Introduction to R, Part II

Some useful functions

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Getting data into, and out of, R

Getting Data Into R

read.table function

- Needs at least 3 arguments
 - Name of the file to read in (in quotes)
 - Whether or not the file has a header row
 - How fields are delimited (commas, tabs, space, etc.)

Getting Data Into R

read.table function

afdata = read.table("agefat.csv", header = TRUE, sep = ",")

x	age	fat	sex
1	24	15.5	male
2	37	20.9	male
3	41	18.6	male
4	60	28	male
5	31	34.7	female
6	39	30.2	female
•••	•••	•••	•••

Getting Data Out of R write.csv function

write.csv(afdata, "myafdata.csv")

Getting Data Out of R

write.csv function

- By default, R will put quotes around anything it thinks is text
- Also adds a column of row "names"
 - This can be annoying

```
"","X","age","fat","sex"
"1",1,24,15.5,"male"
"2",2,37,20.9,"male"
"3",3,41,18.6,"male"
"4",4,60,28,"male"
"5",5,31,34.7,"female"
"6",6,39,30.2,"female"
"7",7,58,21.3,"male"
"8",8,23,9.5,"male"
```

Getting Data Out of R

write.csv function

```
write.csv(afdata, file = "myafdata.csv", row.names = FALSE, quote = FALSE)
```

```
X,age,fat,sex
1,24,15.5,male
2,37,20.9,male
3,41,18.6,male
4,60,28,male
5,31,34.7,female
6,39,30.2,female
7,58,21.3,male
8,23,9.5,male
```

```
summary()
```

• Identifies the type, or "class" of the object, and returns appropriate summary information

summary()

```
summary(afdata)
   X
                               fat
                age
                                             sex
Min.
      :1
           Min. :23.00
                           Min.
                               :7.80 female :15
                           1st Qu. :21.30 male :10
1st Qu. :7 1st Qu. :37.00
Median :13 Median :49.00
                           Median :29.10
Mean :13 Mean :44.96
                           Mean :27.37
3rd Qu. :19 3rd Qu. :57.00
                           3rd Qu. :33.00
                           Max. :42.00
          Max. :61.00
Max. :25
```

```
sd()
```

- Not sure why summary doesn't provide standard deviation
- Can get with sd

```
sd(afdata$age)
[1] 13.14813

sd(afdata$fat)
[1] 8.679272
```

head()

- Shows just the first few lines of an object (the "header")
 - Good for identifying the structure / format of large data files

```
head(afdata)
                fat
        age
                         sex
       24
                15.5
                        male
       37
                20.9
                        male
                18.6
       41
                        male
       60
                28.0
                        male
       31
                34.7
                        female
                        female
                30.2
       39
```

tail()

- Shows just the last few lines of an object (the "header")
 - Good for identifying the structure / format of large data files

```
tail(afdata)
                fat
   X
       age
                        sex
       56
20
   20
                32.5
                        female
21 21
       57
                30.3
                        female
                        female
22
                33.0
  22
       58
23
   23
       58
                33.8
                        female
                41.1
                        female
24 24
       60
25 25
                34.5
                        female
       61
```

```
str()
```

- Summarizes the structure of an object
 - Very useful for complex data sets (*e.g.*, lists and arrays)

```
str(afdata)

'data.frame': 25 obs. of 4 variables:
$ X: int 1 2 3 4 5 6 7 8 9 10 ...
$ age : int 24 37 41 60 31 39 58 23 23 27 ...
$ fat : num 15.5 20.9 18.6 28 34.7 30.2 21.3 9.5 27.9 7.8 ...
$ sex : Factor w/ 2 levels "female", "male": 2 2 2 2 1 1 2 2 1 2
```

length()

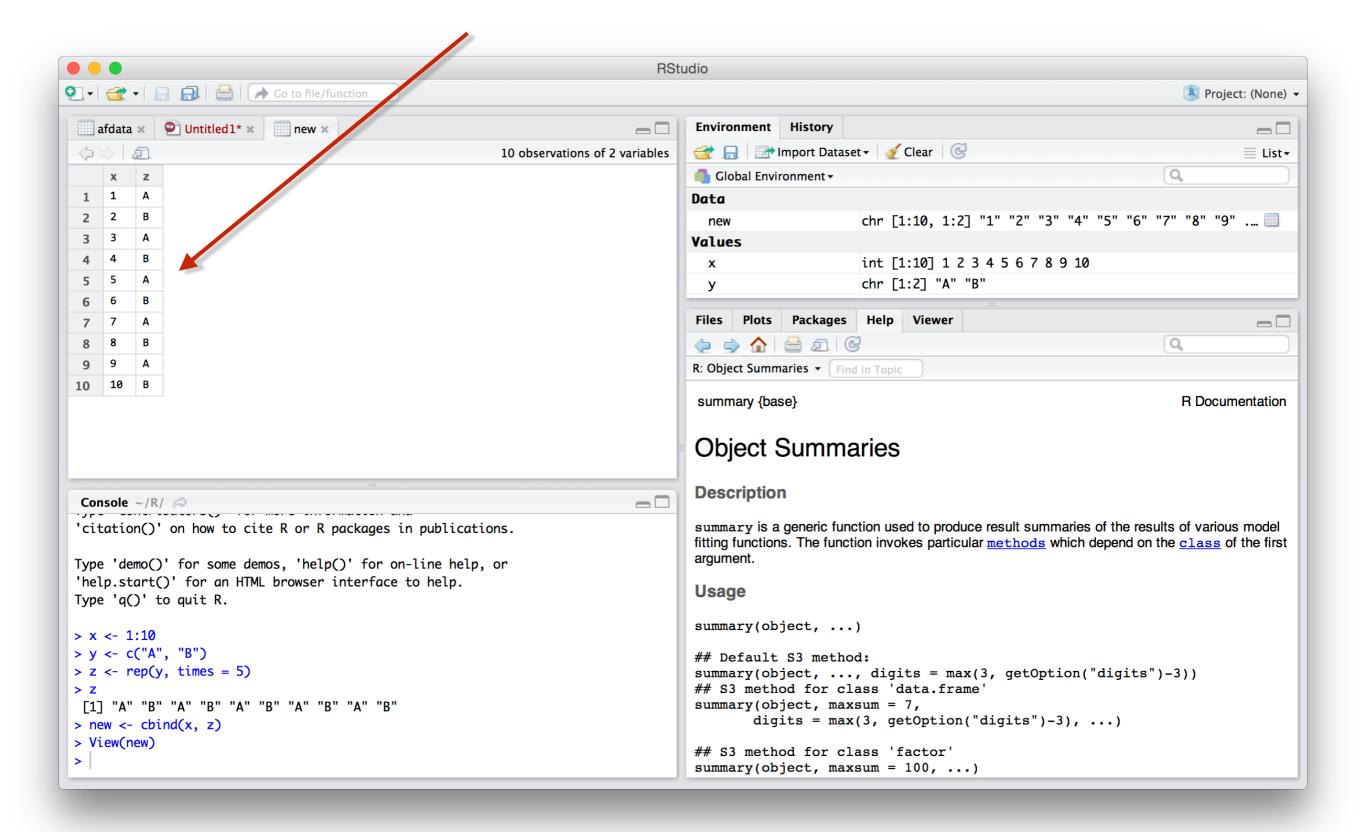
- Returns the length of a vector
 - Very useful when you want to run the same function on files of different sizes

```
length(afdata[ ,1])
[1] 25
```

```
cbind()
```

 Combines different data files as different columns in one new data frame

```
x = 1:10
y = c("A", "B")
z = rep(y, times = 5)
new = cbind(x, z)
```



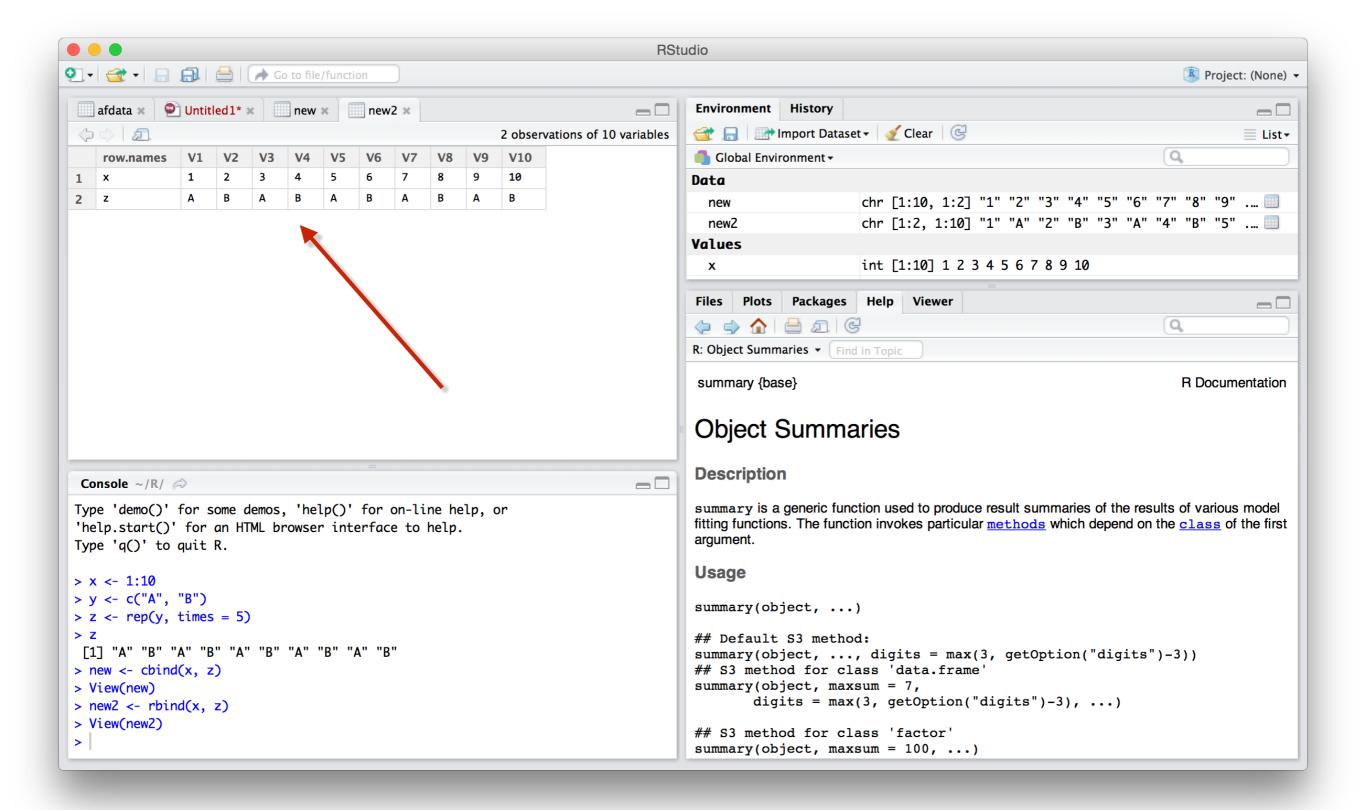
rbind()

• Combines different data files as different rows in one new data frame

```
rbind()
```

• Combines different data files as different rows in one new data frame

```
x = 1:10
y = c("A", "B")
z = rep(y, times = 5)
new2 = rbind(x, z)
```



```
sample()
```

- Can be use to randomize a list (key to simulations!)
 - Can sample with or without replacement

```
sample()
```

- Can be use to randomize a list (key to simulations!)
 - Can sample with or without replacement

```
myList = 1:10

myList
1 2 3 4 5 6 7 8 9 10
```

```
sample(myList, size = 10, replace = FALSE)
[1] 6 2 10 5 4 3 1 9 8 7
```

```
sample(myList, size = 10, replace = TRUE)
[1] 10 10 9 6 3 4 1 8 3 3
```

```
print()
```

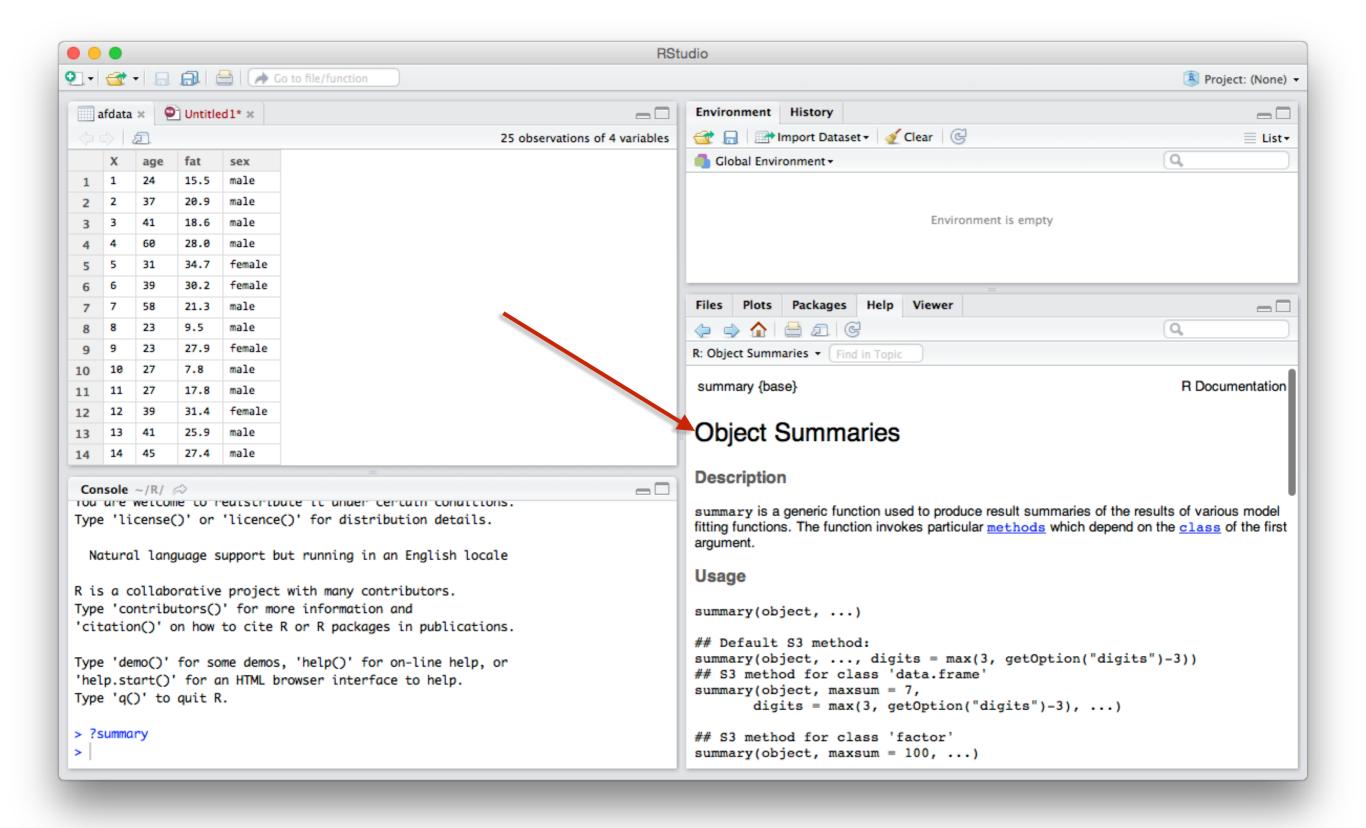
- Prints text to screen
 - May not seem useful now, but key for checking code

```
s = "Bayesian analysis is cool!"
print(s)
[1] "Bayesian analysis is cool!"
```

? - Help

• ? then function name will open the help information for that function

?summary



Google is your friend!

- Can find help for just about anything in R
- Countless number of good online resources for R

Questions?