



# Agenda

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- Finishing up on coercion
- Attributes
- Missing values
- Intro to lists
- Subsetting

# Learning objectives

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- Understand the fundamental difference between lists and atomic vectors
- Understand how atomic vectors are coerced, implicitly or explicitly
- Understand various ways to subset vectors, and how subsetting differs for lists
- Understand what an attribute is, and how to set and modify attributes

# Pop quiz

---

Without actually running the code, predict which type each of the following will coerce to.

```
c(TRUE, 1L, 0L, "False")  
c(1L, FALSE)  
c(7L, 6.23, "eight")  
c(1.25, TRUE, 4L)
```

01:00

# Answers

---

```
typeof(c(TRUE, 1L, 0L, "False"))
```

```
## [1] "character"
```

```
typeof(c(1L, FALSE))
```

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

```
typeof(c(7L, 6.23, "eight"))
```

```
## [1] "character"
```

```
typeof(c(1.25, TRUE, 4L))
```

```
## [1] "double"
```

# Challenge

---

Work with a partner

One of you share your screen:

- Create four atomic vectors, one for each of the fundamental types
- Combine two or more of the vectors. Predict the implicit coercion of each.
- Apply explicit coercions, and predict the output for each.

(basically quiz each other)



08:00

# Attributes

---

# Attributes

---

- What are attributes?
  - metadata... what's metadata?
  - Data about the data



# Other data types

---

Atomic vectors by themselves make up only a small fraction of the total number of data types in R

What are some other data types?

- Data frames
- Matrices & arrays
- Factors
- Dates

Remember, atomic vectors are the atoms of R. Many other data structures are built from atomic vectors.

- We use attributes to create other data types from atomic vectors

# Attributes

---

## Common

- Names
- Dimensions

## Less common

- Arbitrary metadata

# Examples

---

Please follow along!

- See **all** attributes associated with a give object with **attributes**

```
library(palmerpenguins)
attributes(penguins[1:50, ]) # limiting rows just for slides
```

```
## $names
## [1] "species"          "island"            "bill_length_mm"    "bill_de
## [5] "flipper_length_mm" "body_mass_g"       "sex"               "year"
##
## $row.names
## [1] 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 2
## [29] 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50
##
## $class
## [1] "tbl_df"          "tbl"             "data.frame"
```

```
head(penguins)
```

```
## # A tibble: 6 x 8
##   species island    bill_length_mm bill_depth_mm flipper_length_mm body_
##   <fct>    <fct>          <dbl>         <dbl>          <dbl>      <int>
## 1 Big one Torgersen      39.1           18.7           181
## 2 Big one Torgersen      39.5           17.400          186
## 3 Big one Torgersen     40.300           18           195
## 4 Big one Torgersen      NA             NA             NA
## 5 Big one Torgersen      36.7           19.3           193
## 6 Big one Torgersen     39.300           20.6           190
## # ... with 1 more variable: year <int>
```

# Get specific attribute

---

- Access just a single attribute by naming it within `attr`

```
attr(penguins, "class")
```

```
## [1] "tbl_df"      "tbl"        "data.frame"
```

```
attr(penguins, "names")
```

```
## [1] "species"      "island"      "bill_length_mm" "bill_de
## [5] "flipper_length_mm" "body_mass_g" "sex"           "year"
```

Note – this is not generally how you would pull these attributes. Rather, you would use `class()` and `names()`.

# Be specific

---

- Note in the prior slides, I'm asking for attributes on the entire data frame.
- Is that what I want?... maybe. But the individual vectors may have attributes as well

```
attributes(penguins$species)
```

```
## $levels  
## [1] "Big one"      "Little one" "Funny one"  
##  
## $class  
## [1] "factor"
```

```
attributes(penguins$bill_length_mm)
```

```
## NULL
```

# Set attributes

---

- Just redefine them within `attr`

```
attr(penguins$species, "levels") <- c("Big one",  
                                       "Little one",  
                                       "Funny one")
```

```
head(penguins)
```

```
## # A tibble: 6 x 8  
##   species island    bill_length_mm bill_depth_mm flipper_length_mm body_  
##   <fct>    <fct>          <dbl>          <dbl>          <int>  
## 1 Big one Torgersen      39.1            18.7            181  
## 2 Big one Torgersen      39.5            17.400           186  
## 3 Big one Torgersen      40.300           18            195  
## 4 Big one Torgersen      NA              NA              NA  
## 5 Big one Torgersen      36.7            19.3            193  
## 6 Big one Torgersen      39.300           20.6            190  
## # ... with 1 more variable: year <int>
```

Note – you would generally not define levels this way either, but it is a general method for modifying attributes.

# Dimensions

---

- Let's create a matrix (please do it with me)

```
m <- matrix(1:6, ncol = 2)
m
```

```
##      [,1] [,2]
## [1,]    1    4
## [2,]    2    5
## [3,]    3    6
```

- Notice how the matrix fills
- Check out the attributes

```
attributes(m)
```

```
## $dim
## [1] 3 2
```



# Modify the attributes

---

- Let's change it to a 2 x 3 matrix, instead of 3 x 2 (you try first)

```
attr(m, "dim") <- c(2, 3)
m
```

```
##      [,1] [,2] [,3]
## [1,]    1    3    5
## [2,]    2    4    6
```

- is this the result you expected?

# Alternative creation

---

- Create an atomic vector, assign a dimension attribute

```
v <- 1:6  
v
```

```
## [1] 1 2 3 4 5 6
```

```
attr(v, "dim") <- c(3, 2)  
v
```

```
##      [,1] [,2]  
## [1,]    1    4  
## [2,]    2    5  
## [3,]    3    6
```

# Aside

---

- What if we wanted it to fill by row?

```
matrix(6:13,  
       ncol = 2,  
       byrow = TRUE)
```

```
##      [,1] [,2]  
## [1,]    6    7  
## [2,]    8    9  
## [3,]   10   11  
## [4,]   12   13
```

```
vect <- 6:13  
dim(vect) <- c(2, 4)  
vect
```

```
##      [,1] [,2] [,3] [,4]  
## [1,]    6    8   10   12  
## [2,]    7    9   11   13
```

```
t(vect)
```

```
##      [,1] [,2]  
## [1,]    6    7  
## [2,]    8    9  
## [3,]   10   11  
## [4,]   12   13
```

# Names

---

- The following (this slide and the next) are equivalent

```
attr(v, "dimnames") <- list(c("the first", "second", "III"),  
                             c("index", "value"))
```

v

##		index	value
##	the first	1	4
##	second	2	5
##	III	3	6

# Names

---

```
v2 <- 1:6
attr(v2, "dim") <- c(3, 2)
rownames(v2) <- c("the first", "second", "III")
colnames(v2) <- c("index", "value")
v2
```

```
##           index value
## the first      1      4
## second        2      5
## III           3      6
```

# Arbitrary metadata

---

- I don't use this often (wouldn't recommend you do either)

```
attr(v, "matrix_mean") <- mean(v)
v
```

```
##           index value
## the first      1      4
## second        2      5
## III           3      6
## attr(,"matrix_mean")
## [1] 3.5
```

```
attr(v, "matrix_mean")
```

```
## [1] 3.5
```

- Note that *anything* can be stored as an attribute (including matrices or data frames, etc.)

# Real example

---

Fit a multilevel model and pull the variance–covariance matrix

```
m <- lme4::lmer(Reaction ~ 1 + Days + (1 + Days|Subject),  
               data = lme4::sleepstudy)
```

```
lme4::VarCorr(m)$Subject
```

```
##              (Intercept)           Days  
## (Intercept)  612.100158    9.604409  
## Days        9.604409   35.071714  
## attr(,"stddev")  
## (Intercept)           Days  
##   24.740658    5.922138  
## attr(,"correlation")  
##              (Intercept)           Days  
## (Intercept)  1.00000000  0.06555124  
## Days        0.06555124  1.00000000
```

# Matrices vs Data frames

---

Usually we want to work with data frames because they represent our data better.

Sometimes a matrix is more efficient because you can operate on the **entire** matrix at once.

```
set.seed(42)
m <- matrix(rnorm(100, 200, 10), ncol = 10)
m
```

```
##           [,1]      [,2]      [,3]      [,4]      [,5]      [,6]      [,7]
## [1,] 213.7096 213.0487 196.9336 204.5545 202.0600 203.2193 196.3277 189.
## [2,] 194.3530 222.8665 182.1869 207.0484 196.3894 192.1616 201.8523 199.
## [3,] 203.6313 186.1114 198.2808 210.3510 207.5816 215.7573 205.8182 206.
## [4,] 206.3286 197.2121 212.1467 193.9107 192.7330 206.4290 213.9974 190.
## [5,] 204.0427 198.6668 218.9519 205.0496 186.3172 200.8976 192.7271 194.
## [6,] 198.9388 206.3595 195.6953 182.8299 204.3282 202.7655 213.0254 205.
## [7,] 215.1152 197.1575 197.4273 192.1554 191.8861 206.7929 203.3585 207.
## [8,] 199.0534 173.4354 182.3684 191.4909 214.4410 200.8983 210.3851 204.
## [9,] 220.1842 175.5953 204.6010 175.8579 195.6855 170.0691 209.2073 191.
## [10,] 199.3729 213.2011 193.6001 200.3612 206.5565 202.8488 207.2088 189.
##           [,10]
```



```
sum(m)
```

```
## [1] 20032.51
```

```
mean(m)
```

```
## [1] 200.3251
```

```
rowSums(m)
```

```
## [1] 2048.470 1993.774 2041.155 2025.924 1978.173 2007.265 1998.086 1960.173 1978.173 1978.173  
## [10] 2026.901
```

```
colSums(m)
```

```
## [1] 2054.730 1983.654 1982.192 1963.610 1997.978 2001.839 2053.908 1978.173 1978.173 1978.173  
## [10] 1991.281
```

```
# standardize the matrix  
z <- (m - mean(m)) / sd(m)
```

Z

##		[,1]	[,2]	[,3]	[,4]	[,5]	[,6]
##	[1,]	1.28528802	1.2218239	-0.3256841	0.40613865	0.1665940	0.277916
##	[2,]	-0.57349498	2.1646089	-1.7417882	0.64562157	-0.3779416	-0.783932
##	[3,]	0.31748345	-1.3649263	-0.1963133	0.96277141	0.6968297	1.481924
##	[4,]	0.57650528	-0.2989403	1.1352110	-0.61596668	-0.7290676	0.586143
##	[5,]	0.35698951	-0.1592501	1.7887033	0.45367758	-1.3451640	0.054972
##	[6,]	-0.13313334	0.5794704	-0.4445968	-1.68004206	0.3844054	0.234344
##	[7,]	1.42026916	-0.3041875	-0.2782756	-0.78452812	-0.8103926	0.621087
##	[8,]	-0.12212321	-2.5821792	-1.7243635	-0.84833774	1.3555260	0.055041
##	[9,]	1.90703954	-2.3747685	0.4106013	-2.34955213	-0.4455350	-2.905444
##	[10,]	-0.09144695	1.2364622	-0.6458013	0.00346451	0.5983857	0.242345
##		[,8]	[,9]	[,10]			
##	[1,]	-1.0329155	1.42140711	1.30560568			
##	[2,]	-0.1178282	0.21645471	-0.48848642			
##	[3,]	0.5675319	0.05370436	0.59329679			
##	[4,]	-0.9468782	-0.14731870	1.30463971			
##	[5,]	-0.5524942	-1.17812024	-1.09789797			
##	[6,]	0.5266990	0.55646825	-0.85783015			
##	[7,]	0.7064474	-0.23973975	-1.11801576			
##	[8,]	0.4141258	-0.20672212	-1.43248556			
##	[9,]	-0.8818216	0.86505545	0.04558258			
##	[10,]	-1.0873272	0.75791330	0.59603916			

# Stripping attributes

---

- Many operations will strip attributes (generally why it's not a good idea to store important things in them)

`v`

```
##           index value
## the first      1     4
## second        2     5
## III           3     6
## attr(,"matrix_mean")
## [1] 3.5
```

`rowSums(v)`

```
## the first      second      III
##           5           7           9
```

`attributes(rowSums(v))`

```
## $names
## [1] "the first" "second"      "III"
```

- Generally **names** are maintained
- Sometimes, **dim** is maintained, sometimes not
- All else is stripped

# More on `names`

---

- The `names` attribute corresponds to the individual elements within a vector

```
names(v)
```

```
## NULL
```

```
names(v) <- letters[1:6]  
v
```

```
##           index value  
## the first      1      4  
## second        2      5  
## III           3      6  
## attr(,"matrix_mean")  
## [1] 3.5  
## attr(,"names")  
## [1] "a" "b" "c" "d" "e" "f"
```

- Perhaps more straightforward

```
v3a <- c(a = 5, b = 7, c = 12)
v3a
```

```
##  a  b  c
##  5  7 12
```

```
names(v3a)
```

```
## [1] "a" "b" "c"
```

```
attributes(v3a)
```

```
## $names
## [1] "a" "b" "c"
```

# Alternatives

---

```
v3b <- c(5, 7, 12)
names(v3b) <- c("a", "b", "c")
v3b
```

```
##  a  b  c
##  5  7 12
```

```
v3c <- setNames(c(5, 7, 12), c("a", "b", "c"))
v3c
```

```
##  a  b  c
##  5  7 12
```

- Note that **names** is **not** the same thing as **colnames**, but, somewhat confusingly, both work to rename the variables (columns) of a data frame. We'll talk more about why this is momentarily.

# Why names might be helpful

---

```
v
```

```
##           index value
## the first      1      4
## second         2      5
## III            3      6
## attr(,"matrix_mean")
## [1] 3.5
## attr(,"names")
## [1] "a" "b" "c" "d" "e" "f"
```

```
v["b"]
```

```
## b
## 2
```

```
v["e"]
```

```
## e
## 5
```

# Implementation of factors

---

## Quickly

```
fct <- factor(c("a", "a", "b", "c"))  
typeof(fct)
```

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

```
attributes(fct)
```

```
## $levels  
## [1] "a" "b" "c"  
##  
## $class  
## [1] "factor"
```

```
str(fct)
```

```
## Factor w/ 3 levels "a","b","c": 1 1 2 3
```



# More manually

---

```
# First create integer vector
int <- c(1L, 1L, 2L, 3L, 1L, 3L)

# assign some levels
attr(int, "levels") <- c("red", "green", "blue")

# change the class to a factor
class(int) <- "factor"

int
```

```
## [1] red    red    green blue   red    blue
## Levels: red green blue
```

# Implementation of dates

---

## Quickly

```
date <- Sys.Date()  
typeof(date)
```

```
## [1] "double"
```

```
attributes(date)
```

```
## $class  
## [1] "Date"
```

```
attributes(date) <- NULL  
date
```

```
## [1] 18730
```

- This number represents the days passed since January 1, 1970, known as the Unix epoch.

# Missing values

---

- Missing values breed missing values

```
NA > 5
```

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

```
NA * 7
```

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

- What about this one?

```
NA == NA
```

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

It is correct because there's no reason to presume that one missing value is or is not equal to another missing value.

# When missing values don't propagate

---

NA | TRUE

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

```
x <- c(NA, 3, NA, 5)  
any(x > 4)
```

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

# How to test missingness?

---

- We've already seen the following doesn't work

```
x == NA
```

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

- Instead, use `is.na`

```
is.na(x)
```

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

- When does this regularly come into play?

# Lists

---

# Lists

---

- Lists are vectors, but not *atomic* vectors
- Fundamental difference – each element can be a different type

```
list("a", 7L, 3.25, TRUE)
```

```
## [[1]]  
## [1] "a"  
##  
## [[2]]  
## [1] 7  
##  
## [[3]]  
## [1] 3.25  
##  
## [[4]]  
## [1] TRUE
```

# Lists

---

- Technically, each element of the list is a vector, possibly atomic
- The prior example included all *scalars*, which are vectors of length 1.
- Lists do not require all elements to be the same length

```
l <- list(  
  c("a", "b", "c"),  
  rnorm(5),  
  c(7L, 2L),  
  c(TRUE, TRUE, FALSE, TRUE)  
)  
l
```

```
## [[1]]  
## [1] "a" "b" "c"  
##  
## [[2]]  
## [1] 1.2009654 1.0447511 -1.0032086  
##  
## [[3]]  
## [1] 7 2  
##  
## [[4]]  
## [1] TRUE TRUE FALSE TRUE
```



# Check the list

---

```
typeof(l)
```

```
## [1] "list"
```

```
attributes(l)
```

```
## NULL
```

```
str(l)
```

```
## List of 4
## $ : chr [1:3] "a" "b" "c"
## $ : num [1:5] 1.201 1.045 -1.003 1.848 -0.667
## $ : int [1:2] 7 2
## $ : logi [1:4] TRUE TRUE FALSE TRUE
```

# Data frames as lists

---

- A data frame is just a special case of a list, where all the elements are of the same length.

```
l_df <- list(  
  a = c("red", "blue"),  
  b = rnorm(2),  
  c = c(7L, 2L),  
  d = c(TRUE, FALSE)  
)  
l_df
```

```
## $a  
## [1] "red"  "blue"  
##  
## $b  
## [1]  0.1055138 -0.4222559  
##  
## $c  
## [1] 7 2  
##  
## $d  
## [1] TRUE FALSE
```

```
data.frame(l_df)
```

```
##      a      b c      d  
## 1  red  0.1055138 7  TRUE  
## 2 blue -0.4222559 2 FALSE
```

# Subsetting

# Lists

---

# A nested list

---

Lists are often complicated objects. Let's create a somewhat complicated one

```
x <- c(a = 3, b = 5, c = 7)
l <- list(
  x = x,
  x2 = c(x, x),
  x3 = list(
    vect = x,
    squared = x^2,
    cubed = x^3)
)
```

# Subsetting lists

---

## Multiple methods

- Most common: `$`, `[`, and `[[`

```
l[1]
```

```
## $x  
## a b c  
## 3 5 7
```

```
typeof(l[1])
```

```
## [1] "list"
```

```
l[[1]]
```

```
## a b c  
## 3 5 7
```

```
typeof(l[[1]])
```

```
## [1] "double"
```

```
l[[1]]["c"]
```

```
## c  
## 7
```

# Named list

---

- Because the elements of the list are named, we can use `$`

```
l$x2
```

```
## a b c a b c  
## 3 5 7 3 5 7
```

```
l$x3
```

```
## $vect  
## a b c  
## 3 5 7  
##  
## $squared  
## a b c  
## 9 25 49  
##  
## $cubed  
## a b c  
## 27 125 343
```

# Subsetting nested lists

---

- Multiple  $\$$  if all named

```
l$x3$squared
```

```
##  a  b  c  
##  9 25 49
```

- Note this doesn't work on named elements of an atomic vector, just the named elements of a list

```
l$x3$squared$b
```

```
## Error in l$x3$squared$b: $ operator is invalid for atomic vectors
```

But we could do something like...

```
l$x3$squared["b"]
```

```
## b
```

```
## 25
```



# Alternatives

---

- You can always use logical
- Indexing works too

```
l[c(TRUE, FALSE, TRUE)]
```

```
## $x
## a b c
## 3 5 7
##
## $x3
## $x3$vect
## a b c
## 3 5 7
##
## $x3$squared
## a b c
## 9 25 49
##
## $x3$cubed
## a b c
## 27 125 343
```

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

```
## $x
## a b c
## 3 5 7
##
## $x3
## $x3$vect
## a b c
## 3 5 7
##
## $x3$squared
## a b c
## 9 25 49
##
## $x3$cubed
## a b c
## 27 125 343
```

# Careful with your brackets

---

```
l[[c(TRUE, FALSE, FALSE)]]
```

```
## Error in l[[c(TRUE, FALSE, FALSE)]]: recursive indexing failed at level
```

- Why doesn't the above work?

# Subsetting in multiple dimensions

- Generally we deal with 2d data frames
- If there are two dimensions, we separate the `[` subsetting with a comma

```
head(mtcars)
```

```
##           mpg  cyl  disp  hp  drat    wt    qsec vs  am  gear  carb
## Mazda RX4      21.0    6  160  110  3.90  2.620  16.46  0   1     4     4
## Mazda RX4 Wag  21.0    6  160  110  3.90  2.875  17.02  0   1     4     4
## Datsun 710     22.8    4  108   93  3.85  2.320  18.61  1   1     4     1
## Hornet 4 Drive  21.4    6  258  110  3.08  3.215  19.44  1   0     3     1
## Hornet Sportabout 18.7    8  360  175  3.15  3.440  17.02  0   0     3     2
## Valiant        18.1    6  225  105  2.76  3.460  20.22  1   0     3     1
```

```
mtcars[3, 4]
```

```
## [1] 93
```

# Empty indicators

---

- An empty indicator implies "all"

Select the entire fourth column

```
mtcars[,4]
```

```
## [1] 110 110 93 110 175 105 245 62 95 123 123 180 180 180 205 215 230
## [22] 150 150 245 175 66 91 113 264 175 335 109
```

Select the entire 4th row

```
mtcars[4, ]
```

```
##           mpg cyl  disp  hp drat   wt  qsec vs am gear carb
## Hornet  4 Drive 21.4   6  258 110 3.08 3.215 19.44 1  0    3    1
```

# Data types returned

---

- By default, each of the prior will return a vector, which itself can be subset

The following are equivalent

```
mtcars[4, c("mpg", "hp")]
```

```
##                mpg  hp
## Hornet  4 Drive 21.4 110
```

```
mtcars[4, ][c("mpg", "hp")]
```

```
##                mpg  hp
## Hornet  4 Drive 21.4 110
```

# Return a data frame

---

- Often, you don't want the vector returned, but rather the modified data frame.
- Specify `drop = FALSE`

```
mtcars[,4]
```

```
## [1] 110 110 93 110 175 105 245 62 95 123 123 180 180 180 205 215 230
## [22] 150 150 245 175 66 91 113 264 175 335 109
```

```
mtcars[,4, drop = FALSE]
```

```
##                hp
## Mazda RX4      110
## Mazda RX4 Wag  110
## Datsun 710      93
## Hornet 4 Drive  110
## Hornet Sportabout 175
## Valiant        105
## Duster 360     245
## Merc 240D       62
## Merc 230       95
```

# tibbles

---

- Note dropping the data frame attribute is the default for a `data.frame` but NOT a `tibble`.

```
mtcars_tbl <- tibble::as_tibble(mtcars)
mtcars_tbl[,4]
```

```
## # A tibble: 32 x 1
##       hp
##   <dbl>
## 1    110
## 2    110
## 3     93
## 4    110
## 5    175
## 6    105
## 7    245
## 8     62
## 9     95
## 10   123
## # ... with 22 more rows
```

# You can override this

---

```
mtcars_tbl[,4, drop = TRUE]
```

```
## [1] 110 110 93 110 175 105 245 62 95 123 123 180 180 180 205 215 230
## [22] 150 150 245 175 66 91 113 264 175 335 109
```



# More than two dimensions

---

- Depending on your applications, you may not run into this much

```
array <- 1:12
dim(array) <- c(2, 3, 2)
array
```

```
## , , 1
##
##      [,1] [,2] [,3]
## [1,]    1    3    5
## [2,]    2    4    6
##
## , , 2
##
##      [,1] [,2] [,3]
## [1,]    7    9   11
## [2,]    8   10   12
```

# Subset array

---

Select just the second matrix

```
array[ , 2]
```

```
##      [,1] [,2] [,3]  
## [1,]    7    9   11  
## [2,]    8   10   12
```

Select first column of each matrix

```
array[ , 1, ]
```

```
##      [,1] [,2]  
## [1,]    1    7  
## [2,]    2    8
```

# Back to lists

---

Why are they so useful?

- Fairly obviously, they're much more flexible
- Often returned by functions, for example, `lm`

```
m <- lm(mpg ~ hp, mtcars)
str(m)
```

```
## List of 12
## $ coefficients : Named num [1:2] 30.0989 -0.0682
##   ..- attr(*, "names")= chr [1:2] "(Intercept)" "hp"
## $ residuals    : Named num [1:32] -1.594 -1.594 -0.954 -1.194 0.541 ...
##   ..- attr(*, "names")= chr [1:32] "Mazda RX4" "Mazda RX4 Wag" "Datsun 7
## $ effects      : Named num [1:32] -113.65 -26.046 -0.556 -0.852 0.67 ...
##   ..- attr(*, "names")= chr [1:32] "(Intercept)" "hp" "" "" ...
## $ rank         : int 2
## $ fitted.values: Named num [1:32] 22.6 22.6 23.8 22.6 18.2 ...
##   ..- attr(*, "names")= chr [1:32] "Mazda RX4" "Mazda RX4 Wag" "Datsun 7
## $ assign       : int [1:2] 0 1
## $ qr          : List of 5
##   ..$ qr      : num [1:32, 1:2] -5.657 0.177 0.177 0.177 0.177 ...
```

# Summary

---

- Atomic vectors must all be the same type
  - implicit coercion occurs if not (and you haven't specified the coercion explicitly)
- Lists are also vectors, but not atomic vectors
  - Each element can be of a different type and length
  - Incredibly flexible, but often a little more difficult to get the hang of, particularly with subsetting

# Any time left?

---

Practice: Fit the model on the slide two previous

```
m <- lm(mpg ~ hp, mtcars)
```

Pull each of the following from the model object

- **effects**
- **names** of the **effects**
- **qr** matrix
- third row of the **qr** matrix
- fifth **rowname** of the **qr** matrix
- **data.frame** used in the estimation of the model

# Next time

---

Loops with base  $R$