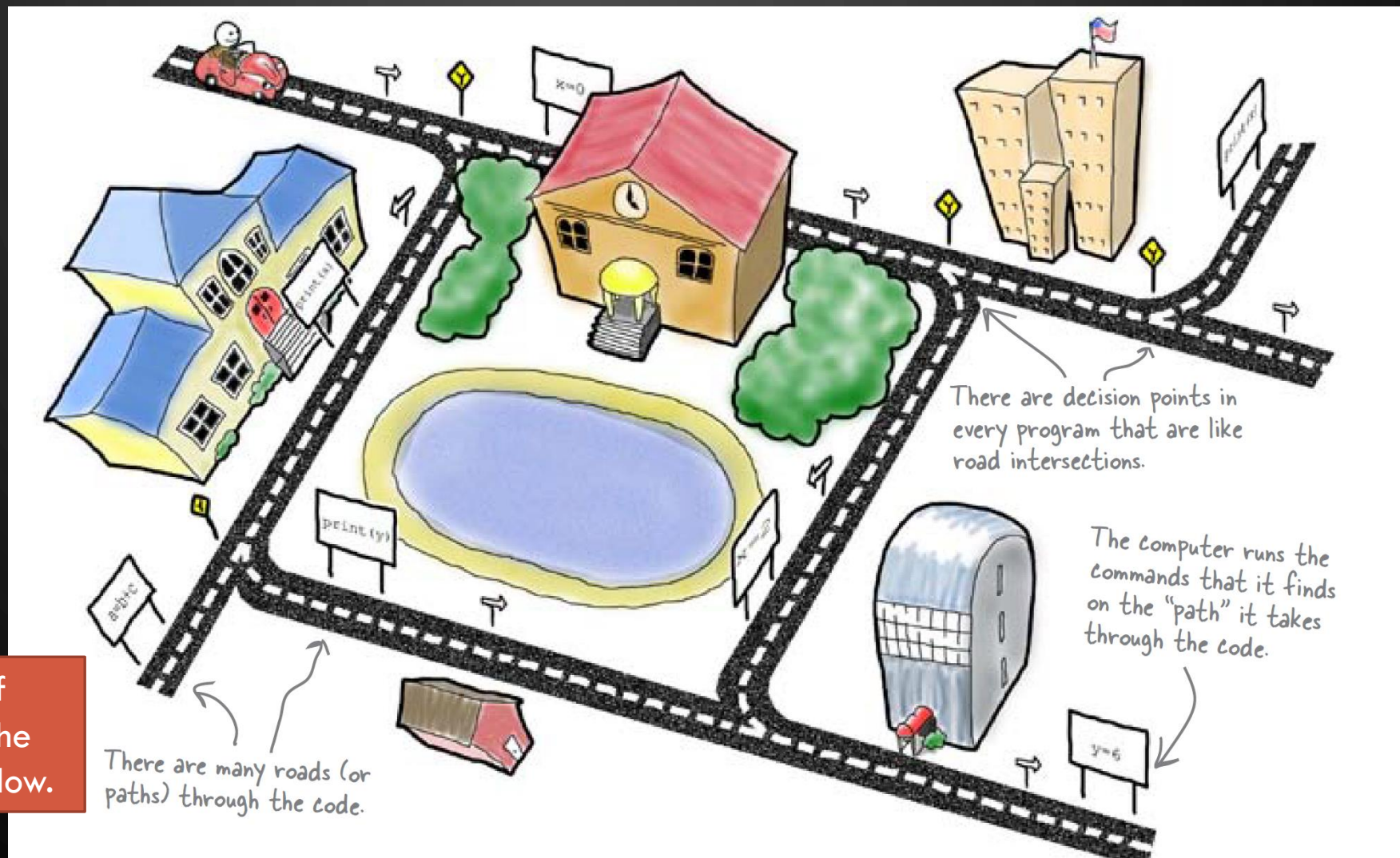
- Programs are usually written by humans and executed by computers. Hence a program should be clear, concise and direct.
- Programs are usually lists of consecutive steps.
- Programs have well-defined initiations and terminations.
- Programs have well-defined constants, inputs, outputs and variables.
- Programs are written by using a finite set of operations defined by programming languages.

# HOW DOES PROGRAMMING WORKS?





# CODEVILLE



A path is a set of instructions that the computer will follow.

# ALGORITHMS AND FLOWCHARTS

- A typical programming task can be divided into two phases:
- Problem solving phase:
  - Produce an ordered sequence of steps that describe solution of problem
  - This sequence of steps is called an algorithm
- Implementation phase:
  - Implement the program in some programming language




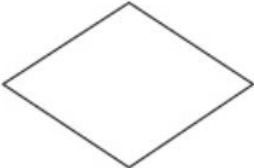


# ALGORITHMS AND FLOWCHARTS

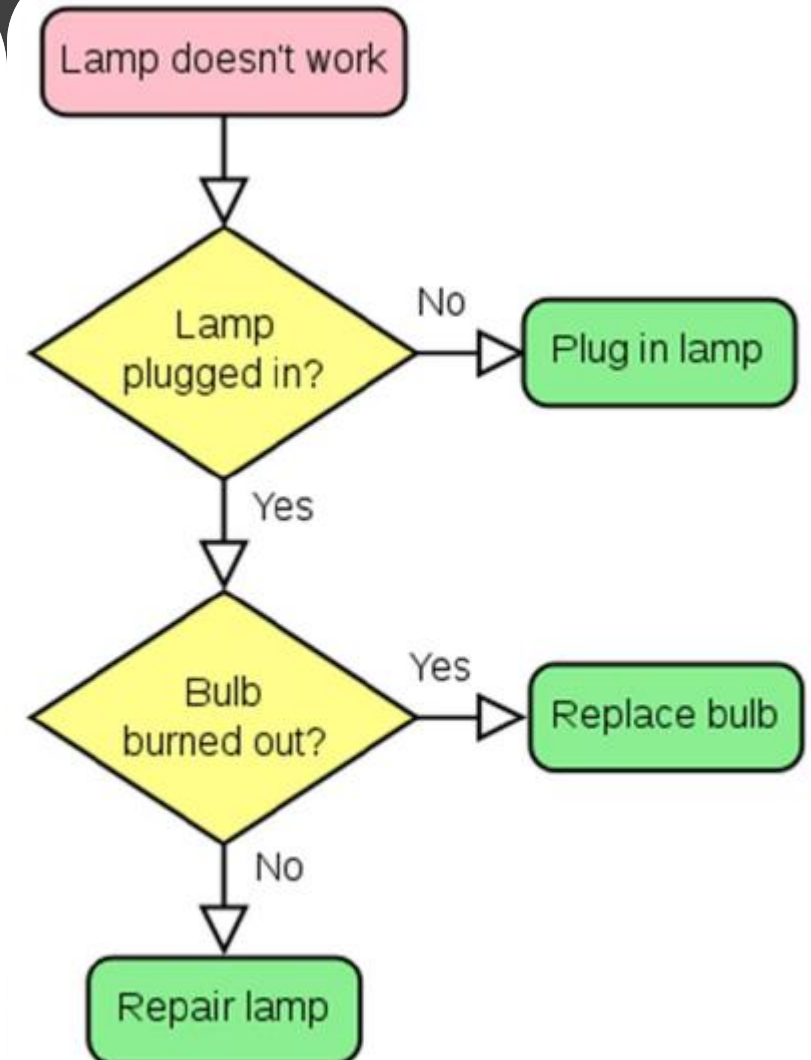
- First step in problem solving is to produce a general algorithm (one can use pseudo code).
- Pseudo code is an artificial and informal language that helps programmers develop algorithms.
- Pseudo code is very similar to everyday English.
- Program flowcharts are graphical representations show the sequence of instructions in a single program.
- A flowchart emphasizes individual steps and their interconnections.



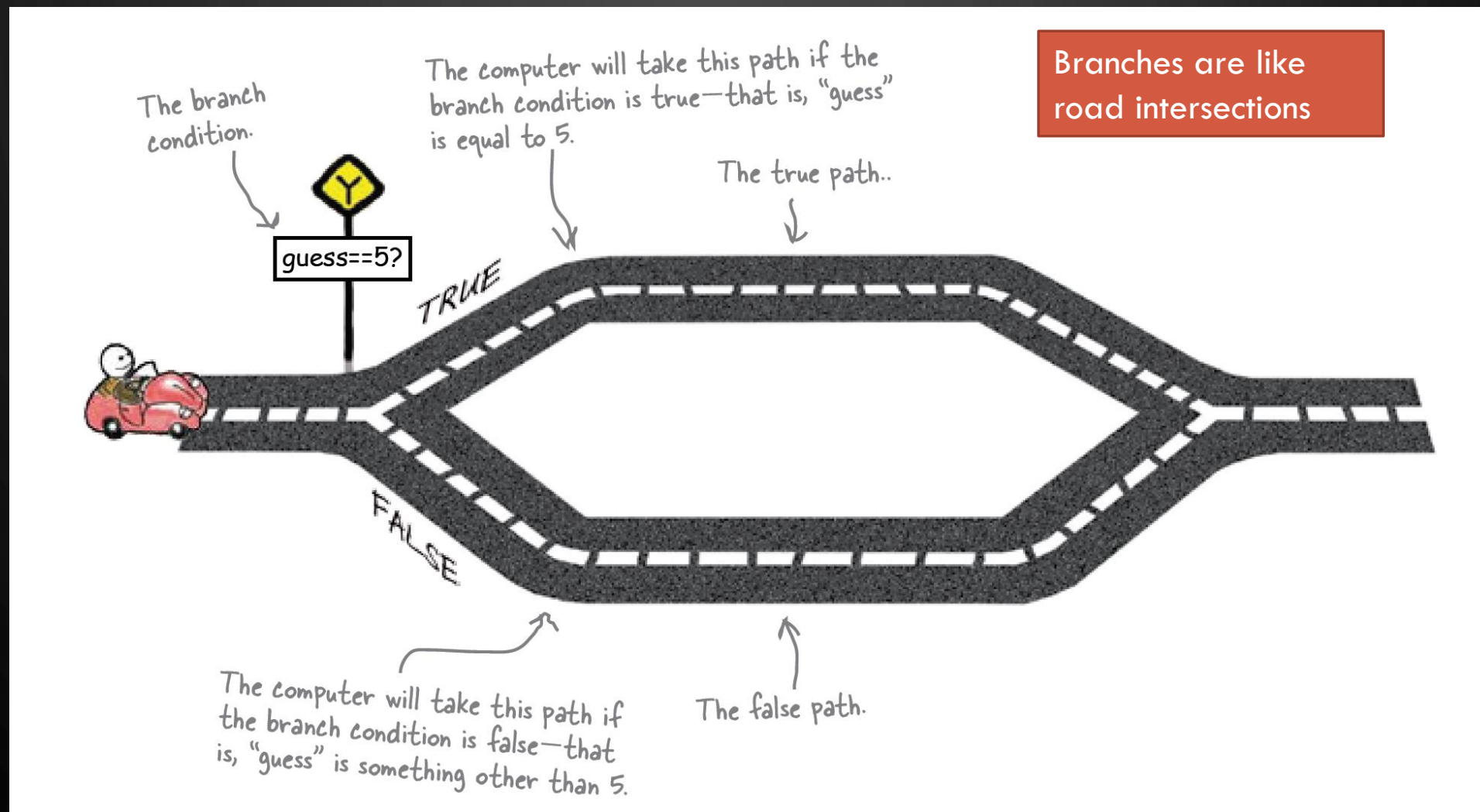
# FLOWCHART SYMBOLS

## Basic

Name	Symbol	Use in Flowchart
Oval		Denotes the beginning or end of the program
Parallelogram		Denotes an input operation
Rectangle		Denotes a process to be carried out e.g. addition, subtraction, division etc.
Diamond		Denotes a decision (or branch) to be made. The program should continue along one of two routes. (e.g. IF/THEN/ELSE)
Hybrid		Denotes an output operation
Flow line		Denotes the direction of logic flow in the program



# IF/ ELSE BRANCHES



# R IF STATEMENT

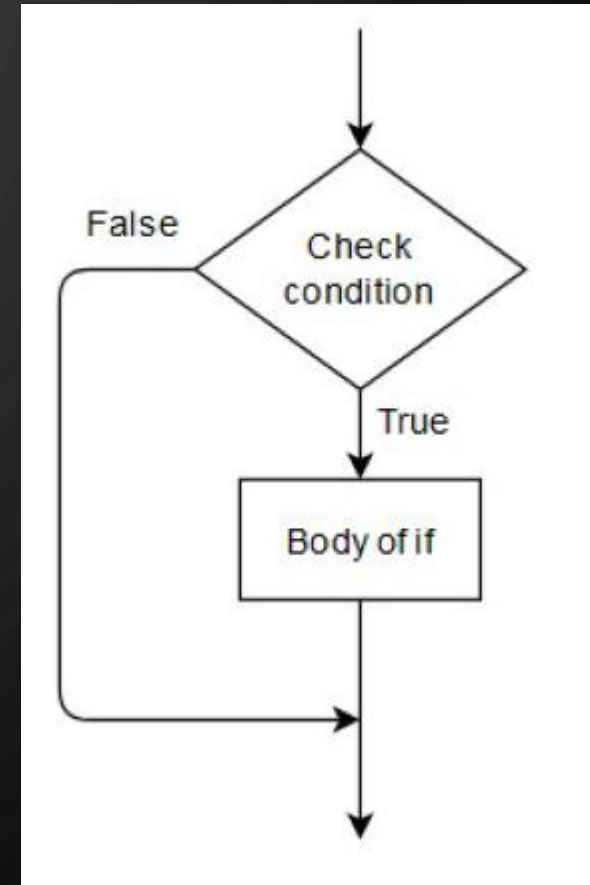
```
if (test_expression) {  
    statement  
}
```

- If the “test\_expression” is TRUE, the statement gets executed. But if it is FALSE, nothing happens.

- Example:

```
x < -5
```

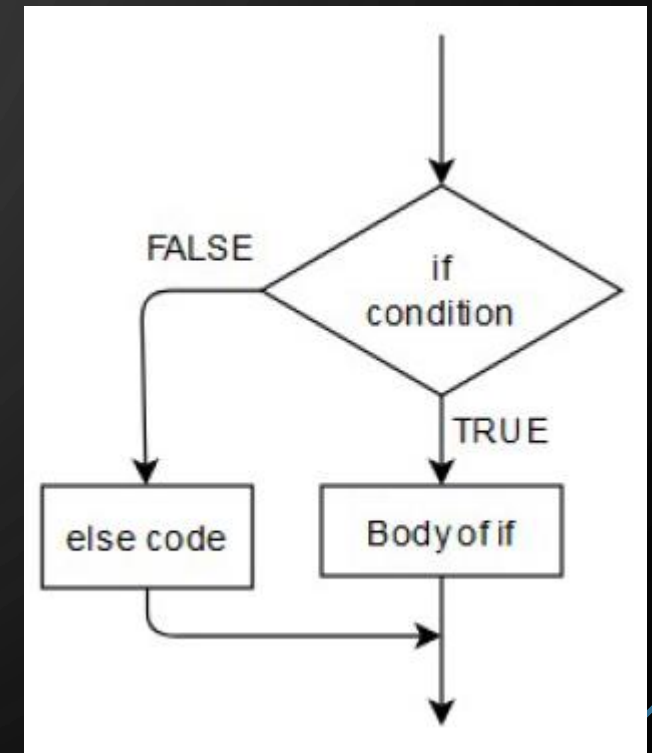
```
if (x > 0) {  
    print("x is positive")  
}
```



# R IF...ELSE STATEMENT

```
if (test_expression) {  
    statement1  
} else {  
    statement2  
}
```

- The “else” part is optional and is only evaluated if “test\_expression” is FALSE.
- It is important to note that else must be in the same line as the closing braces of the if statement.



# R IF...ELSE STATEMENT

- **Example:**

```
x<-5
```

```
if (x>=0) {
```

```
    print("x is non-negative")
```

```
} else{
```

```
    print("x is negative")
```

```
}
```

# R IF...ELSE IF STATEMENT

```
if ( test_expression1) {  
    statement1  
} else if ( test_expression2) {  
    statement2  
} else if ( test_expression3) {  
    statement3  
} else  
    statement4
```

- We can use as many if...else if statement as we want as follows.
- Only one statement will get executed depending upon the “test\_expressions”.



# R IFELSE STATEMENT

- There is a vector equivalent form of the if...else statement in R, the `ifelse()` function.

```
ifelse(test_expression,x,y)
```

- The returned vector has element from “x” if the corresponding value of “test\_expression” is TRUE or from y if the corresponding value of “test\_expression” is FALSE.

# R IFELSE STATEMENT

- Example:

```
a<-c(3,-2,5)
```

```
> ifelse(a>=0,"non-negative","negative")
```

```
[1] "non-negative" "negative" "non-negative"
```

# R SWITCH STATEMENT

- The `switch(expr,...)` function evaluates “expr” and accordingly chooses one of the further arguments.

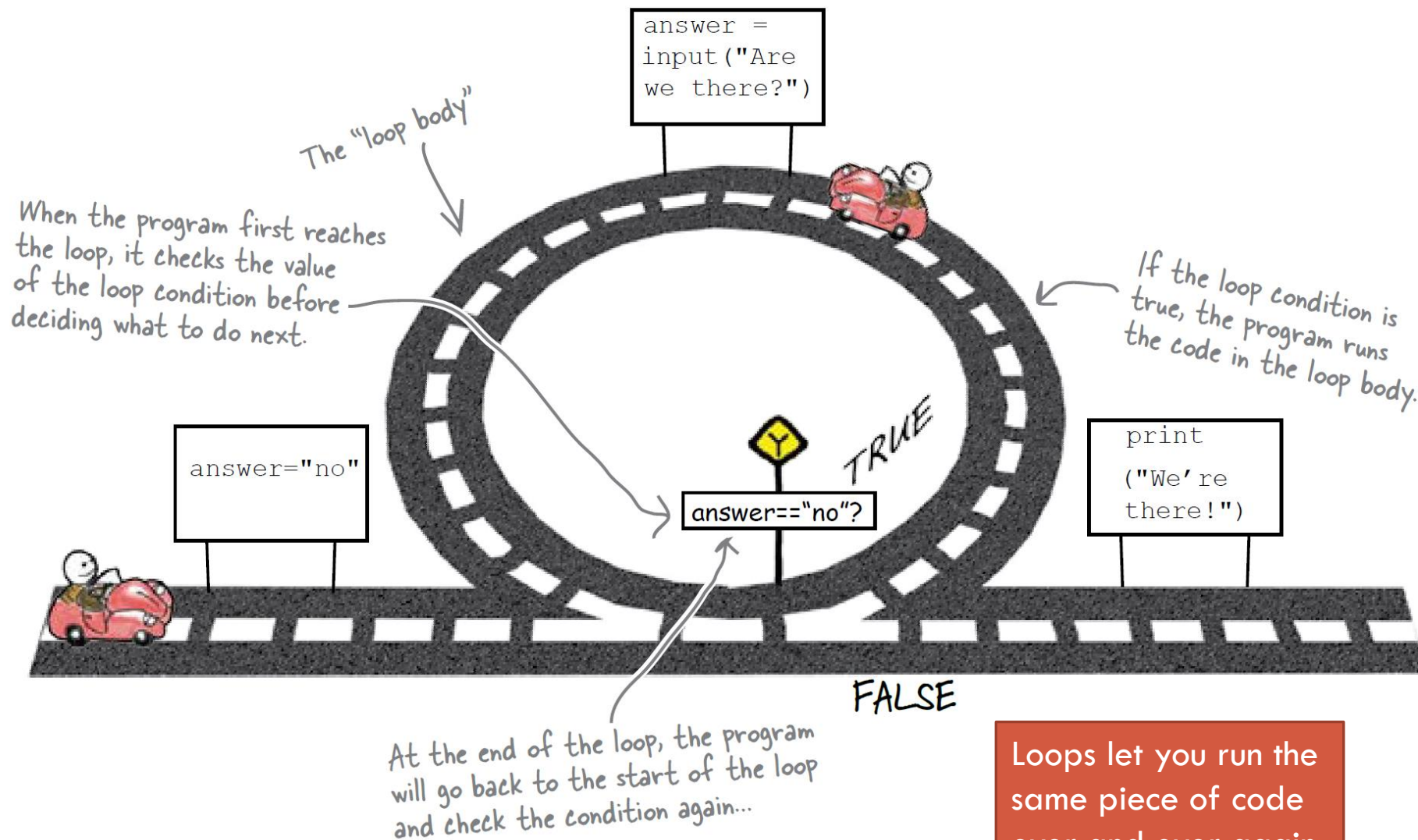
- Example:

```
x<-3
```

```
switch(x, "red", "blue", "green", "yellow")
```

```
[1] "green"
```

# LOOP



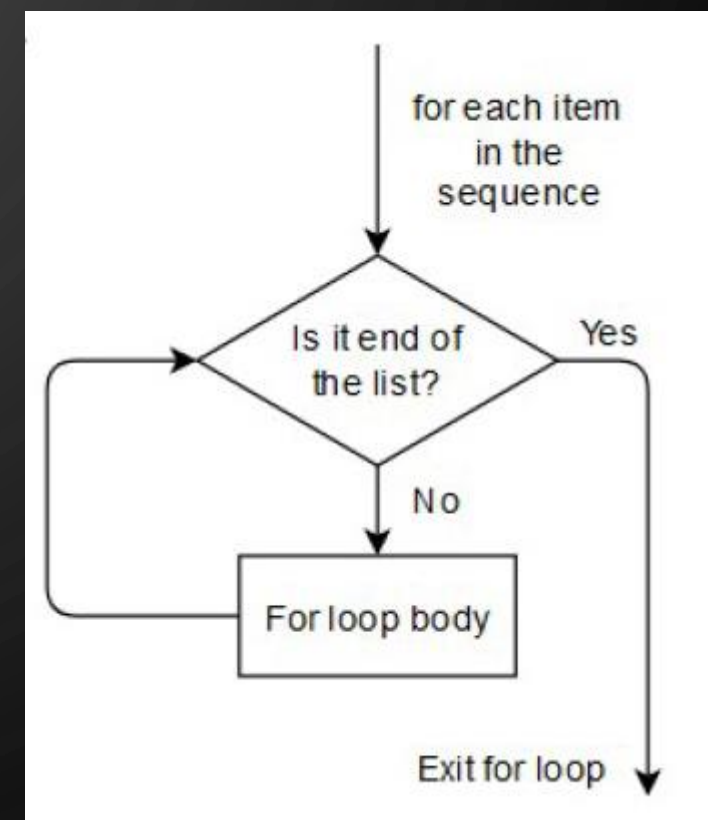
# R FOR STATEMENT

```
for (val in sequence)
{
    statement
}
```

- A for loop is used to iterate over a vector.
- In each iteration, statement is evaluated.

• **Example:**

```
x<-1
for(i in 1:3){
x<-x+1
}
x
[1] 4
```



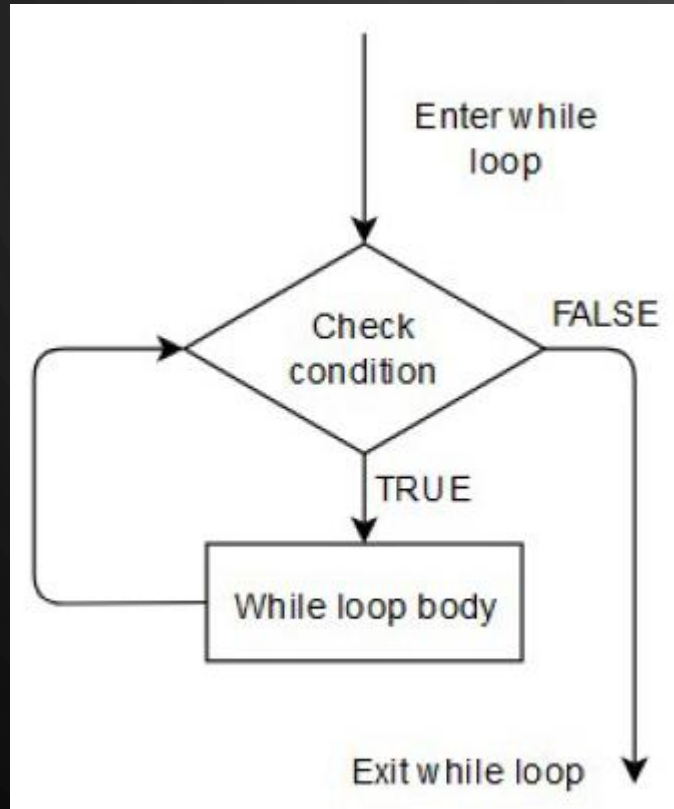
# R WHILE STATEMENT

```
while (test_expression)
{
    statement
}
```

- The “test\_expression” is evaluated and the body of the loop is entered if the result is TRUE.
- The statements inside the loop are executed and the flow returns to evaluate the “test\_expression” again.
- This is repeated each time until “test\_expression” evaluates to FALSE, in which case, the loop exits.



# R WHILE STATEMENT



- Example:

```
x<-4
```

```
while (x<=10) {
```

```
  x<-x+1
```

```
}
```

```
x
```

```
[1] 11
```

# R BREAK STATEMENT

- A break statement is used inside a loop to stop the iterations and flow the control outside of the loop.

- Example:

```
x <- 1:5  
for (val in x) {  
  if (val == 3){  
    break  
  }  
  print(val)  
}  
[1] 1  
[1] 2
```

# R NEXT STATEMENT

- A next statement is useful when we want to skip the current iteration of a loop without terminating it.
- Example:

```
x <- 1:5  
for (val in x) {  
    if (val == 3) {  
        next  
    }  
    print(val)  
}
```

[1] 1  
[1] 2  
[1] 4  
[1] 5

# PROGRAMMING IN R USING FUNCTIONS

- Programming in R is organized around functions.
- The basic declaration or definition of a function looks like below:

```
myfunction<- function(argument1, argument2, ...) {  
  # clever manipulations of arguments  
  return(values to return)  
}
```

input

statements

output

The name of the function

```
myfunction(argument1, argument2, ...)
```

Call the function

# PROGRAMMING IN R USING FUNCTIONS

- Objects that are created within the function are local to the environment of the function – they don't exist outside of the function.
- You can pass the values into the global environment with the `return()` function.
- The argument can be any type of object (scalar, matrix, vector, data frame or logical)
- A function needs to have a name and a body of code that does something.

## EXAMPLE

```
square.it<-function(x) {  
  square<-x*x  
  return(square)  
}  
  
#square a number  
square.it(5)  
  
#square a vector  
square.it(c(1,4,2))
```



# EXAMPLE

```
my.fun<-function(X.matrix, y.vec, z.scalar){  
  #use previous function  
  sq.scalar<-square.it(z.scalar)  
  
  mult<-X.matrix%*%y.vec  
  Final<-mult*sq.scalar  
  return(Final)  
}  
  
my.mat<-cbind(c(1,2,3),c(3,4,5))  
my.vec<-c(5,6)  
  
my.fun(X.matrix=my.mat, y.vec=my.vec, z.scalar=9)
```

# GOOD FUNCTION WRITING PRACTICES

- Keep your functions short.
  - If things start to get long, you can probably split up the function into several functions.
  - It also makes your code easier to update.
- Put in comments on what are the inputs, what the function does and what is the output.
- Check for errors along the way.

# THANK YOU