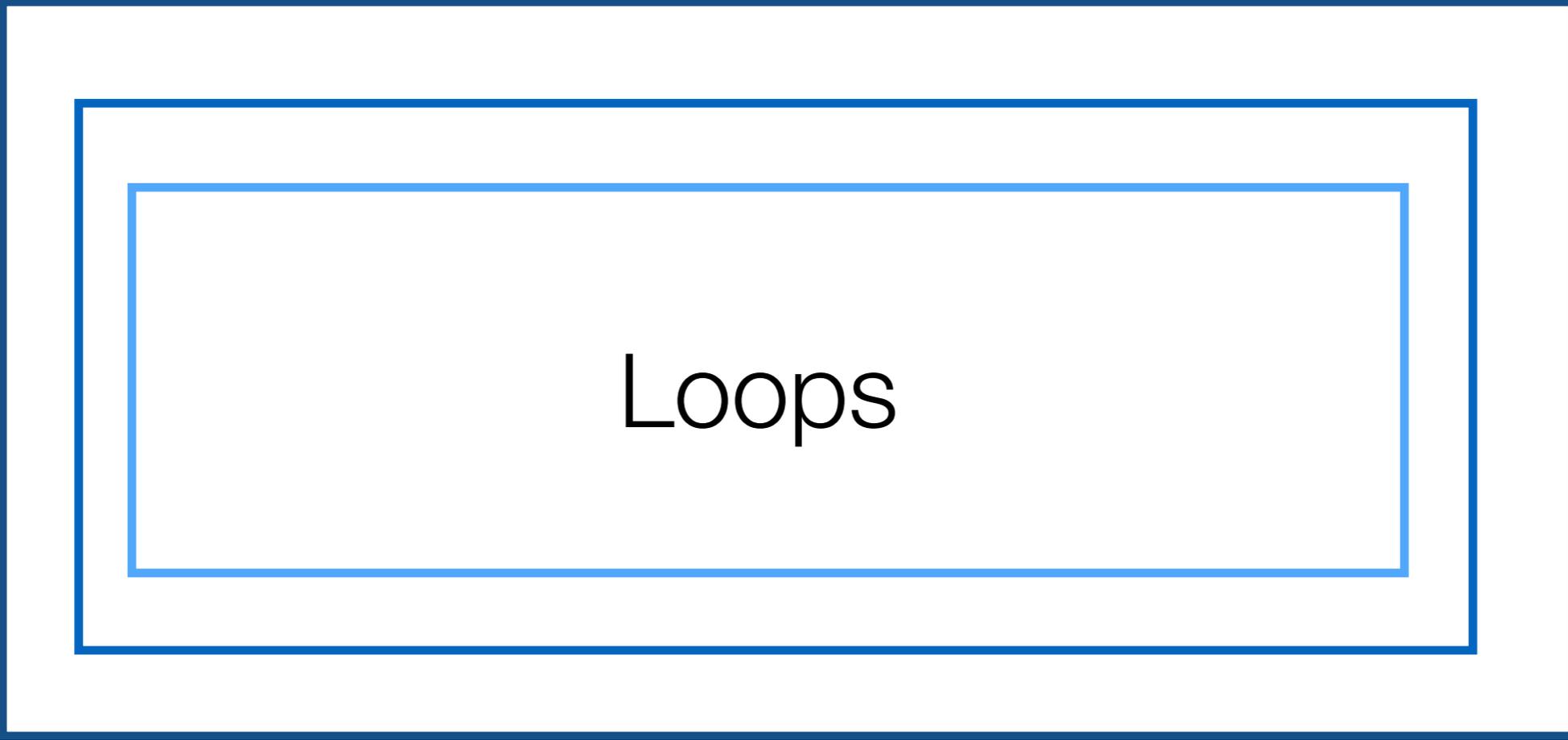


R Bootcamp Part 3

Dani Navarro
Amy Perfors

Today's Plan

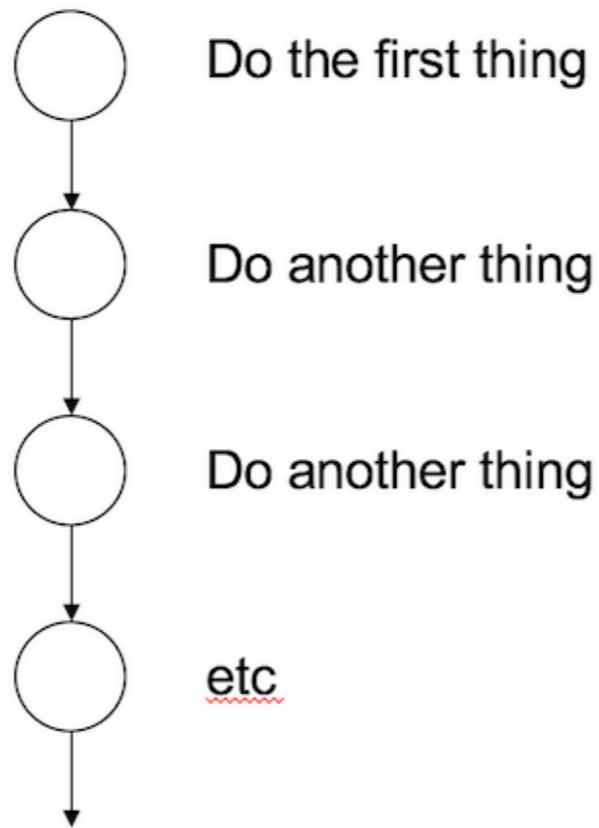
1. Loops
2. Branches
3. Functions
4. Programming
5. File system



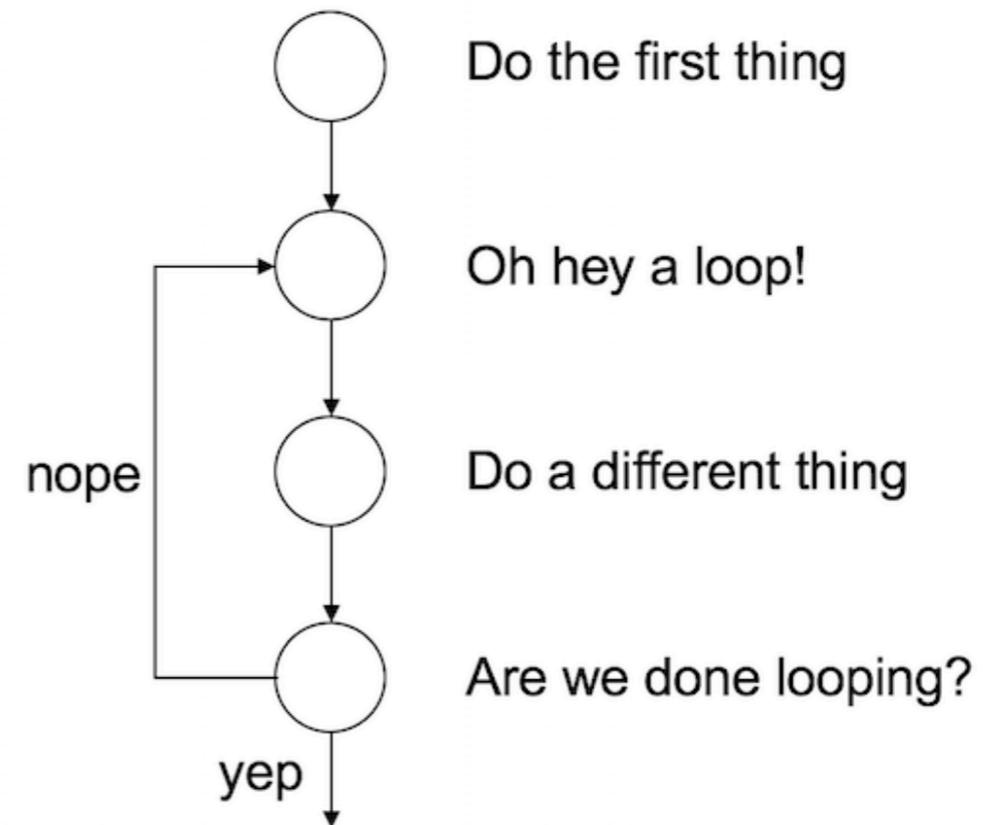
Loops

The purpose of a loop

How R reads a script without loops



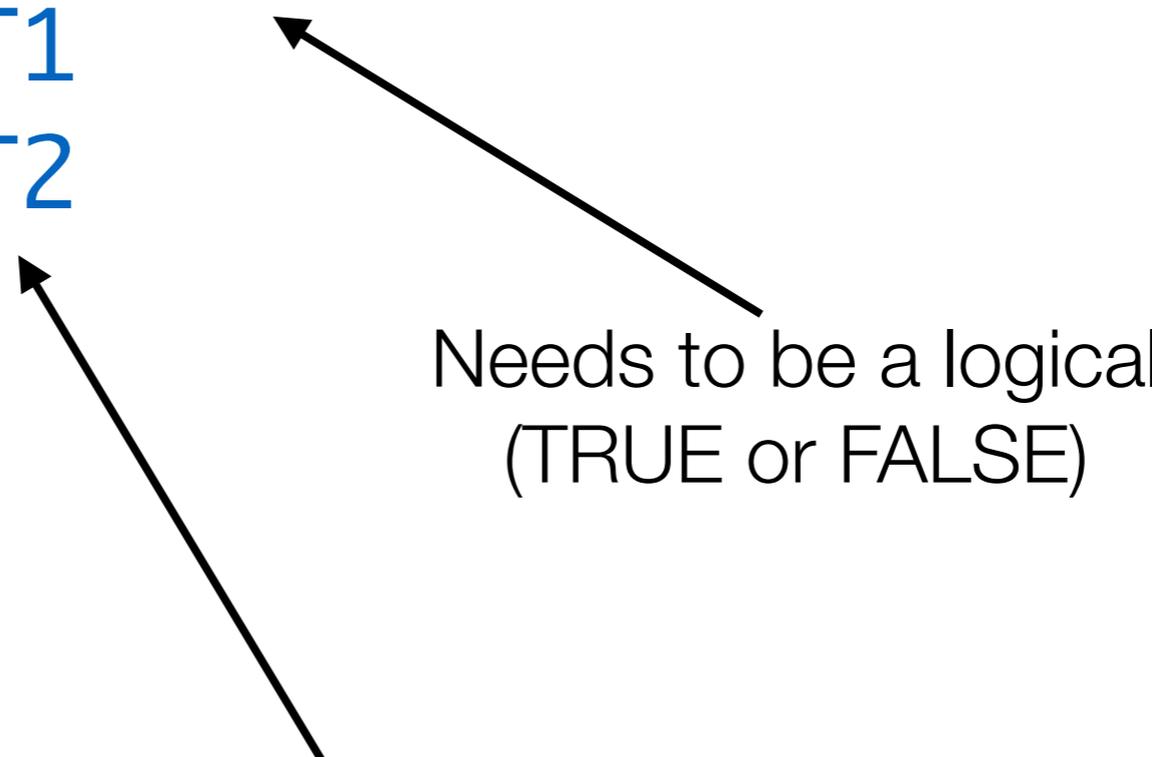
How R reads a script with a loop



While loops

```
while ( CONDITION ) {  
    STATEMENT1  
    STATEMENT2  
    ETC  
}
```

Needs to be a logical
(TRUE or FALSE)

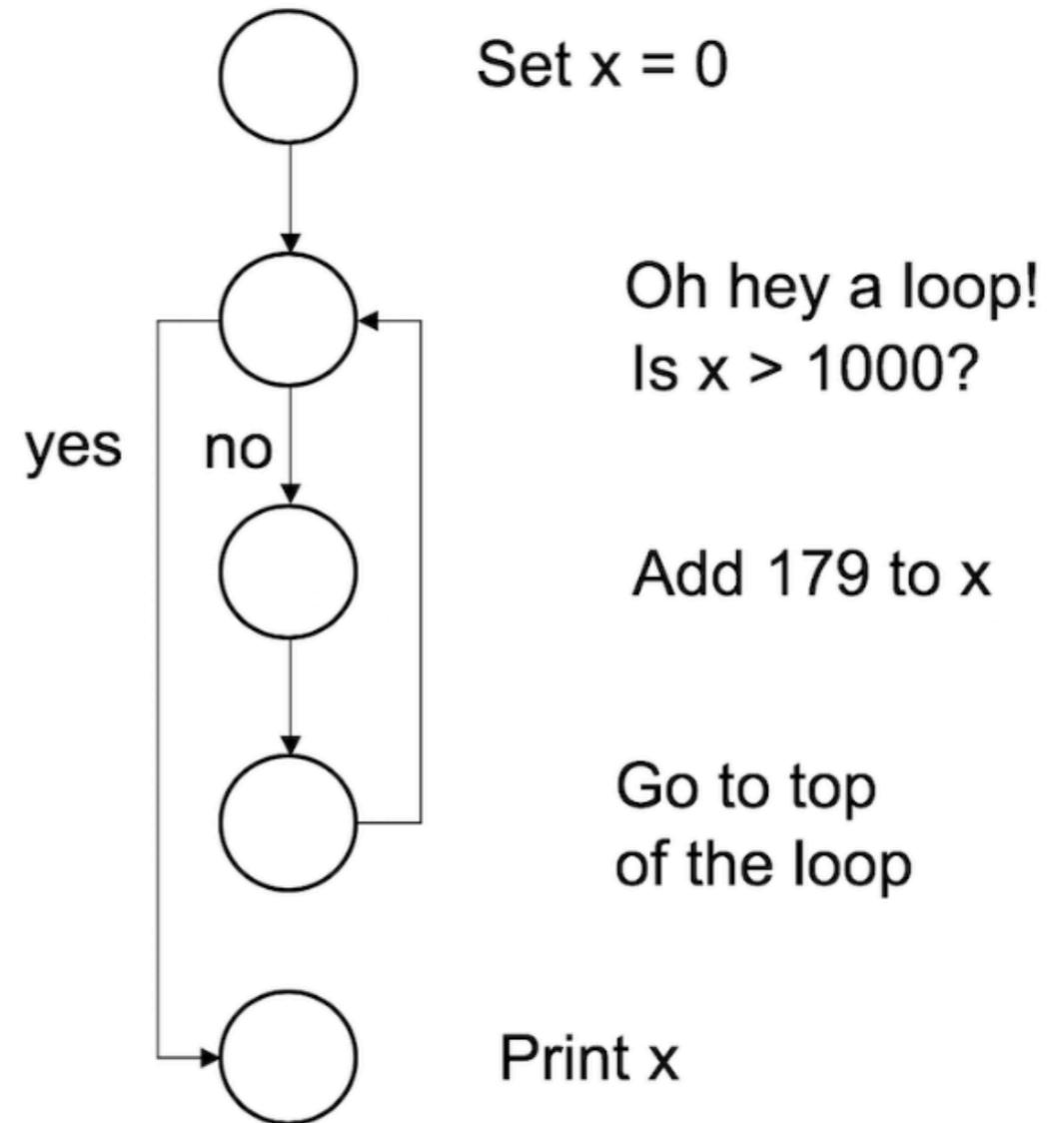


Does all of these
things as long as the
condition is TRUE

While loops

```
x <- 0  
while (x < 1000) {  
  x <- x + 179  
}  
print(x)
```

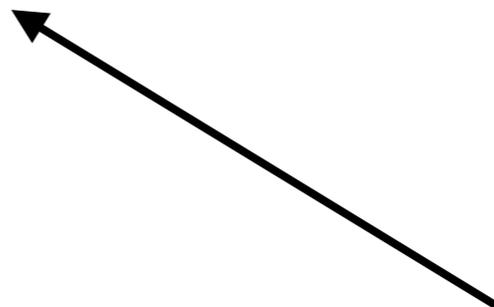
```
## [1] 1074
```



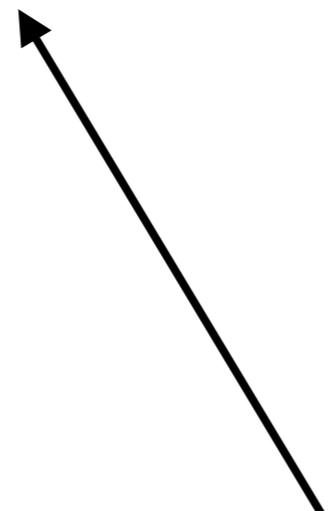
For loops

```
for ( VAR in VECTOR ) {  
    STATEMENT1  
    STATEMENT2  
    ETC  
}
```

Counts through each
of the things in the
vector



Does all of these
things as long as the
condition is TRUE



For loops

```
for ( value in 1:10 ) {  
  answer <- 137*value  
  print(answer)  
}  
# 137  
# 274  
# 411  
# 548  
# 685  
# 822  
# 959  
# 1096  
# 1233  
# 1370
```

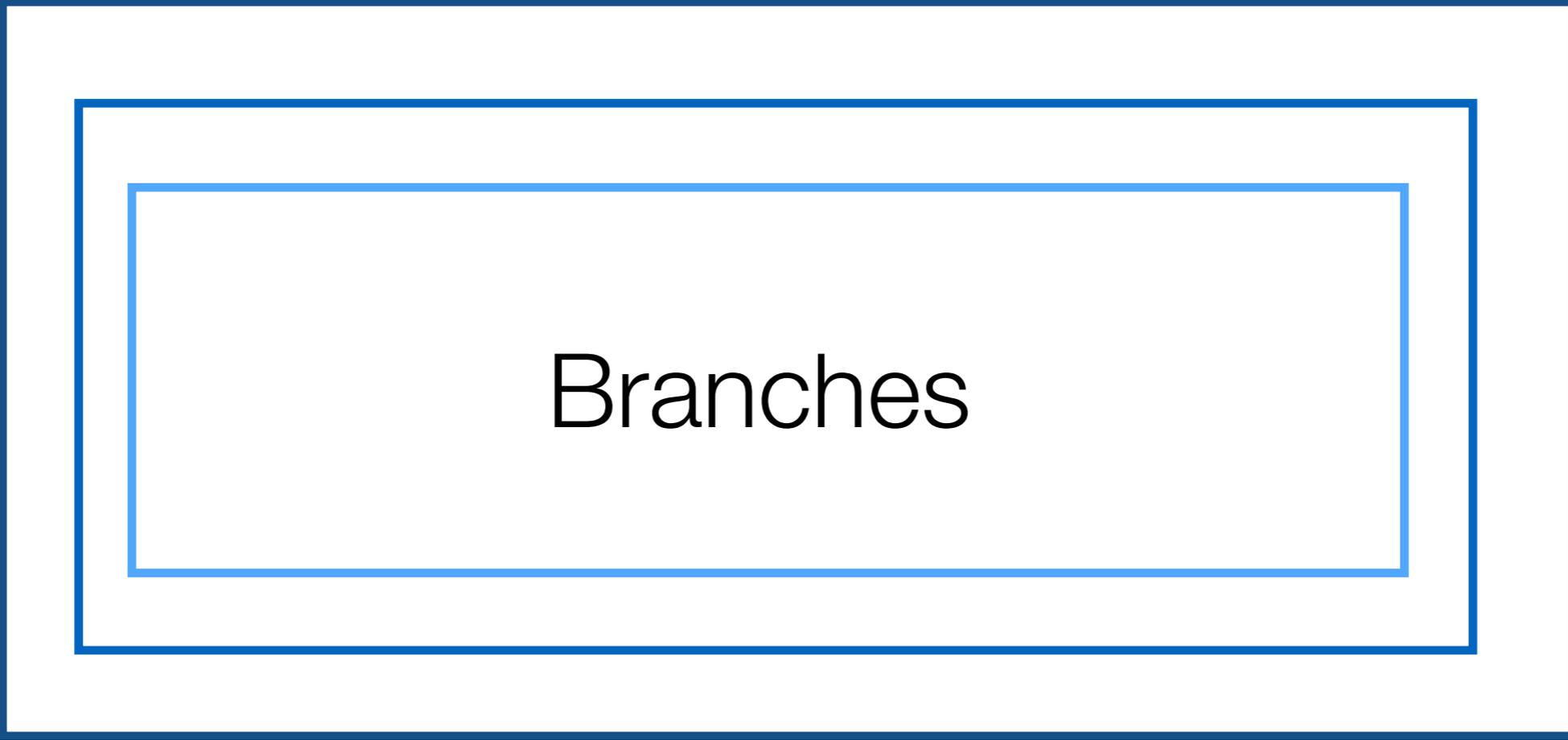
Looping over vectors

```
words <- c("farewell", "cruel", "world")
for (thisWord in words) {
  nLetters <- nchar(thisWord)
  blockWord <- toupper(thisWord)
  cat(blockWord, "has", nLetters, "letters\n")
}
```

```
# FAREWELL has 8 letters
```

```
# CRUEL has 5 letters
```

```
# WORLD has 5 letters
```



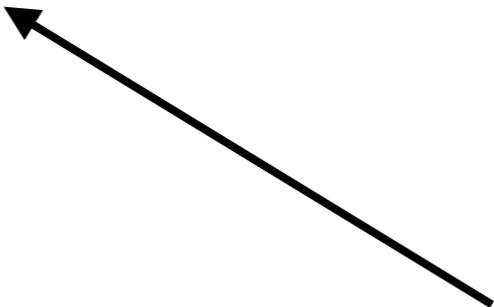
Branches

Branches

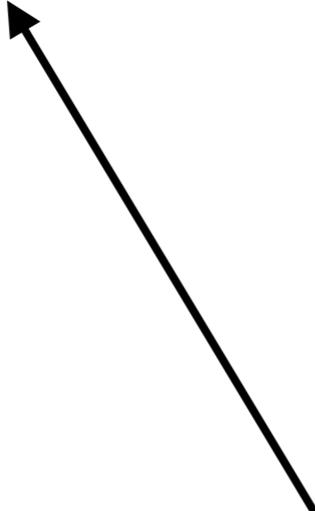
These let you evaluate conditional statements and do different things depending on the outcome

```
If ( CONDITION ) {  
    STATEMENT1  
    STATEMENT2  
    ETC  
}
```

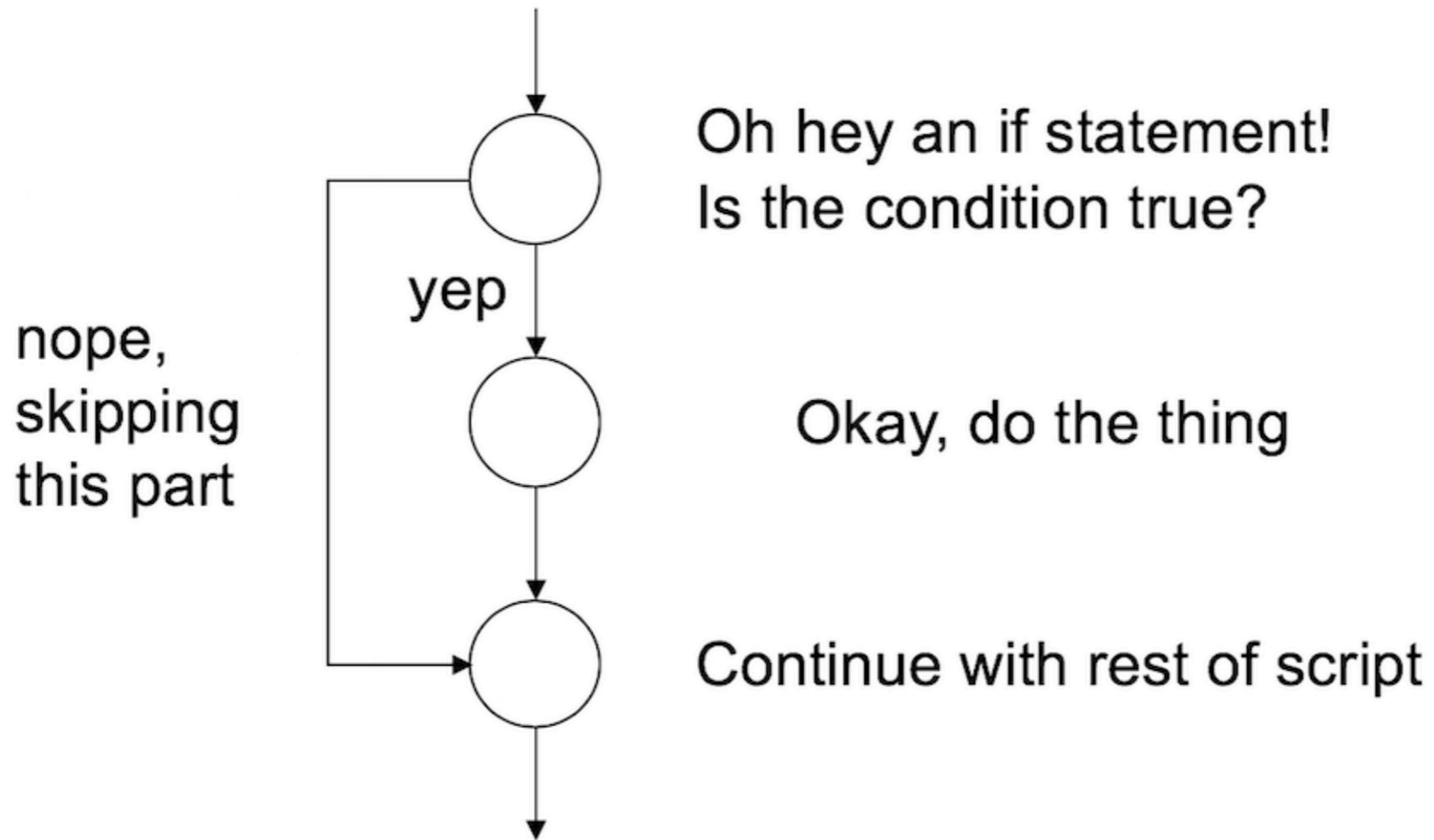
Needs to be a logical
(TRUE or FALSE)



Does all of these
things only if the
condition is TRUE

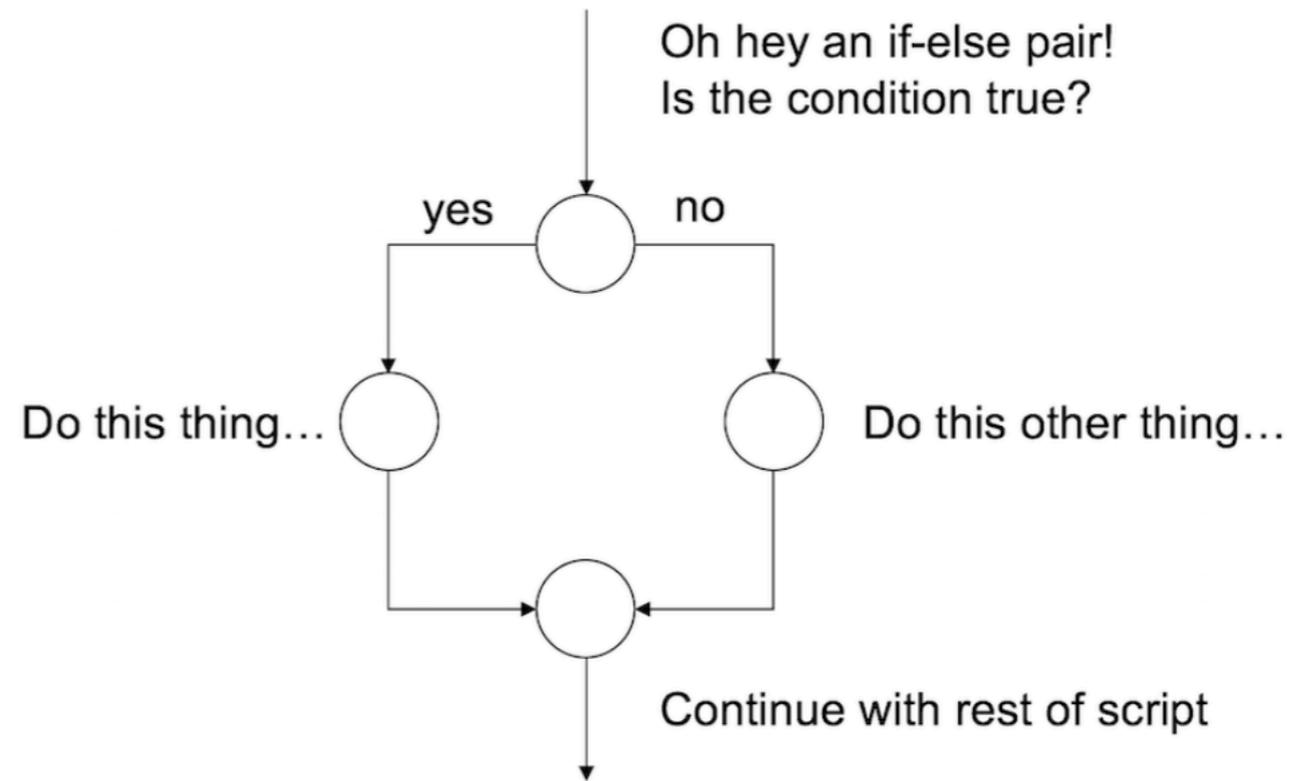


Branches



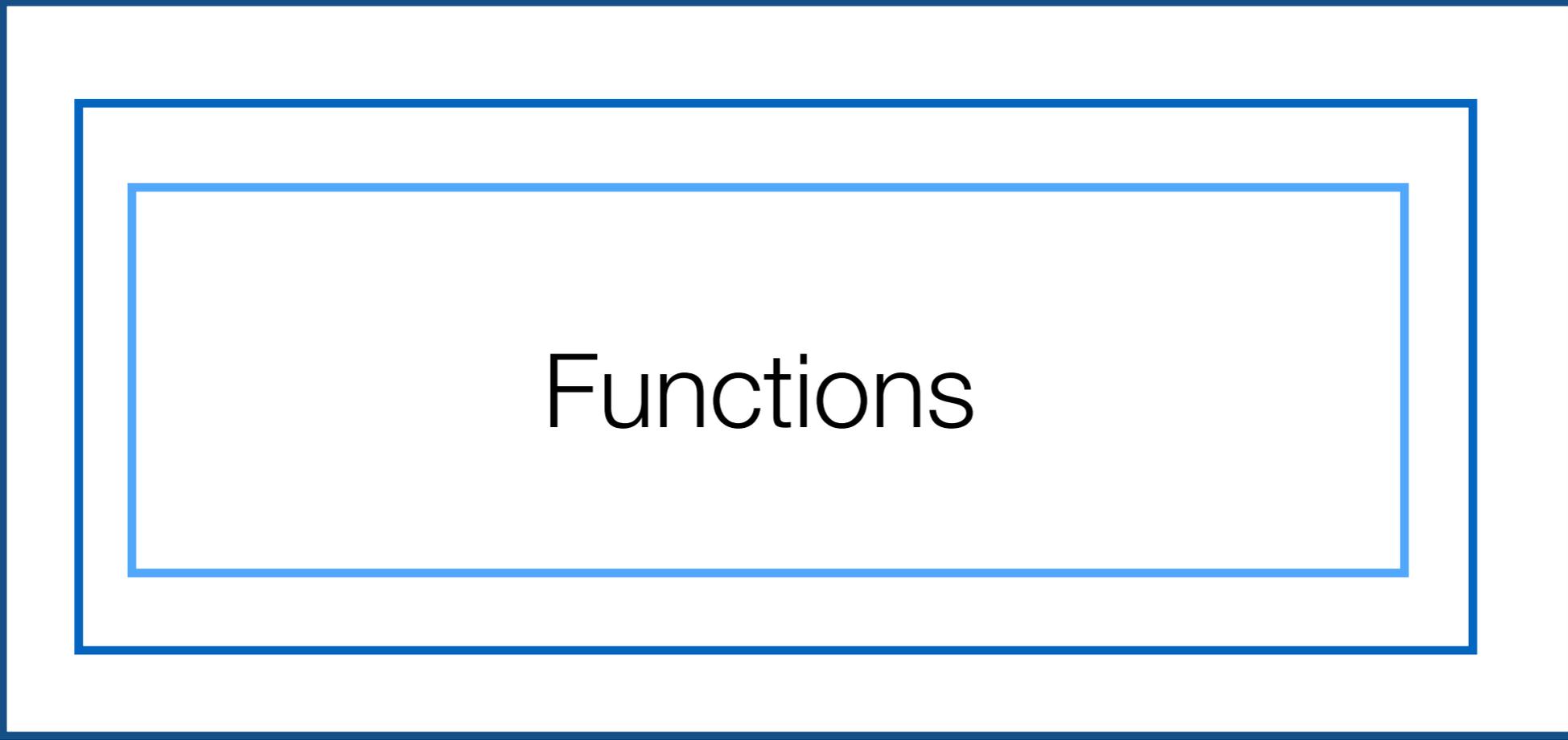
If-Else

```
if ( CONDITION ) {  
    STATEMENT1  
    STATEMENT2  
    ETC  
} else {  
    STATEMENT3  
    STATEMENT4  
}
```



Example

```
if (today=="Saturday") {  
    print("Yay! Weekend!")  
} else if (today=="Sunday") {  
    print("Uh oh, Monday is coming")  
} else {  
    print("I need coffee.")  
}
```



Functions

Functions

You can actually create your *own* functions with arguments. Whenever it is called R will execute the statements within it. Creating a function means R creates a temporary environment with it while it's in practice, and only “keeps” the value in the `return()` statement.

```
FNAME <- function (ARG1, ARG2, ARG3, ETC) {  
  STATEMENT1  
  STATEMENT2  
  STATEMENT3  
  ETC  
  return (VALUE)  
}
```

Functions

Here's an example of a function that will square any number.

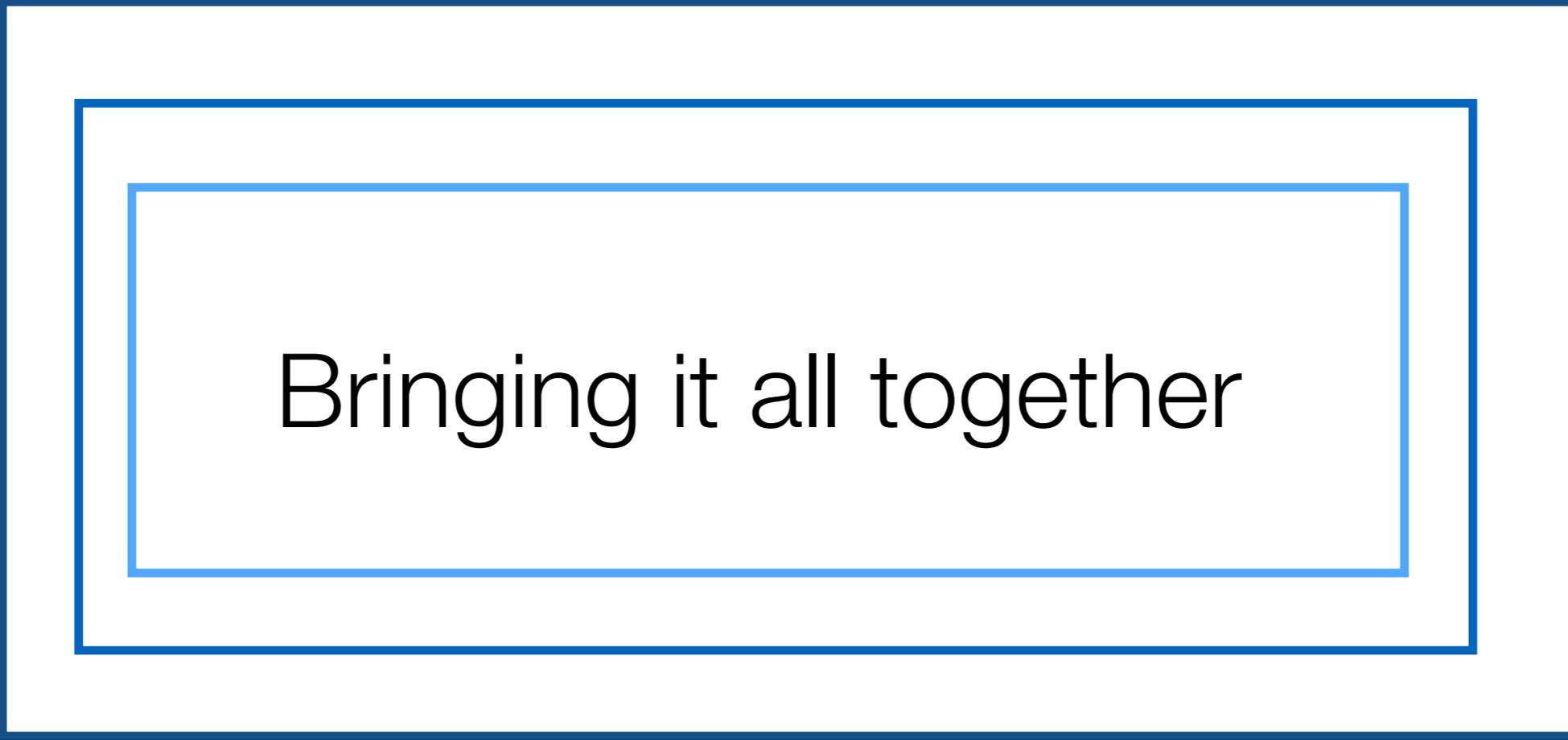
```
square <- function(x) {  
  y <- x*x  
  return(y)  
}
```

```
> square(4)  
# 16
```

Functions

The ... argument lets the user enter as many arguments as they would like, as in the example below.

```
doubleMax <- function(...) {  
  maxVal <- max(...)  
  out <- 2*maxVal  
  return(out)  
}
```



Bringing it all together

What is all this about?????

Suppose we present a compound stimulus AB, which consists of two things, a tone (A) and a light (B). This compound is presented together with a shock. In associative learning studies, this kind of trial is denoted AB+ to indicate that the outcome (US) was present at the same time as the two stimuli that comprise the CS. According to the Rescorla-Wagner model, the rule for updating the associative strengths v_A and v_B between the originally neutral stimuli and the shock is given by:

$$\begin{aligned}v_A &\leftarrow v_A + \alpha_A \beta_U (\lambda_U - v_{AB}) \\v_B &\leftarrow v_B + \alpha_B \beta_U (\lambda_U - v_{AB})\end{aligned}$$

where the associative value v_{AB} of the compound stimulus AB is just the sum of the values of the two items individually. This is expressed as:

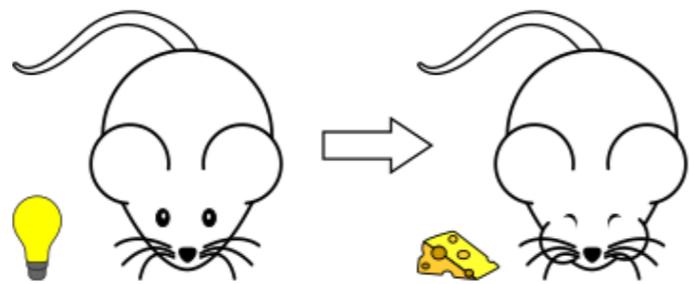
$$v_{AB} = v_A + v_B$$

To understand this rule, note that:

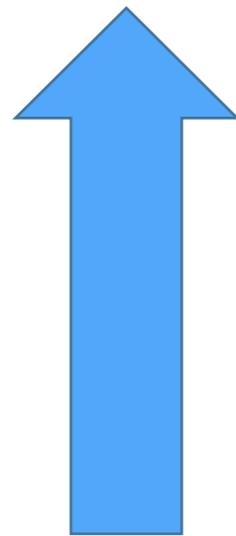
- λ_U is a variable that represents the “reward value” (or “punishment value”) of the US itself, and as such represents the maximum possible association strength for the CS.
- β_U is a learning rate linked to the US (e.g. how quickly do I learn about shocks?)
- α_A is a learning rate linked to the CS (e.g. how quickly do I learn about tones?)
- α_B is also a learning rate linked to the CS (e.g. how quickly do I learn about lights?)



Associative learning



In the simplest design (forward conditioning) the CS is presented slightly before the US, so that it can serve as a signal that reward is coming

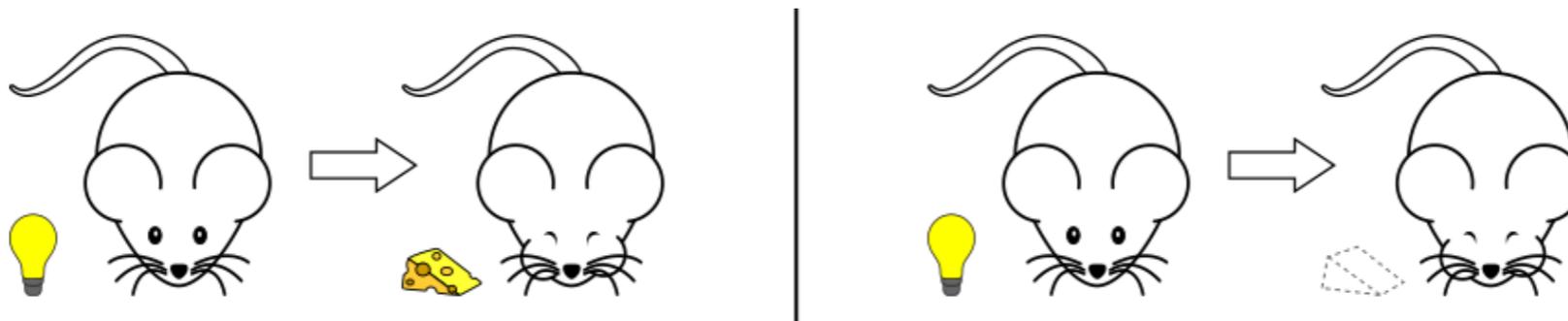


Unconditioned stimulus (US) – something inherently rewarding

Conditioned stimulus (CS) – something initially neutral

Associative learning

(well, Pavlovian anyway)



After some number of presentations, the learner starts to respond to the CS in the same way they would respond to the US

They have a learned association between the CS and the US

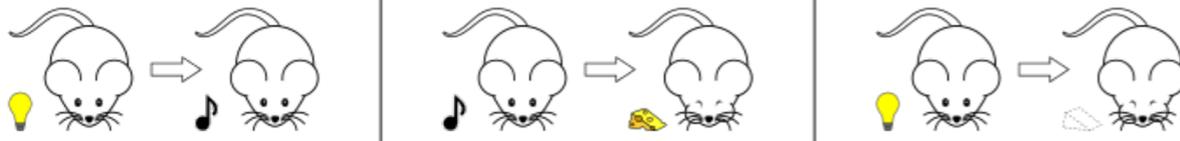
FORWARD CONDITIONING



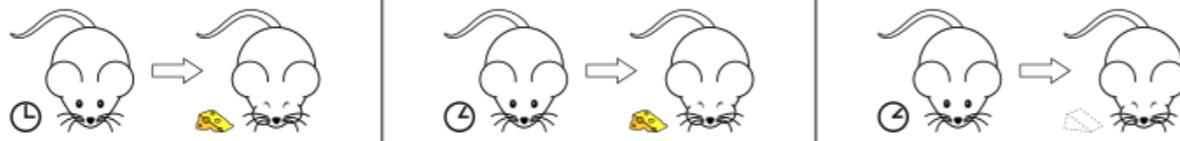
SIMULTANEOUS CONDITIONING



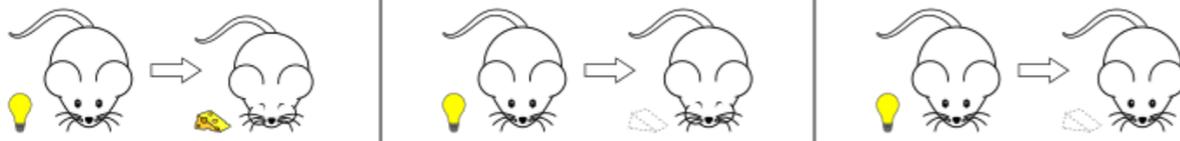
SECOND ORDER CONDITIONING



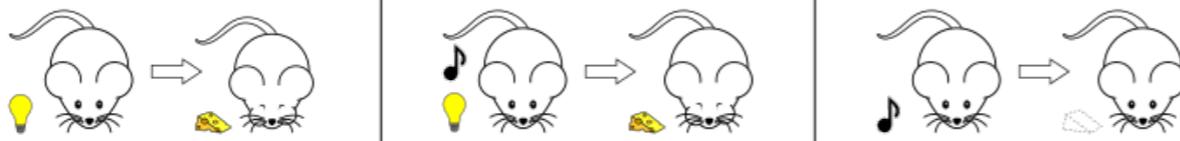
TEMPORAL CONDITIONING



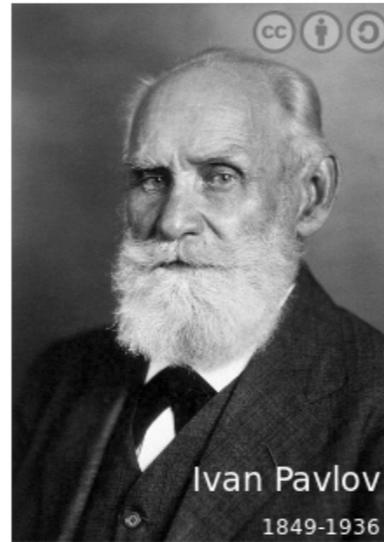
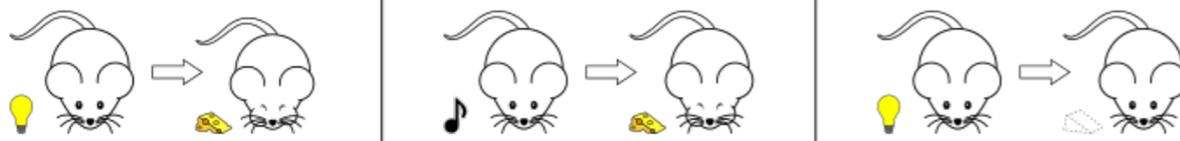
EXTINCTION



BLOCKING



INHIBITION

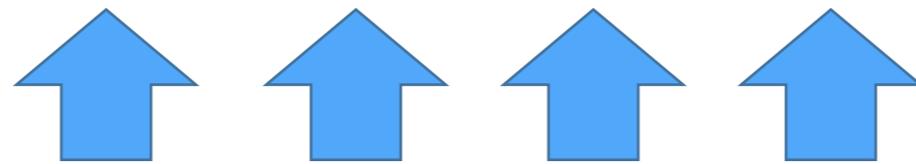


There are many variations on this!

(Long list of empirical effects to account for)

The Rescorla-Wagner model

$$V_x \leftarrow V_x + \alpha\beta(\lambda - V_{xy})$$



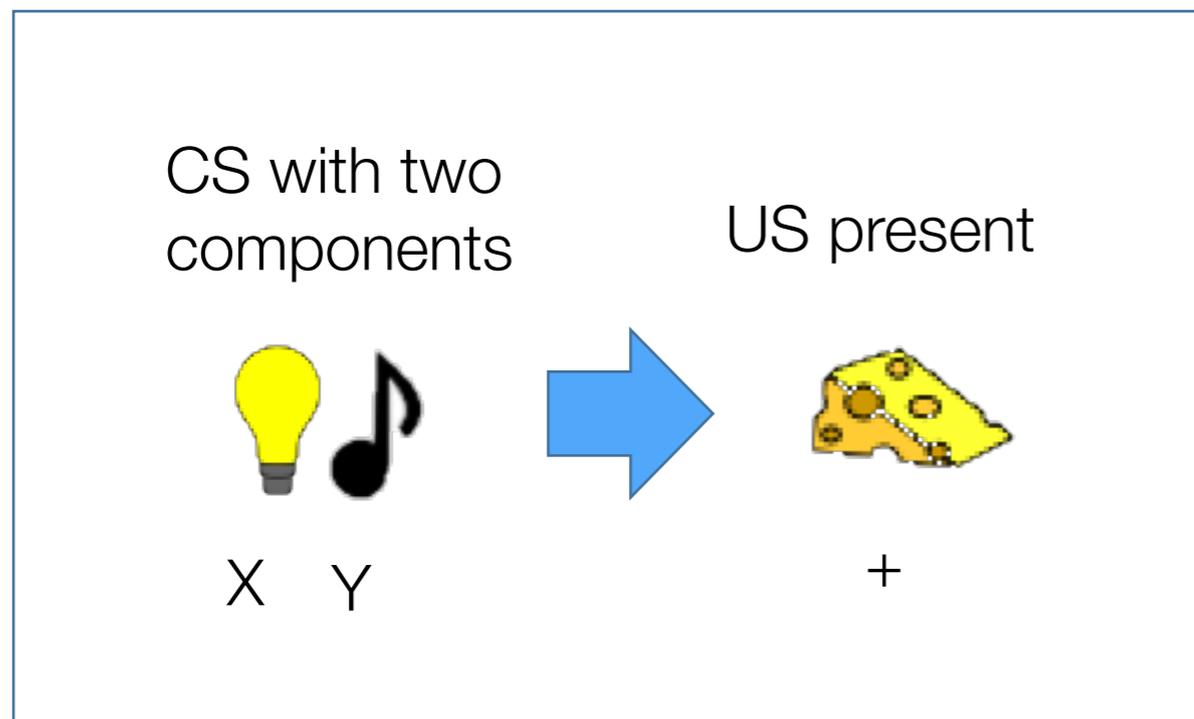
One popular (though flawed & incomplete) account of associative learning is the Rescorla-Wagner model



But what does this strange inscription mean?????

The Rescorla-Wagner model

$$V_x \leftarrow V_x + \alpha\beta(\lambda - V_{xy})$$



Consider a design in which there are two features present (X and Y) and the learner needs to predict an outcome that might be present (+) or absent (-)

An XY+ trial

The Rescorla-Wagner model

$$V_x \leftarrow V_x + \alpha\beta(\lambda - V_{xy})$$

The old strength of association for stimulus X 

The new strength of association for stimulus X after seeing XY+



The difference between the old and the new. By convention “differences” are denoted “delta”, so we call this “delta-V”, ΔV

The Rescorla-Wagner model

$$\Delta V_x = \alpha\beta(\lambda - V_{xy})$$

This delta-V describes “how much we learn about X from the current trial/event”

The “alpha” and “beta” terms here are parameters describing learning rates.

- alpha depends on the CS
- beta depends on the US



The Rescorla-Wagner model

$$\Delta V_x = \alpha \beta (\lambda - V_{xy})$$



This difference term here is called the “reward prediction error”

The Rescorla-Wagner model

$$\Delta V_x = \alpha\beta(\lambda - V_{xy})$$

lambda is represents the “intrinsic” value of the outcome (unconditioned stimulus), sometimes referred to as the “reward”, r

The Rescorla-Wagner model

$$\Delta V_x = \alpha\beta(\lambda - V_{xy})$$



V_{xy} is the “predicted reward”: the amount of reward/punishment that the learner expects to receive upon seeing the compound stimulus XY

In the Rescorla-Wagner model, expectations are additive, which means that:

$$V_{xy} = V_x + V_y$$

(But not all learning models assume additivity)

Error driven learning!

$$\Delta V_x = \alpha \beta (\lambda - V_{xy})$$

reward

expected reward

the difference between outcomes and expectations is the prediction error, and it is this error that “drives” learning

how much does the learner change their beliefs?

learning is gradual, and depends on a learning rate

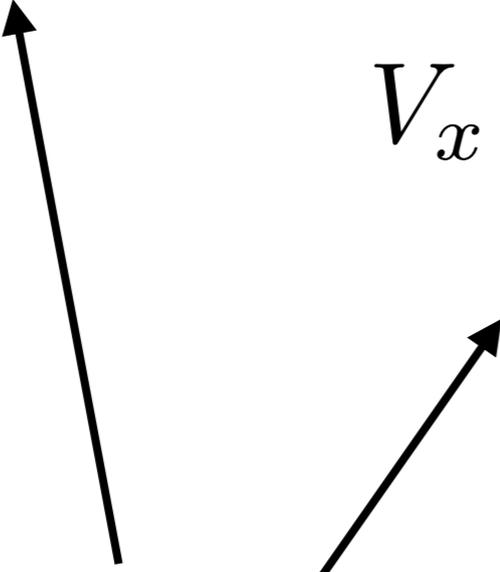


Let's do this!

Step 1: Skeleton

```
updateRW <- function(value, alpha, beta, lambda) {  
}
```

$$V_x \leftarrow V_x + \alpha\beta(\lambda - V_{xy})$$



The design of our R function mirrors the structure of the Rescorla-Wagner model that it implements

Step 1: Skeleton

```
updateRW <- function(value, alpha, beta, lambda) {  
}
```

Vector specifying
associative strength
between US and each
element of the CS

Vector with the
saliency
parameters

Single learning
rate (since only
one US
presented)

Single max
associability

Reminder:

$$V_x \leftarrow V_x + \alpha\beta(\lambda - V_{xy})$$

Step 2: Make a plan

```
updateRW <- function(value, alpha, beta, lambda) {  
  
  # compute the value of the compound stimulus  
  # compute the prediction error  
  # compute the change in strength  
  # update the association value  
  # return the new value  
  
}
```

Reminder:

$$V_x \leftarrow V_x + \alpha\beta(\lambda - V_{xy})$$

Step 3: Put in the details

```
updateRW <- function(value, alpha, beta, lambda) {  
  # compute the value of the compound stimulus  
  valueCompound <- sum(value)  
  
  # compute the prediction error  
  predictionError <- lambda - valueCompound  
  
  # compute the change in strength  
  valueChange <- alpha * beta * predictionError  
  
  # update the association value  
  value <- value + valueChange  
  
  # return the new value  
  return(value)  
}
```

Reminder:

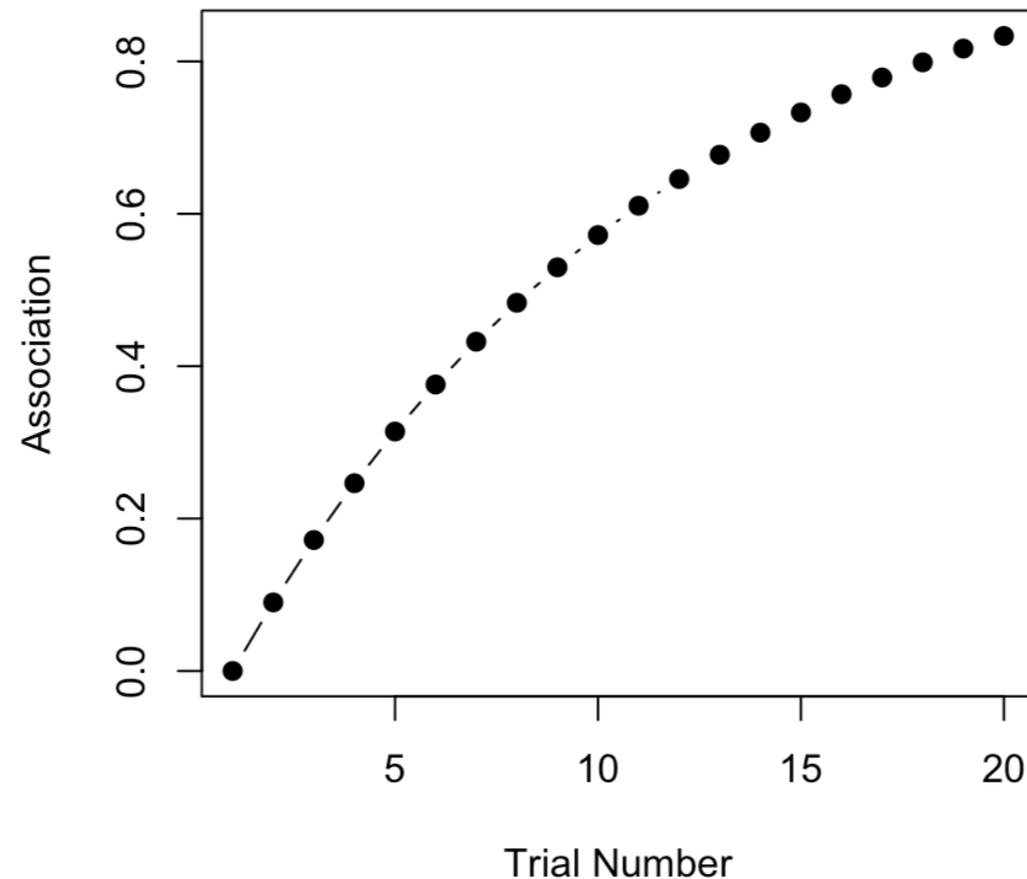
$$V_x \leftarrow V_x + \alpha\beta(\lambda - V_{xy})$$

Step 4: Model predictions

1. Conditioning
2. Extinction
3. Blocking

Conditioning

```
nTrials <- 20  
strength <- numeric(nTrials)  
  
for (trial in 2:nTrials) {  
  strength[trial] <- updateRW(strength[trial-1])  
}
```



Extinction

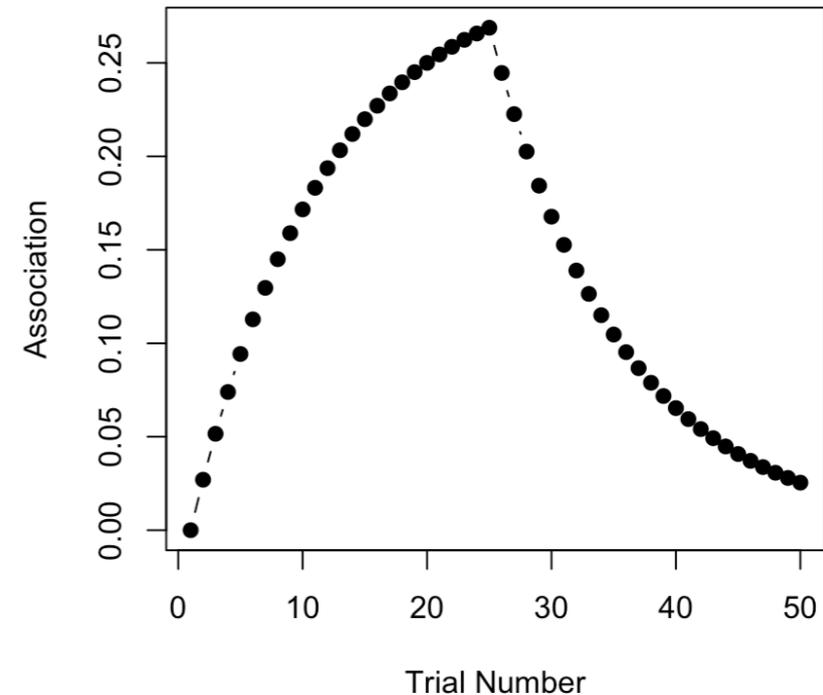
```
nTrials <- 50
strength <- numeric(nTrials)
lambda <- 0.3

for (trial in 2:nTrials) {

  # remove the shock after trial 25
  if (trial>25) {
    lambda <- 0
  }

  # update associative strength on each trial
  strength[trial] <- updateRW(value=strength[trial-1],
                              lambda=lambda)

}
```



Blocking

```
# total number of trials across  
# both phases of the task  
n_trials <- 50  
  
# vectors of zeros  
strength_A <- rep(0,n_trials)  
strength_B <- rep(0,n_trials)
```

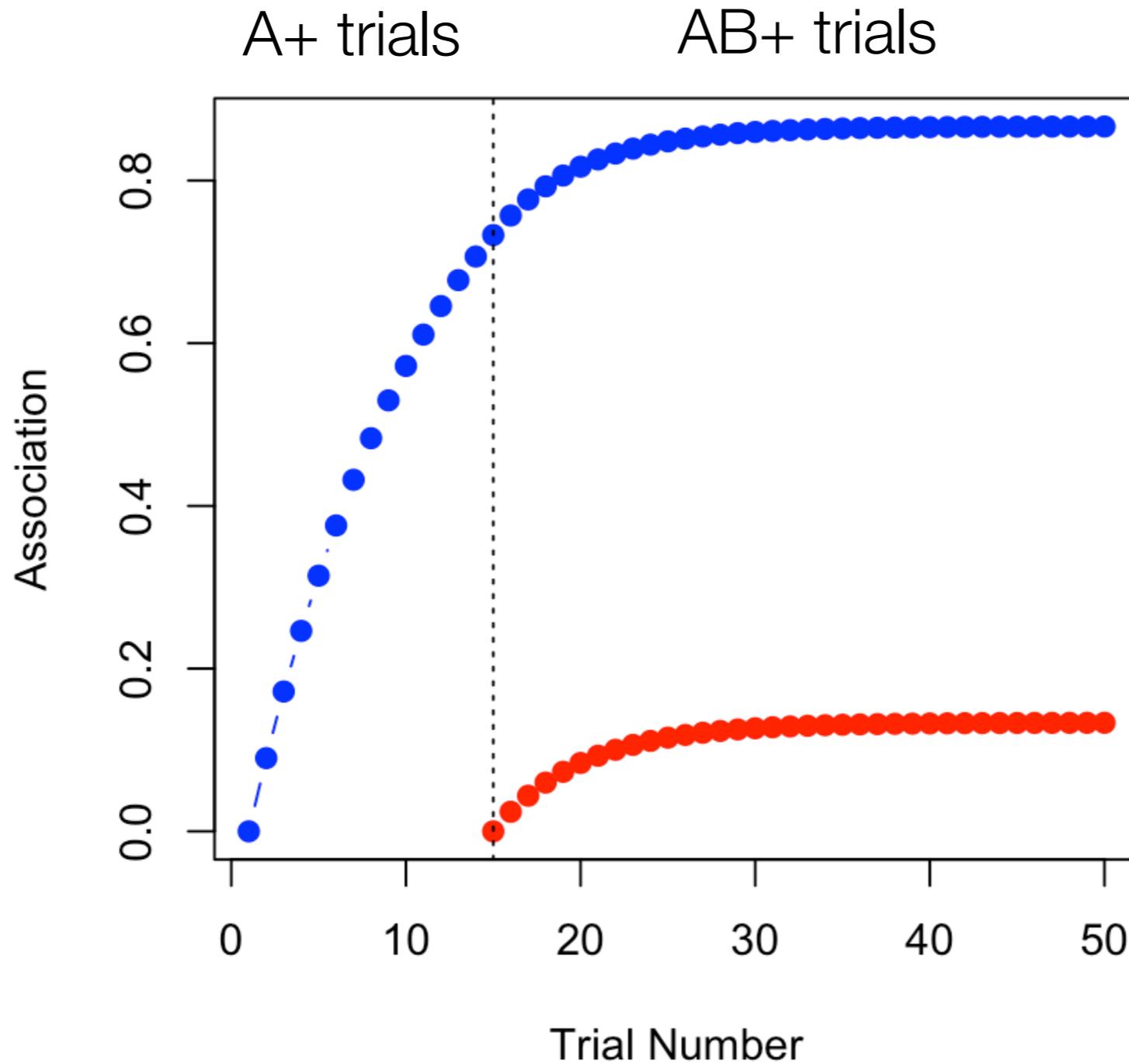
Blocking

```
# total number of trials across  
# both phases of the task  
n_trials <- 50  
  
# vectors of zeros  
strength_A <- rep(0,n_trials)  
strength_B <- rep(0,n_trials)  
  
# learning rate for the CS at the  
# start of the experiment is .3 for  
# A and 0 for B (b/c it's absent)  
alpha <- c(.3, 0)
```

Blocking

```
for(trial in 2:n_trials) {  
  
  # after trial 15, both stimuli are present  
  if(trial > 15) alpha <- c(.3, .3)  
  
  # vector of current associative strengths  
  v_old <- c(strength_A[trial-1], strength_B[trial-1])  
  
  # vector of new associative strengths  
  v_new <- update_RW(  
    value = v_old,  
    alpha = alpha  
  )  
  
  # record the new strengths  
  strength_A[trial] <- v_new[1]  
  strength_B[trial] <- v_new[2]  
}
```

Blocking



Strong association to A is formed early and maintained

There is learning to B, but greatly reduced and it asymptotes at a low level

Intro to R cheat sheet

1 Saving and importing

- Save as .RData, using menu or `save.image()`
- Can load .csv, using menu or `read.csv()`

12 Scripts let you run and save series of commands

```
1 # this is my first script
2 # it's just for DRIP class
3 #
4 # author: Amy Perfors
5
6 # define some variables
7 age <- 19
8 box <- "cat"
9
10 # print something
11 print( box )
12 print( age )
```

13

`help(functionName)`
e.g. `help(print)`

Files Plots Packages Help Viewer

R: Print Values Find in Topic

print {base}

Print Values

Description

print prints its argument and returns it in a function which means that new printing method

Usage

```
print(x, ...)
```

Arguments

x	an object used to select
...	further arguments passed
quote	logical, indicating whether quotes.
max.levels	integer, indicating how many extra "Levels" lines will be printed.
width	only used when max.levels is not null.