

Topic 3

# Loading data into R

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Klaipeda, 28-30 September 2011

## 3. Loading data into R

### 3.1. Using Excel for data preparation

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# Preparing data for R

- *Microsoft Excel* is one of the most common software used to prepare datasets for R
- An *Excel* file is usually saved as a `.txt` or `.csv` file before loading it into R
- Other formats (e.g., SPSS, SAS, Access, SQL, etc.) are possible – see package `foreign`

# Some rules to remember when preparing data for R

- No empty cells; missing values should be replaced with as NA
- Name the columns, e.g.:
- No spaces in the variable names
- No names starting with numbers or dots
- Put the file into the current R's working directory

	A	B	C	D
1	Treatment	Barrel	Length	Weight
2	Control	Control3	29.28	0.992
3	Control	Control3	29.83	0.772
4	Control	Control3	31.93	0.894
5	Control	Control3	26.63	0.822

# Example dataset we will begin with

- Experiment on the influence of pH on the shell length / tissue dry weight in *Dreissena polymorpha*
- Four pH treatments: Control (Lake Ontario water), A (7.3), B (7.1), and C (6.9)
- Each treatment has three replicates: Control1-Control3, A1-A3, B1-B3, and C1-C3



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### 3.2. Reading data in

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# Set up the working directory

- Use the command

```
> setwd("~/Introductory R  
Course/R_Course_Datasets")
```

- Or in **RStudio** do

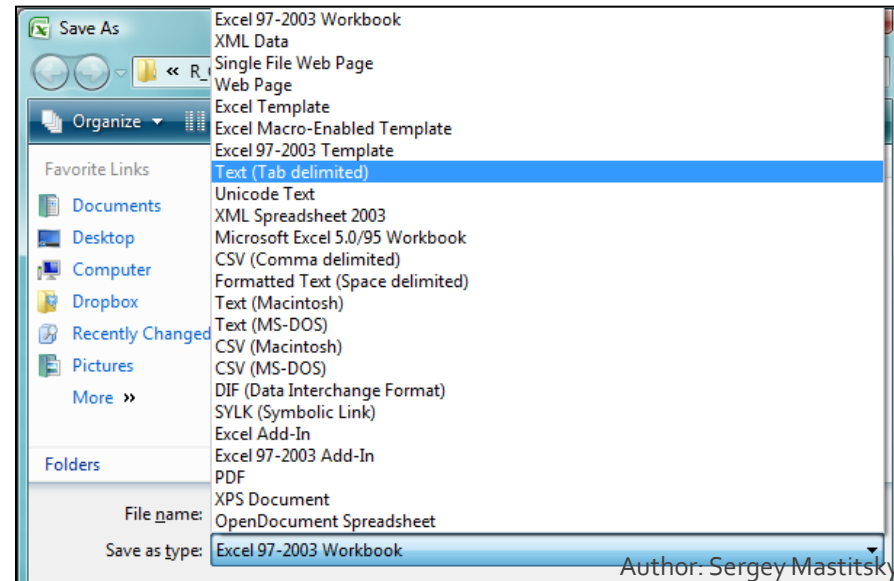
**Tools -> Set Working Directory -> Choose  
Directory -> ...your Desktop -> folder  
“Introductory R Course” -> folder  
“R\_Course\_Datasets”**

# Preparing the file in *Excel*

- Open the file `pH_down_experiment.xls`
- Quickly examine the data
- Save the file as a tab-delimited text file with the same name:

Save as -> Save as type -> Text (Tab delimited)

- Open the file in a text editor to see how it looks like





# The `read.table()` function

- The main “working horse” to read data in R
- Its main arguments:
  - `file = "some.name.txt"`: name of the file to read in (can be a URL link as well)
  - `header = TRUE`: indicates if the columns have names
  - `sep = "\t"` or `sep = ", "`: separator of values in the file

# Reading our example dataset in:

```
> LWdata <- read.table(  
  file = "pH_down_experiment.txt",  
  header = TRUE,  
  sep = "\t")  
> names(LWdata)  
> head(LWdata)  
> LWdata
```

# Saving the workspace

```
> save.image(file =  
  "pH_experiment.rda")
```

- Or in **RStudio** do

**Workspace -> Save Workspace As...**

pH\_experiment.rda