# Visualization of omics data

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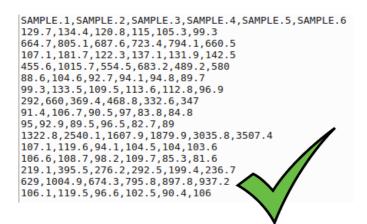
# Input files

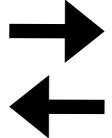
- A text file is a file formed by alphanumeric characters, line breaks and tabs.
- Text files can be opened without any special software.

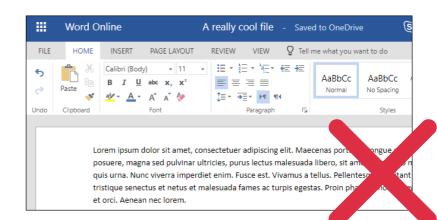
#### >SEQUENCE 1

MTEITAAMVKELRESTGAGMMDCKNALSETNGDFDKAVQLLREKGLGKAAKKADRLAAEG LVSVKVSDDFTIAAMRPSYLSYEDLDMTFVENEYKALVAELEKENEERRRLKDPNKPEHK IPQFASRKQLSDAILKEAEEKIKEELKAQGKPEKIWDNIIPGKMNSFIADNSQLDSKLTL MGQFYVMDDKKTVEQVIAEKEKEFGGKIKIVEFICFEVGEGLEKKTEDFAAEVAAQL >SEQUENCE 2

SATVSEINSETDFVAKNDQFIALTKDTTAHIQSNSLQSVEELHSSTINGVKFEFYLKS ATIGENLVVRRFATLKAGANGVVNGYIHTNGRVGVVIAAACDSAEVASKSRD

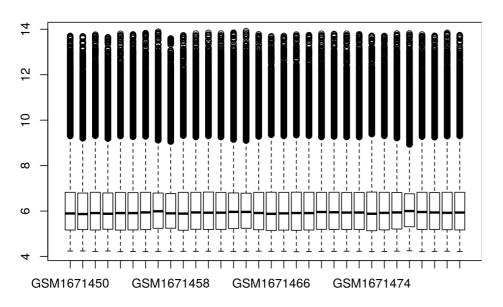


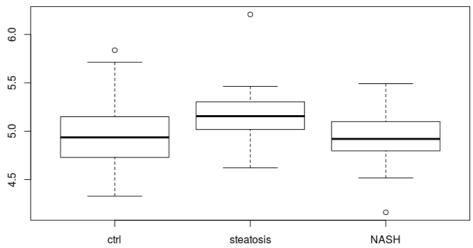




010		▼ fx 2					
	Α	В	С	D	E	F	
1	SAMPLE.1	SAMPLE.2	SAMPLE.3	SAMPLE.4	SAMPLE.5	SAMPLE.6	
2	129.7	134.4	120.8	115	105.3	99.3	
3	664.7	805.1	687.6	723.4	794.1	660.5	
4	107.1	181.7	122.3	137.1	131.9	142.5	
5	455.6	1015.7	554.5	683.2	489.2	580	
6	88.6	104.6	92.7	94.1	94.8	89.7	
7	99.3	133.5	109.5	113.6	112.8	96.9	
8	292	660	369.4	468.8	332.6	347	
9	91.4	106.7	90.5	97	83.8	84.8	
10	95	92.9	89.5	96.5	82.7	89	
11	1322.8	2540.1	1607.9	1879.9	3035.8	3507.4	
12	107.1	119.6	94.1	104.5	104	103.6	
13	106.6	108.7	98.2	109.7	85.3	81.6	
14	219.1	395.5	276.2	292.5	199.4	236.7	
15	629	1004.9	674.3	795.8	897.8	937.2	
16	106.1	119.5	96.6	102.5	90.4	106	
17							

Box plot (or whisker plot) are commonly used to compare datasets.

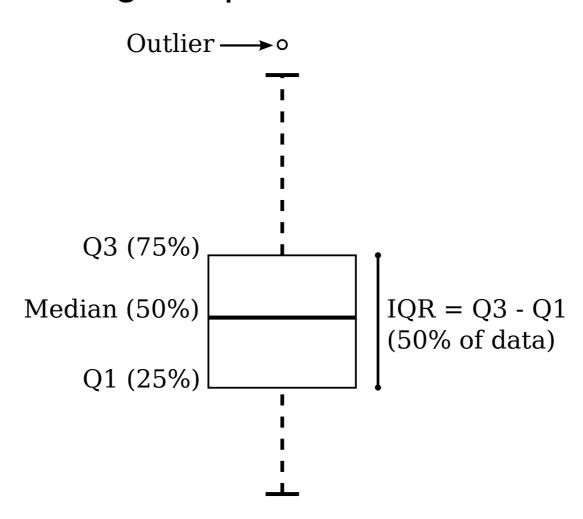




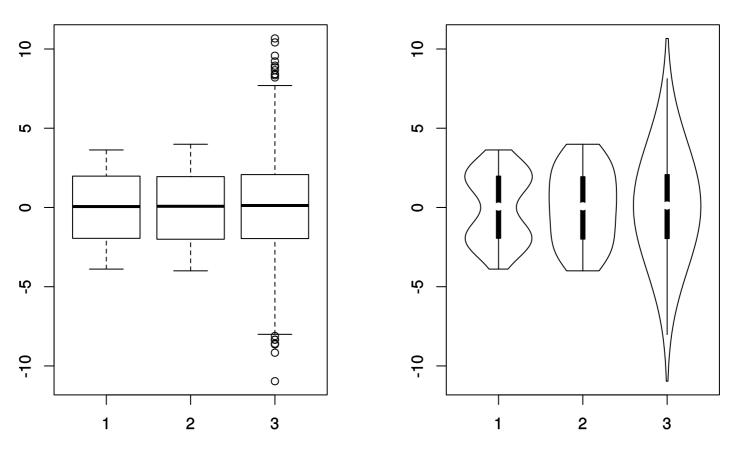
Normalized data from microarray experiment GSE68421

Expression of the gene ISG15 among several group of patients (GSE48452)

### **Understanding Boxplots**

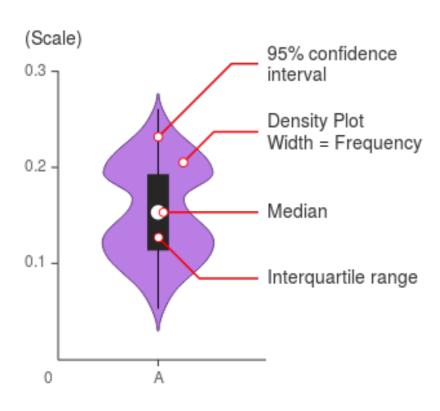


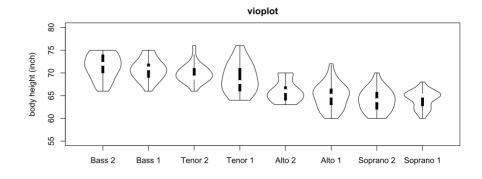
### Variants of box plot: violin plot and bean plot

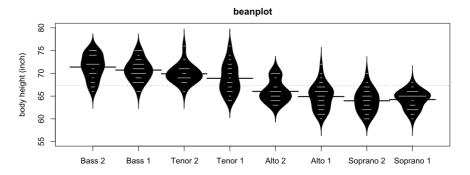


Box plot (left) and violin plot (right) from bimodal (1), uniform (2) and normal (3) distributions

### Variants of box plot: violin plot and bean plot







Source: Kampstra, P. (2008). Beanplot: A Boxplot Alternative for Visual Comparison of Distributions.

Source: https://datavizcatalogue.com/methods/violin\_plot.html

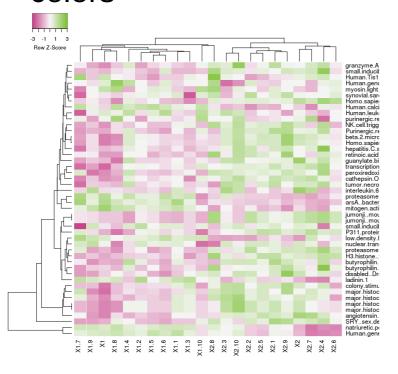
- Accepted file formats:
  - Tab, comma or semicolon separated files
- Data Structure
  - The first row is used to assign column names.
  - The other rows only with numeric values
- It is possible to paste data directly.

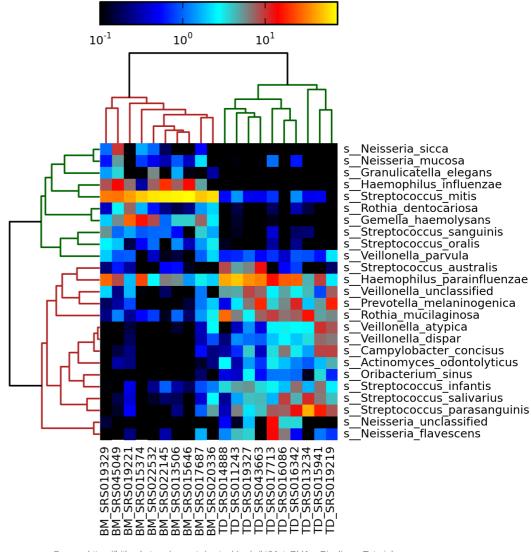
http://shiny.chemgrid.org/boxplotr/

http://bioinfo.cipf.es/BoxPlotR.shiny/

# Heatmapper

A heatmap is a graphical representation of data where the individual values contained in a matrix are represented as colors





# Heatmapper

- Accepted file formats:
  - Tab separated files with extensions .txt, .dat, .tsv or .tab.
  - First sheet of a Excel file with extension .xlsx
- Data Structure
  - The first (header) row is used to assign column names.
  - A column labeled UNIQID is required. (or a NAME column)
  - If a column labeled NAME exists (optional), it will be used to assign row names.
  - Any additional columns containing non-numeric data will be ignored.
  - Data values can be positive or negative numbers.
- Input data can have up to 2500 rows and 300 columns

http://heatmapper.ca/

# Exercises

