#### Lecture outline:

- · Data frame basics
- Loading data into data frames
- Basic statistical operations

In [1]:

%load\_ext rmagic

# **Data frame basics**

# Constructing and manipulating data frames

The R data.frame class

The central class in R is the class

data.frame

that represents data tables in a similar way than that of Pandas DataFrame.

**Remark:** The period in the class name data.frame has no programming meaning; it is a mere convention among R programmers, used to group related variables, functions, methods, and classes. (We already saw that convention with the naming of class methods.)

As we saw in the last lecture, classes in R are just "enhanced lists" with

- a class name stored in the list hidden "class string", which is set by using the class (x) function
- a class constructor, which is a regular function used to contruct class instances (or objects)
- a collection of methods and generic functions acting on the class instances

Since data frames are R classes, one can think of them as enhanced lists

- with the hidden class string changed to 'data.frame',
- with a contructor named data.frame,
- with a collection of methods and a collections of associated generic functions.

The underlying list of a data frame contains the vectors representing the data table columns (i.e. the statistical variables).

One passes these vectors to the data frame constructor to populate its underlying list:

```
X = c(a=1, b=2, c=3)
Y = c(a=3, b=9, c=1)
Z = c(a='u', b='v', c='w')

df1 = data.frame(X, Y, Z)

print(df1)
```

```
X Y Z
a 1 3 u
b 2 9 v
c 3 1 w
```

d 12 33 k

## Data frame transposition

One can always transpose a data frame using the **transpose generic function** t(x):

**Remark:** In a data frame, the **values in a given column** should all be of the **same type**. **After transposition**, some column values may end up beeing of different types. In this case, all the **column values will be converted (casted, or coerced) to a matching type** (in the example above: character vectors, since it is possible to cast a number into a string ('1' instead of 1), but the opposite operation does not make sense.

#### Adding new observations to an existing data frame

Suppose we have two data frames, representing **two samples** of the **same population** for **the same variables**. For instance, the data frame df1 above and the following new one:

```
In [13]: %%R

X = c(d=12, e=12, f=63)
Y = c(d=33, e=96, f=0)
Z = c(d='k', e='l', f='m')

df2 = data.frame(X, Y, Z)

print(df2)

X Y Z
```

```
e 12 96 1
f 63 0 m
```

We can combine these two sets of observations into a single data frame using the generic **row binding function** 

```
rbind(df1, df2)
```

that will return a data frame with the combined sets of observations:

### Adding new variables to an existing data frame

Suppose we have two data frames, representing the **same sample** but containing values for **different population variables**. For instance, the data frame df3 above and the following new one:

```
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In [16]:
         df4 = data.frame(sex=c(a='M', b='F', c='M', d='F', e='M', f='F'))
         print(df4)
           sex
            Μ
         а
         b
            F
            M
         d
            F
            Μ
         е
         f
             F
```

We add additional variables to a data frame by using the generic column bind function

```
cbind(df1, df2)
```

that returns a data frame with additional column representing the new variables:

```
In [17]: %%R

df5 = cbind(df3, df4)

print(df5)
```

```
X Y Z sex
a 1 3 u M
b 2 9 v F
c 3 1 w M
d 12 33 k F
e 12 96 1 M
f 63 0 m F
```

### Bracket operator to retrieve columns

One accesses the **columns of a data frame**, the very same way one accesses the **elements of a list**. So you know how to do that from the previous lecture.

Since the **data frame columns are vectors**, one can do **vectorized arithmetic** with them. This allows us to produce new variables by adding, multipliying, taking square roots, etc. from data frame columns.

# Example: a grade data frame (as always)

### Formal setting: universe and variables

Let's have a look at our favorite grade example. So we have a population (or universe) of three students in a certain class:

$$\Omega = \Big\{ \mathrm{Bob}, \, \mathrm{Aline}, \, \mathrm{Agnes} \Big\}$$

**Remark:** The element in the set  $\Omega$  are meant to represent actual students, with all their characteristics/features/variables, and not only the student names.

Since the students have taken three exams in this class (the first and second midterm, denoted by M1 and M2, and the final exam denoted by F), we have to three **quantitative** variables associated with these exams: that is, we have three mathematical functions:

$$M1,\,M2,\,F:\Omega o [0,100]$$

For instance, the grade of student  $s \in \Omega$  at the final exam is F(s), which is a number between 0 and 100.

#### Data frame instanciation from vectors

Let's now construct three R vectors representing the values of our variables M1, M2, and F:

```
M1 = c(Bob=67, Aline=88, Agnes=99)

M2 = c(Bob=82, Aline=91, Agnes=100)

F = c(Bob=3, Aline=38, Agnes=97)
```

and construct an R data frame with them:

```
In [19]: %%R
grades = data.frame(M1, M2, F)
```

R data frames have a method print.data.frame method implemented, so we can invoke the generic function print on data frame instances:

```
In [20]: %%R

print(grades)

M1 M2 F

Bob 67 82 3

Aline 88 91 38

Agnes 99 100 97
```

# Maliciously changing the class string of a data frame...

One sees that the print function is not the same as the print function for R lists: it's much better.

By curiosity, let's change the class string of our data.frame object grades, and see if it changes the behavior of the generic function print:

```
In [21]: %%R
    class(grades) = 'list'
    print(grades)

$M1
    [1] 67 88 99

$M2
    [1] 82 91 100

$F
    [1] 3 38 97

attr(,"row.names")
    [1] "Bob" "Aline" "Agnes"
```

We see that the generic function print now understand that our object is no longer a data frame but simply a list. As a result, it prints the underlying list the old fashion way.

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Let's restore the status our fallen data frame grades:

## Accessing and computing with columns

Accessing data frame columns amount to accessing the elements of the underlying list, which we know how to do. We have two syntax for that:

- the dollar sign operator syntax
- the bracket operator syntax

The return value of these operators is the corresponding data frame column: i.e. a R vector. Thus we can perform arithmetical vectorized operation, in the very same way as we did with Pandas Series. For instance, let's compute the total class grade with the following crazy and mean formula:

$$TG=\sqrt{M_1\cos^2(F)+M_2\sin^2(F)}$$

#### Accessing rows and single variable values

R data frames enjoy a similar **double entry bracket syntax** as Numpy arrays.

 The data frame rows are indexed from 1 to the total number of rows (i.e. the number of individuals in our sample\*\*) • The data frame columns are indexed from 1 to the total number of columns (i.e. the number of variables we are considering).

Now, given a data frame data the bracket operator

```
data[i, j]
```

data[i,]

will return the value of variable j (i.e. in the  $j^{th}$  column) for individual i (i.e. in the  $i^{th}$  row).

will return the  $i^{th}$  row as a **one row data frame**.

# Loading data into a data frame

As for Pandas data frames, we have functions that allow us to load data tables located either on the **internet** at a given **url** or on the **local file system** at a given **path**.

The functions

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```
read.table(address, row.names, col.names, header, sep)
read.csv(address, row.names, col.names, header)
```

will return a data frame using the **data table** located at the value passed to the argument address.

The **column names** will be inferred from the first line of the data table if the parameter header is set to TRUE (the default is FALSE). They can also be specified by passing to the argument col.name a character vector.

The row names can be specified by passing to the argument row.names either

- a character vector
- the column index to use for the row names

The difference between these two functions is that

- read.table expects a blanck character (i.e. spaces, tabs, etc.) as separator by default, although the separator can be specified by setting the argument sep
- read.csv expects a comma separated value (csv) data table

Have a look <a href="http://stat.ethz.ch/R-manual/R-devel/library/utils/html/read.table.html">http://stat.ethz.ch/R-manual/R-devel/library/utils/html/read.table.html</a>) for a description of all the possible arguments that can be passed to these functions, as well as for their default values.

# **Examples: loading tabular data**

#### Comma separated values

Consider the comma separated file located at:

```
In [36]: csv_url = 'http://www.stat.berkeley.edu/classes/s133/data/world.txt'
    !curl $csv_url 2>/dev/null | head -4

    country,gdp,income,literacy,military
    Albania, 4500, 4937, 98.7,56500000
    Algeria,5900,6799,69.8,2.48e+09
    Angola,1900,2457,66.8,183580000
```

This variable <code>csv\_url</code> is defined in a Python cell, and thus not accessible by default to R cells. To have access to it in R cells, one can use the <code>-i</code> (i for "input") option of the magic command <code>%%R</code>:

To display the first or last entries of a data frame, one has the generic functions:

```
head(x) tail(x)
```

```
In [38]: %%R
print(head(countries))
```

```
gdp income literacy
                               military
          4500 4937
                         98.7 5.6500e+07
Albania
Algeria
          5900
                6799
                         69.8 2.4800e+09
Angola
                         66.8 1.8358e+08
         1900 2457
                       97.2 4.3000e+09
Argentina 11200 12468
Armenia
                       99.4 1.3500e+08
         3900 3806
Australia 28900 29893
                       99.9 1.6650e+10
```

#### **Custom separator**

rank|name|box|date

- 1|Avatar|\$759.563|December 18, 2009
- 2|Titanic|\$600.788|December 19, 1997
- 3|The Dark Knight|\$533.184|July 18, 2008

```
In [40]: %%R -i custom_url
movies = read.table(custom_url, header=TRUE, sep='|')
```

```
In [41]: %%R
print(head(movies))
```

```
rank
                                     name
                                               box
                                                                date
1
    1
                                   Avatar $759.563 December 18, 2009
2
                                  Titanic $600.788 December 19, 1997
3
                                                       July 18, 2008
                          The Dark Knight $533.184
    4 Star Wars: Episode IV - A New Hope $460.998
                                                      May 25, 1977
4
                                                       May 19, 2004
5
                                  Shrek 2 $437.212
              E.T. the Extra-Terrestrial $434.975
                                                       June 11, 1982
```

#### Blanck-space separated tables

In [43]: table\_url = 'http://www.stat.berkeley.edu/classes/s133/data/pop.txt'
 !curl \$table\_url 2>/dev/null | head -4

State	pop2004	pop2003
Alabama	4530182	4500752
Alaska	655435	648818

Arizona 5743834 5580811

```
In [47]: %%R -i table_url
states = read.table(table_url, header=TRUE, , row.names=1)
print(head(states))
```

```
pop2004 pop2003
Alabama 4530182 4500752
Alaska 655435 648818
Arizona 5743834 5580811
Arkansas 2752629 2725714
California 35893799 35484453
Colorado 4601403 4550688
```

# **Basic statistical operations**

## **Summary statistics**

One gets **summary statistics** for the variables stored in a data frame, very much the same way as we did with Pandas. The difference is that we use R data frame methods through the generic function:

```
summary (data)
```

The argument data can be

- a whole data frame, in which case summary will return a table containing the summary statistics for all data frame variables
- a numeric vector, such as a data frame column

Let's print the summary statistics of our data frame containing the population of all states in 2004 (first column) and in 2003 (second column):

```
In [48]: %%R
stats_all = summary(states)
summary_2004 = summary(states$pop2004)
```

```
In [49]: %%R
print(stats_all)
```

```
pop2004 pop2003

Min. : 506529 Min. : 501242

1st Qu.: 2375472 1st Qu.: 2323889

Median : 4522976 Median : 4498543

Mean : 6123315 Mean : 6060413

3rd Qu.: 6282303 3rd Qu.: 6255088

Max. :35893799 Max. :35484453
```

```
In [29]: %%R

print(summary_2004)

Min. 1st Qu. Median Mean 3rd Qu. Max.
506500 2375000 4523000 6123000 6282000 35890000
```

#### R basic statistical functions

R has a number of useful numerical built-in functions.

Since the basic type representing numbers in R is vectorized, all the numeric functions take **numeric vectors as input**.

Therefore, all these built-in functions can take data frame columns representing quantitative variables as their input.

Explore by yourself the following list of functions:

#### Statistics:

#### Some other useful math functions:

```
abs(x)
sqrt(x)
ceiling(x)
floor(x)
trunc(x)
round(x, digits=n)
signif(x, digits=n)
cos(x), sin(x), tan(x)
acos(x), cosh(x), acosh(x), etc.
log(x)
log10(x)
exp(x)
```