

# Naming conventions in R

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# Naming rules

# Why do we need names?

Everything in R is an object that can be named

Unless you name an object, it doesn't stay in the R environment

You name an object using the assignment operator `<-`  
I read this as a left arrow

For example

```
spine_data <- readr::read_csv('data/Dataset_spine.csv')
```

Here, we read the CSV file into R, and then save it in the current R session by naming it `spine_data`

# Syntactic names

Following these rules will always give you valid object names without the need for additional trouble.

1. Contains letters, numbers, `.` or `_` characters
2. Starts with a letter or `.` not followed by a number
3. Is not a reserved word ()

So `.2way` is not a syntactically valid name

# Going wild

You can actually use **any** name for an object, as long as you encase it in backticks

The name can then include spaces, special characters, start with a number, etc.

Still, don't use reserved words.

You can do

```
`First variable`
```

```
`2 many bikes`
```

**But should you?!!**

# Naming conventions

There are several prevalent naming conventions in R

- snake case, e.g., `my_variable`
- lower camel case, e.g., `myVariable`
- upper camel case, e.g., `MyVariable`
- period separated, e.g., `my.variable`
- lower case, e.g., `myvariable`

Choose one and stick with it. It will save you some sanity.

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The `janitor` package has a nice function called `clean_names` which will transform the column names in a `data.frame` to follow a particular naming convention, with the default being snake case.

# janitor::clean\_names

convert all column names to `_` case!

(not so awesome column names)



clean\_names(  
your data frame +  
case choice here)

- CHOOSE:
- ✓ snake
  - lower\_camel
  - big\_camel
  - screaming\_snake
  - + more!

WAY MORE  
DEAL WITHABLE  
COLUMN  
NAMES!



# Resources

The following commands, typed in the R console, will get you to the corresponding help files.

- `?make.names`
- `?assign`



# Brackets

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There are three kinds of brackets in R

`[]` are used for extracting elements from arrays, matrices, data frames.

Arrays

Lists

- `x[3]` is the 3rd element of an array `x`
- `d[1,3]` is the element in the 1st row and 3rd column of a matrix/data frame `d`
- `d[2,]` is the entire first **row** of a matrix/data frame `d`
- `d[,4]` is the entire 4th **column** of a matrix/data frame `d`
- `d[, 'gender']` is the column named *gender* in a *data frame d*

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Arrays   Lists

To extract single elements, you use a **double bracket**:

- `lst[[3]]` is the third element of the list `lst`
- `lst[['apple']]` is the element named *apple* in a named list `lst`

To extract a sub-list, you use **single brackets**:

- `lst[1:3]` creates a sub-list with the first 3 elements of `lst`

# Brackets

There are three kinds of brackets in R

`()` are used for specifying arguments to functions

- `mean(x)` gives the mean of an array of numbers `x`
- `summary(d)` gives a summary representation of a data frame `d`
- There are several functions used to make the following visualization

```
ggplot(beaches, aes(x = temperature, y = rainfall)) +  
  geom_point() +  
  geom_smooth() +  
  theme_classic()
```

# Brackets

There are three kinds of brackets in R

`{ }` are used to contain groups of commands/statements

A conditional statement

```
if (age < 18){
  person <- 'Minor'
} else if (age > 65) {
  person <- 'Senior'
} else {
  person <- 'Adult'
}
```

A function definition

```
my_mean <- function(x, na.rm = T){
  if(na.rm){
    x <- x[!is.na(x)]
  }
  s <- sum(x)
  n <- length(x)
  mn <- s / n # There is a built-in function mean, s
  return(mn)
}
```