

Introduction to R

BIOE 498/598 PJ

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Everything is a vector

Vectors are built with the `c()` function (for “combine”)

```
a <- c(10, 20, 30, 40)
a
```

```
## [1] 10 20 30 40
```

Vectors are 1-indexed (like Matlab, Fortran, or Julia)

```
a[1]
```

```
## [1] 10
```

```
a[2:4]
```

```
## [1] 20 30 40
```

Some R artifacts

For historical reasons, R assigns objects to variables using `<-` (said “gets”)

```
a <- c(10, 20, 30, 40)
```

However, the equals sign also works

```
a = c(11, 22, 33, 44)
```

Also, periods are allowed in variables names

```
a.var.name <- 7  
is.na(a.var.name)
```

```
## [1] FALSE
```

It's best to avoid this and use underscores (`_`) instead

Vector Types

Numeric

```
r <- rnorm(4)
r
```

```
## [1] 1.3493629 -0.2646543 1.9097851 0.4067838
```

Logical (TRUE and FALSE can be abbreviated T and F)

```
r < 0.0
```

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

Character

```
c <- c("this", "is", "a", "character")
```

Notes about characters

Everything is a vector of “strings”; there are no individual “letters”

```
s <- "a string"  
length(s)
```

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

```
s[1]
```

```
## [1] "a string"
```

```
s[2]
```

```
## [1] NA
```

Notes about characters (continued)

```
nchar("a string")
```

```
## [1] 8
```

```
c <- c("this", "is", "a", "character")  
length(c)
```

```
## [1] 4
```

```
nchar(c)
```

```
## [1] 4 2 1 9
```

Everything in a vector is the same type

R will convert objects into the same type

```
c(1, 2, "three")
```

```
## [1] "1"      "2"      "three"
```

For mixed types, use a list

```
list(1, 2, "three")
```

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

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

```
##
```

```
## [[2]]
```

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

```
##
```

```
## [[3]]
```

```
## [1] "three"
```

Factors are special vectors

Factors look like regular vectors

```
x1 <- as.factor(c("wt", "ko", "wt"))
```

```
x1
```

```
## [1] wt ko wt
```

```
## Levels: ko wt
```


Factors are special vectors

Factors look like regular vectors

```
x1 <- as.factor(c("wt", "ko", "wt"))  
x1
```

```
## [1] wt ko wt  
## Levels: ko wt
```

But they are stored as integers and a “key” of levels

```
as.integer(x1)
```

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

```
levels(x1)
```

```
## [1] "ko" "wt"
```

Functions

Most functions are *vectorized* and operate elementwise

```
a <- runif(4)
```

```
a
```

```
## [1] 0.7240759 0.7865879 0.2458798 0.1849558
```

```
10*a + 1
```

```
## [1] 8.240759 8.865879 3.458798 2.849558
```

```
sqrt(a) + cos(a)
```

```
## [1] 1.600038 1.593163 1.465786 1.413009
```

You can always ask for help: `?sqrt`.

Data Frames

Data frames hold tables of data.

```
data <- read.csv("MonkeyThrow.csv")  
data
```

```
##   run  hand hat boots distance  
## 1    5 left yes   yes     4.5  
## 2    6 right yes   yes     6.0  
## 3    2 left  no    yes     7.0  
## 4    7 right no    yes     9.5  
## 5    1 left yes   no     5.0  
## 6    8 right yes   no     6.5  
## 7    4 left  no    no     4.0  
## 8    3 right no    no     7.5
```

Size of Data Frames

```
nrow(data)
```

```
## [1] 8
```

```
ncol(data)
```

```
## [1] 5
```

The length of a data frame is the number of *columns*.

```
length(data)
```

```
## [1] 5
```

What are data frames?

```
head(data, n=5) # just the first 5 rows
```

```
##   run  hand hat boots distance
## 1   5  left yes  yes     4.5
## 2   6 right yes  yes     6.0
## 3   2  left no   yes     7.0
## 4   7 right no   yes     9.5
## 5   1  left yes  no     5.0
```

- ▶ Each column in a data frame is a vector (and must be the same type).
- ▶ Each row contains data of different types

What types (“classes”) of data are in our data frame?

```
lapply(data, class) # applies `class` to each column
```

```
## $run
```

```
## [1] "integer"
```

```
##
```

```
## $hand
```

```
## [1] "factor"
```

```
##
```

```
## $hat
```

```
## [1] "factor"
```

```
##
```

```
## $boots
```

```
## [1] "factor"
```

```
##
```

```
## $distance
```

```
## [1] "numeric"
```

Working with columns

```
data$hat
```

```
## [1] yes yes no no yes yes no no
```

```
## Levels: no yes
```

```
log10(data$distance)
```

```
## [1] 0.6532125 0.7781513 0.8450980 0.9777236 0.6989700 0.7781513 0.8450980 0.9777236
```

```
## [8] 0.8750613
```

Making new columns

```
data$log_dist <- log10(data$distance)
data
```

```
##   run  hand hat boots distance  log_dist
## 1   5 left yes  yes    4.5 0.6532125
## 2   6 right yes  yes    6.0 0.7781513
## 3   2 left  no   yes    7.0 0.8450980
## 4   7 right no   yes    9.5 0.9777236
## 5   1 left yes   no    5.0 0.6989700
## 6   8 right yes   no    6.5 0.8129134
## 7   4 left  no   no    4.0 0.6020600
## 8   3 right no    no    7.5 0.8750613
```

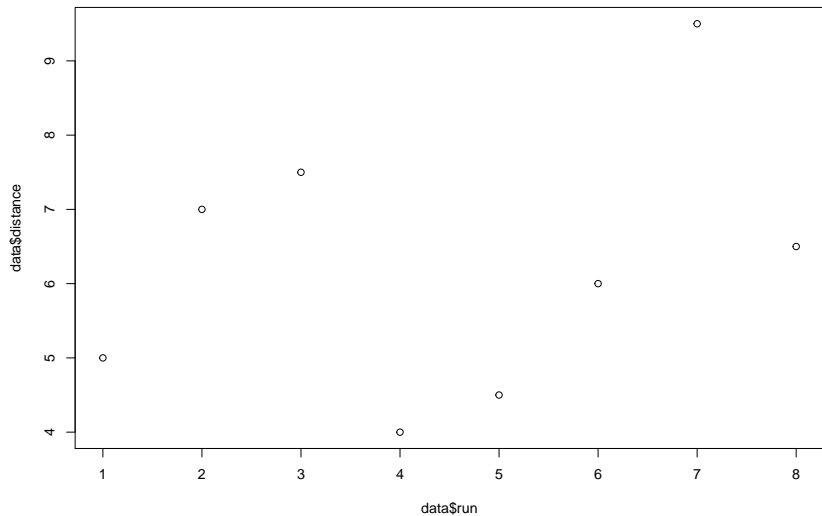

Nevermind, let's delete that column

```
data$log_dist <- NULL  
data
```

```
##   run  hand hat boots distance  
## 1    5 left yes  yes    4.5  
## 2    6 right yes  yes    6.0  
## 3    2 left  no   yes    7.0  
## 4    7 right no   yes    9.5  
## 5    1 left yes   no    5.0  
## 6    8 right yes   no    6.5  
## 7    4 left  no   no    4.0  
## 8    3 right no    no    7.5
```

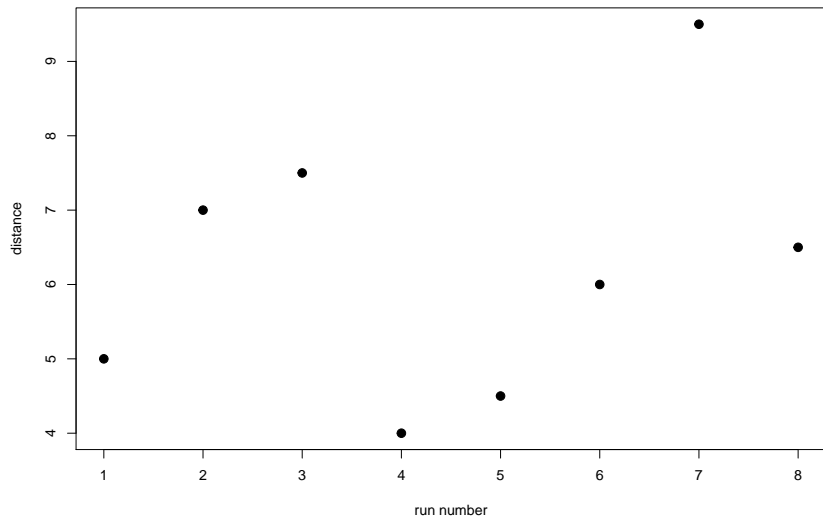
Visualizing data

```
plot(data$run, data$distance)
```



We can do better

```
plot(data$run, data$distance, xlab="run number",  
      ylab="distance", cex=2, pch=20)
```



Libraries

Many of the functions we use are in libraries. We need to load the library first.

```
library("doetools") # include `farplot` function  
farplot(data, response="distance", factors=c("hand", "hat", "boots"))
```

