

Initiation to R software Session III

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Import/Export text files

Import text files

To load text files stored in your computer, use the function `read.table()`.

```
read.table(file, sep, header)
```

- ▶ `file`: the name of the file (character)
- ▶ `sep`: separator used in file (" " by default)
- ▶ `header`: TRUE if file contains columns names, FALSE by default

```
tab = read.table("../data/age_gender.txt", header = T)  
head(tab, 3)
```

```
##    age gender  
## 1   28      F  
## 2   36      H  
## 3   45      F
```

Import text files: variants of `read.table()`

- ▶ `file.choose()`: choose a file through the GUI.
- ▶ `read.csv()`: read CSV files.
- ▶ `read.delim()`: read delimited text files.
- ▶ `read.fwf()`: read fixed-width-formatted files.

R can read files in specific software formats (Excel, SAS, SPSS, Stata) using the functions from the packages `foreign`.

```
# Try this  
install.packages("foreign")  
library(foreign)  
?read.dta
```

Export text files

To export a `data.frame` into text files on your working directory, use the function `write.table()`.

```
write.table(x, file, append, col.names, row.names)
```

- ▶ `x`: `data.frame`
- ▶ `file`: name of the file in which to write
- ▶ `append`: if `TRUE`, add to an (eventually existing) file, if `FALSE`, overwrite the existing file (default)
- ▶ `col.names` and `row.names`: if `TRUE`, write the columns/rows names

```
write.table(tab, "age_gender.txt", row.names = T,  
            col.names = T)
```

`write()` is the same as `write.table()` with less options.

Save/Load R objects

R objects can be saved in both ascii and binary formats:

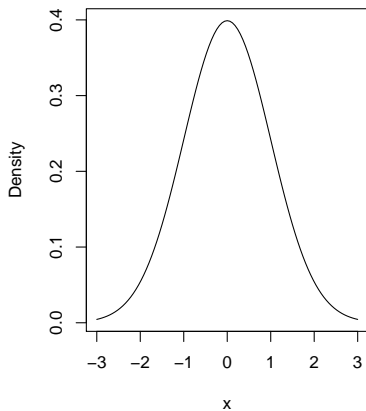
- ▶ `dump()`: save R objects in ascii format
- ▶ `source()`: load R objects saved with `dump()`
- ▶ `save()`: save R objects in binary format
- ▶ `load()`: load R objects saved with `save()`

```
# Try this  
dump(ls(), file = "objects.txt")  
source("objects.txt")
```

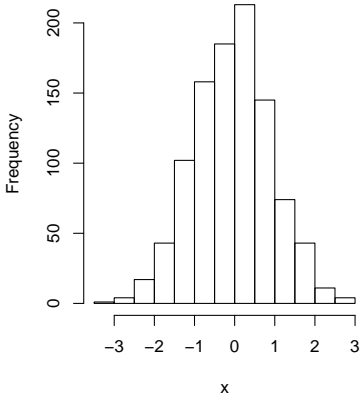
Probability distributions and sample simulation

Probability distribution and sample simulation

Density of normal distribution $N(0,1)$

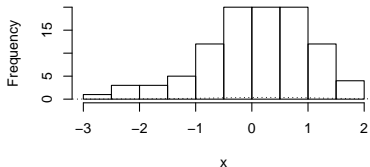


Distribution of 1000 random samples

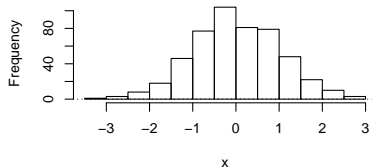


Probability distribution and sample simulation

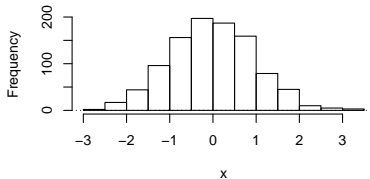
N = 100



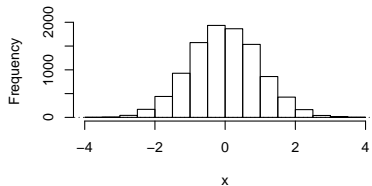
N = 500



N = 1000



N = 10000



Usual probability distributions

Distributions for **continuous** variables:

- ▶ **Normal** distribution: $X \rightsquigarrow N(\mu, \sigma)$:

$$f(x) = \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{1}{2\sigma^2}(x-\mu)^2}, \forall x \in \mathbb{R}.$$

- ▶ **Uniform** distribution: $X \rightsquigarrow U(a, b)$: if $x \in [a, b]$, $f(x) = \frac{1}{b-a}$, elsewhere $f(x) = 0$.

Distributions for **discrete** variables:

- ▶ **Poisson** distribution: $X \rightsquigarrow P(\lambda)$: $P(X = x) = e^{-\lambda} \frac{\lambda^x}{x!}, \forall x \in \mathbb{N}$.

- ▶ **Binomial** distribution: $X \rightsquigarrow B(n, p)$:
 $P(X = x) = C_n^x p^x (1-p)^{n-x}, \forall x \in [0, n]$.

Probability distributions in R

R can evaluate probabilistic quantities following usual probability distributions.

Usual distributions: `*norm()`, `*binom()`, `*chisq()`, `*unif()`, `*pois()`, `*t()`, `*exp()`, ...

* should be replaced by one of the following:

- ▶ p distribution function, $F(x) = P(X \leq x)$, use `pnorm()`, `pbinom()`, `pt()`, ...
- ▶ d density $P(X = x)$ or $f(x)$, use `dnorm()`, `dt()`, `dpois()`, ...
- ▶ q quantiles of order q , $\operatorname{argmin}_x \{P(X \leq x) > q\}$, use `qnorm()`, `qbinom()`, `qchisq()`, ...
- ▶ r sample simulation, use `runif()`, `rpois()`, ...

Probability distributions examples

```
dbinom(3, 10, 0.2)
```

```
## [1] 0.2013266
```

```
rbinom(10, 10, 0.2)
```

```
## [1] 0 3 2 3 3 1 3 2 2 3
```

```
pbinom(1, 10, 0.2)
```

```
## [1] 0.3758096
```

Probability distributions examples

```
pbinom(2, 10, 0.2)
```

```
## [1] 0.6777995
```

```
qbinom(0.5, 10, 0.2)
```

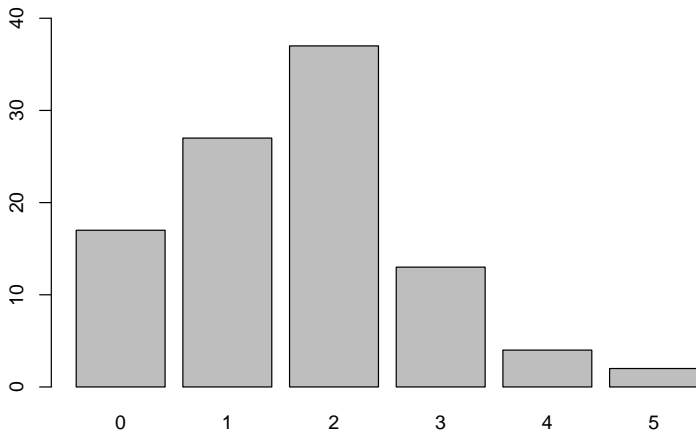
```
## [1] 2
```

```
qchisq(0.1, 8)
```

```
## [1] 3.489539
```

Probability distributions examples

```
x=rbinom(100, 5, 1/3)
par(mfrow = c(1,1))
barplot(table(x), ylim = c(0, 40))
```

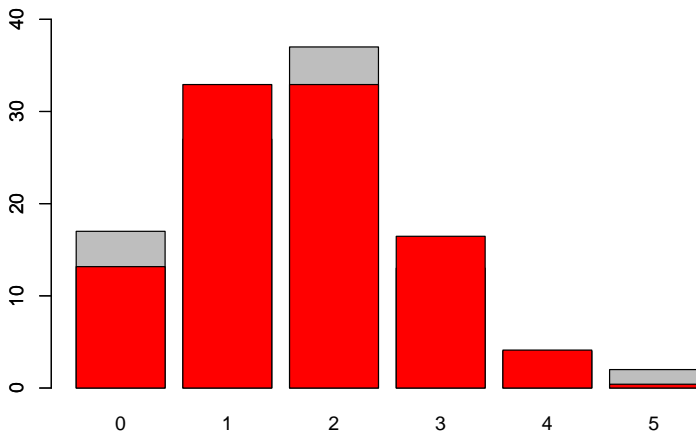


Probability distributions examples

```
# add a barplot to an existing barplot
```

```
barplot(table(x), ylim = c(0, 40))
```

```
barplot(dbinom(0:5, 5, 1/3)*100, add=T, col=2)
```



Descriptive statistics

Characteristics of a statistical serie

Let (x, y) be a pair of statistical series (vector or data.frame).

- ▶ `table(x)` returns the value counts of x (for x discrete or character).
- ▶ `summary(x)` returns a summary of descriptives statistics of x (minimum, 1st quartile, mean, median, 3rd quartile, maximum), for x numeric.
- ▶ `mean(x)`, `median(x)`, `var(x)`, `sd(x)` for mean, median, variance and standard-deviation (x numeric).
- ▶ `quantile(x, probs)` returns the quantiles of x (numeric) corresponding to parameter vector of probabilities (% of the population)/ Returns quantiles by default.
- ▶ `cor(x,y)`, `cov(x,y)` return the correlation/covariance matrix between x and y (numeric).

Characteristics of a statistical serie: examples

Try this

```
age = c(18, 15, 12, 16, 20, 17)
weight = c(55, 57, 46, 54, 60, 57)
name = c("a", "b", "c", "a")
table(weight); table(name)
summary(weight); summary(name)
mean(weight)
quantile(age)
quantile(weight)
cor(age, weight)
quantile(weight, probs = 0.25)
quantile(age, probs = c(0.1, 0.4))
```

Visualization of a statistical serie

- ▶ `hist(x)` plot the histogram of x .
- ▶ `density(x)` computes the kernel density estimator of x .
- ▶ `ecdf(x)` computes the empirical distribution function of x .
- ▶ `barplot(x)` bar chart of x (discrete).
- ▶ `stem(x)` tree of values of x (discrete).
- ▶ `boxplot(x)` boxplot of values of x (discrete).
- ▶ `qqnorm(x)` plot the quantiles of x in function of the quantiles of the normal distribution.
- ▶ `qqplot(x,y)` plot the quantiles of x in function of the quantiles of y .
- ▶ `plot(x)` plot the values of x .
- ▶ `plot(x,y)` scatter plot of coordinates (x,y) .

Examples: estimation of the distribution function using the empirical cumulative distribution function (`ecdf()`)

The **empirical distribution function** of a series of observations x_1, \dots, x_n is the **step function** between points $(x_i, \frac{i}{n})$, it is defined as follows:

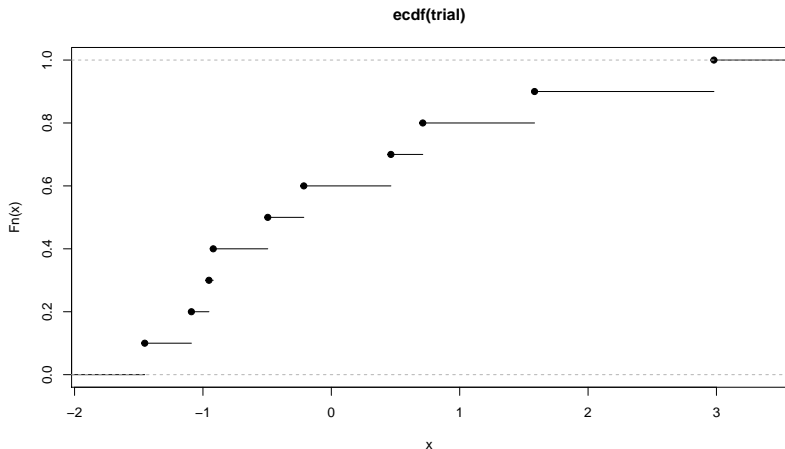
$$F_n(x) = \frac{\#\{i : x_i \leq x\}}{n} = \frac{1}{n} \sum_{i=1}^n \mathbb{I}_{\{x_i \leq x\}}$$

If x_1, \dots, x_n are realizations of random variable X , $F_n(x)$ is an estimation of the distribution function of X : $F(x) = P(X \leq x)$.

$\forall x, F_n(x) \rightarrow F$ when $n \rightarrow +\infty$.

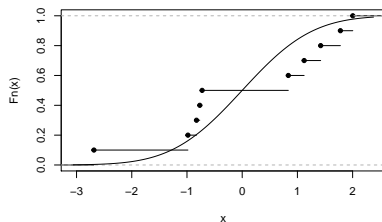
Examples: estimation of the distribution function of a probability distribution

```
trial = rnorm(10); plot(ecdf(trial))
```

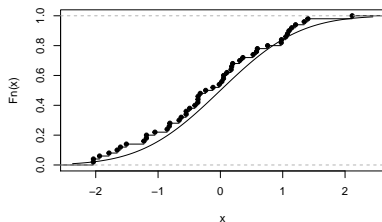


Examples: estimation of the distribution function of a probability distribution

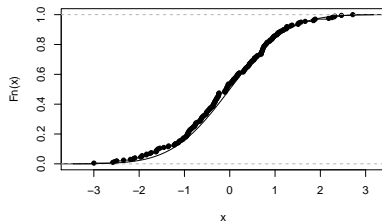
ECDF for $n = 10$



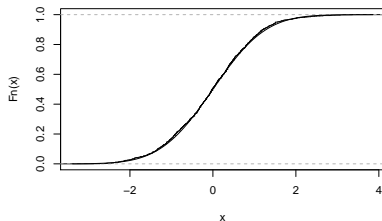
ECDF for $n = 50$



ECDF for $n = 200$



ECDF for $n = 1000$



Examples: estimation of a discrete probability distribution

For a discrete series x_1, \dots, x_n , consider the proportion of observations that take each value x of the series (equivalent to a bar plot).

$$p_n(x) = \frac{\#\{i : x_i = x\}}{n} = \frac{1}{n} \sum_{i=1}^n \mathbb{1}_{\{x_i=x\}}$$

If x_1, \dots, x_n are realizations of a discrete random variable X , $p_n(x)$ is an approximation of the probability distribution of X , $P(X = x)$.

The bar plot of the series estimates the graph of the probability distribution of X . Greater is n , better is the estimation.

Examples: estimation of a discrete probability distribution

```
trial = rbinom(10, 10, 0.3);  
t = table(trial); t
```

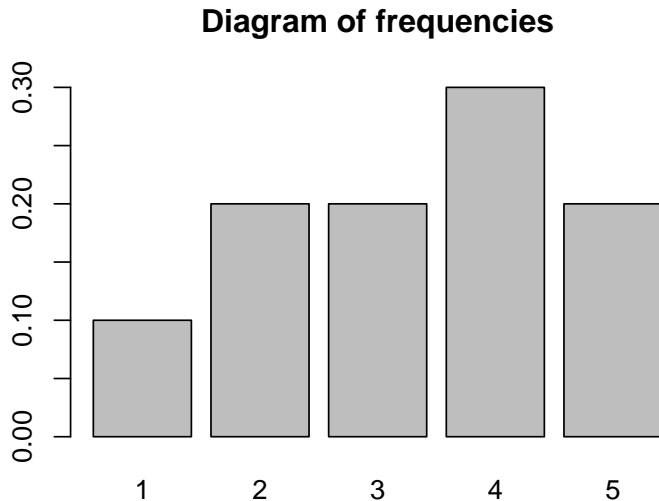
```
## trial  
## 1 2 3 4 5  
## 1 2 2 3 2
```

```
stem(trial)
```

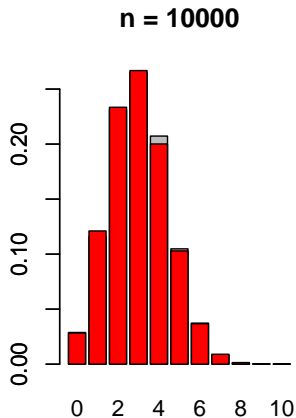
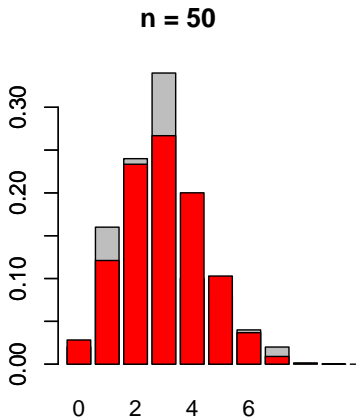
```
##  
## The decimal point is at the |  
##  
## 1 | 0  
## 2 | 00  
## 3 | 00  
## 4 | 000  
## 5 | 00
```


Examples: estimation of a discrete probability distribution

```
par(mfrow = c(1,1))  
barplot(t/length(trial), main = "Diagram of frequencies")
```



Examples: estimation of a discrete probability distribution



Examples: density estimation of a continuous probability distribution

The **density** f of a random variable X can be estimated with a **histogram**: let I be the interval of a series' values and $(I_j)_{j \leq k}$ a partition of I in k classes. The histogram based on this partition is the step function which is the proportion of observations in each class, normalized by class amplitude. Class amplitudes I_j should be adjusted to better match with the real distribution of observations.

$$\hat{f}_n(x) = \sum_{j=1}^k \left(\frac{\#\{i : x_i \in I_j\}}{n I_j} \right) 1_{\{x \in I_j\}}$$

In general, classes have the same amplitude, determined by the number of classes k (you can also use `cut()`).

Greater is n , more $\hat{f}_n(x)$ is similar to f .

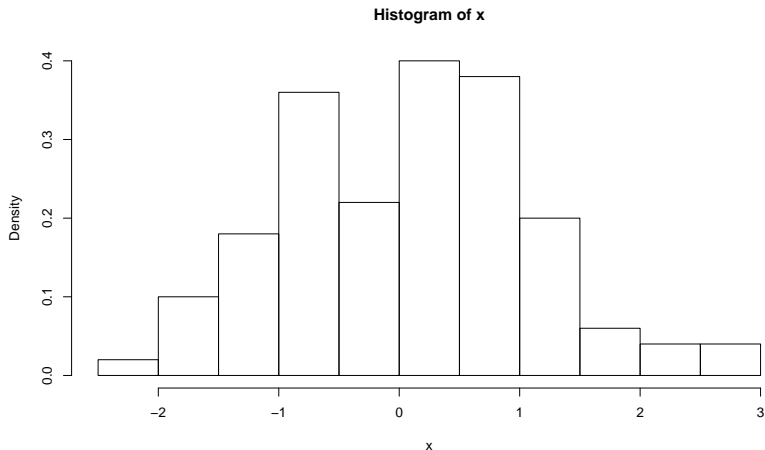
Examples: density estimation of a continuous probability distribution

```
hist(x, breaks, prob, right, col, main, xlab, ylab,  
...)
```

- ▶ `x` is a statistical series
- ▶ `breaks` is a vector of class breakpoints, or a number corresponding to $k + 1$, or a function to compute the number of classes. Default uses the Sturge rule ($k = 1 + 1.322 \log_{10} n$)
- ▶ `prob` returns the histogram of counts if `prob = FALSE`, returns the histogram of relative counts (frequencies) if `prob = TRUE`. Default is `FALSE`.
- ▶ `right` is `TRUE` if the classes of the histogram are right-closed and left-opened, default is `FALSE`
- ▶ `col` is the color
- ▶ `main`, `xlab` and `ylab` are the main title, x-axis title and y-axis title

Examples: density estimation of a continuous probability distribution

```
x = rnorm(100); hist(x, prob=T, breaks=12)
t = hist(x, prob = T, breaks = 12)
```



Examples: density estimation of a continuous probability distribution

```
names(t)
```

```
## [1] "breaks" "counts" "density" "mids" "xname" "equidist"
```

```
t$breaks
```

```
## [1] -2.5 -2.0 -1.5 -1.0 -0.5 0.0 0.5 1.0 1.5 2.0 2.5 3.0
```

```
t$density
```

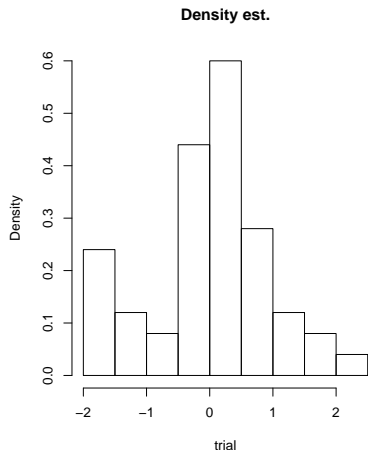
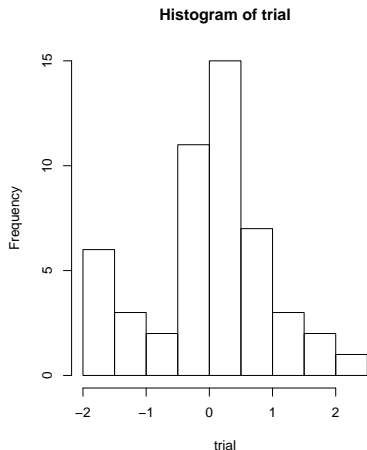
```
## [1] 0.02 0.10 0.18 0.36 0.22 0.40 0.38 0.20 0.06 0.04 0.04
```

```
t$counts
```

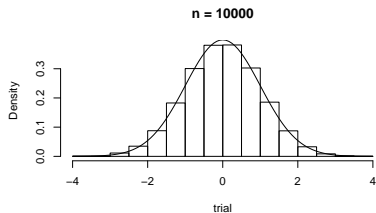
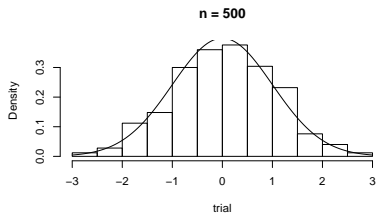
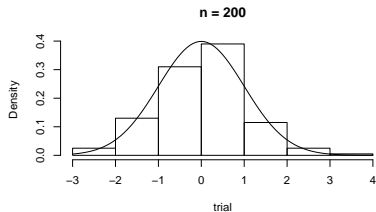
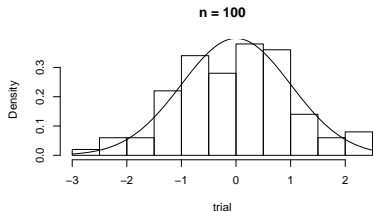
```
## [1] 1 5 9 18 11 20 19 10 3 2 2
```

Examples: density estimation of a continuous probability distribution

```
trial = rnorm(50); par(mfrow = c(1,2))  
hist(trial); hist(trial, prob=T, main="Density est.")
```



Examples: density estimation of a continuous probability distribution



Examples: density estimation of a continuous probability distribution

The **density** f of a random variable X can be estimated with a **kernel estimator**: the “derivative” of the empirical cumulative distribution function is a classical estimator of the distribution function. $\forall x \in I$, and $\epsilon > 0$ (a small value), the “derivative” is:

$$\frac{\hat{F}_n(x + \epsilon) - \hat{F}_n(x - \epsilon)}{2\epsilon} = \frac{\#\{i : x_i \in [-\epsilon x, \epsilon x]\}}{2n\epsilon} = \frac{1}{2n\epsilon} \sum_{i=1}^n 1_{-1 \leq \frac{x_i - x}{\epsilon} \leq 1}$$

This estimator assumes that the observations are uniformly drawn around each x_i . One can use a smoother distribution, of density K , with k is the smoothness parameter. We thus consider the following estimator:

$$\hat{f}_n(x) = \frac{1}{n\epsilon} \sum_{i=1}^n K\left(\frac{x - x_i}{\epsilon}\right)$$

Examples: density estimation of a continuous probability distribution

```
density(x, bw, kernel, ...)
```

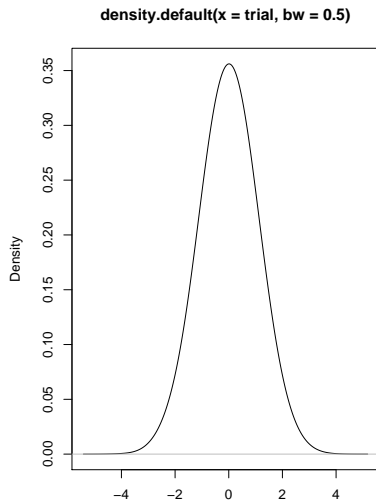
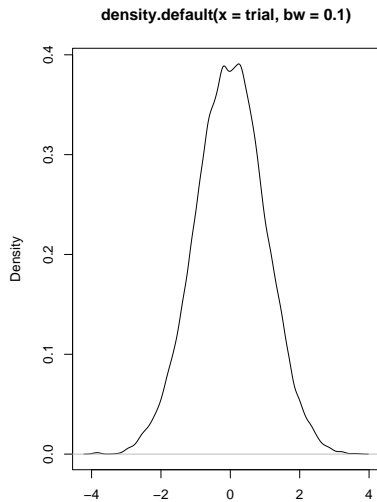
- ▶ `x` is a series
- ▶ `bw` is the size of the smoothing window. It is defined by a number. Default is automatic.
- ▶ `kernel` is a character string giving the kernel used (gaussian, rectangular, ...). Default is gaussian.

The output of `density()` corresponds to descriptive statistics of x and the density estimated on values of x .

Use `plot(density())` to plot the density curve.

Examples: density estimation of a continuous probability distribution

The smoothing window is important, greater it is, smoother is the estimation. Lower it is, noisier is the estimation.



Examples: density estimation

```
density(trial)
```

```
##
```

```
## Call:
```

```
## density.default(x = trial)
```

```
##
```

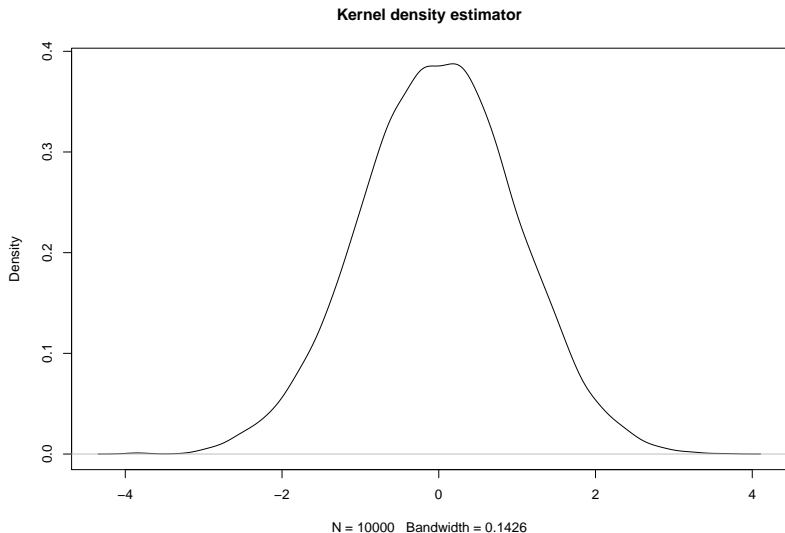
```
## Data: trial (10000 obs.); Bandwidth 'bw' = 0.1426
```

```
##
```

##	x	y
##	Min. :-4.3469	Min. :0.0000033
##	1st Qu.:-2.2336	1st Qu.:0.0021840
##	Median :-0.1204	Median :0.0439594
##	Mean :-0.1204	Mean :0.1181849
##	3rd Qu.: 1.9929	3rd Qu.:0.2254036
##	Max. : 4.1062	Max. :0.3875813

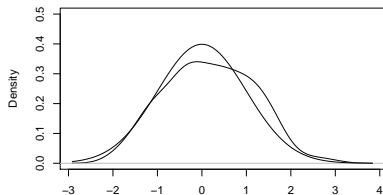
Examples: density estimation

```
plot(density(trial), main="Kernel density estimator")
```



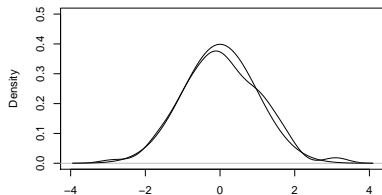
Examples: density estimation

n = 100



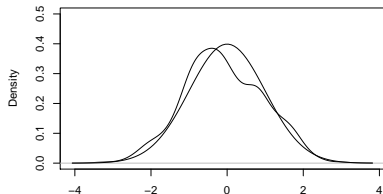
N = 100 Bandwidth = 0.3533

n = 200



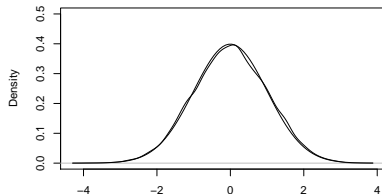
N = 200 Bandwidth = 0.3253

n = 500



N = 500 Bandwidth = 0.269

n = 10000



N = 10000 Bandwidth = 0.1443

Examples: characteristics of a series

`boxplot()` plots a **boxplot**, which aims to visualize the characteristics of a series (outliers, symmetry, dispersion).

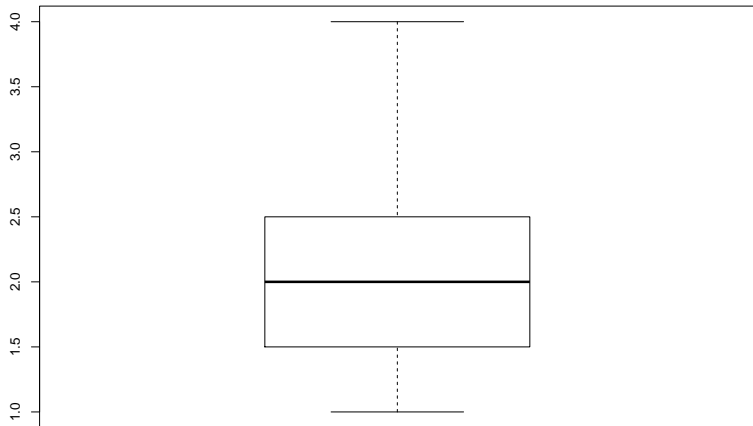
`qqnorm()` aims to compare a series' distribution to a standard gaussian distribution.

```
x=c(1,1,2,2,2,3,4); summary(x)
```

##	Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
##	1.000	1.500	2.000	2.143	2.500	4.000

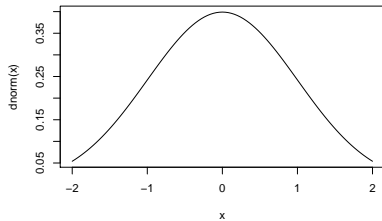
Examples: characteristics of a series

```
boxplot(x)
```

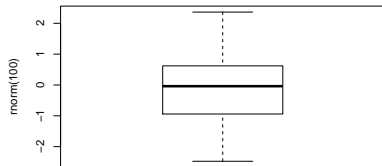


Examples: characteristics of a series

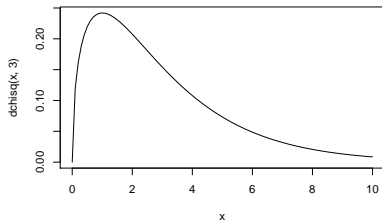
Density plot



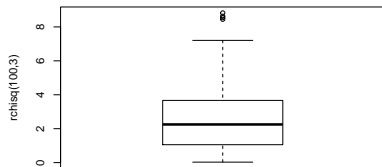
Boxplot



Density plot

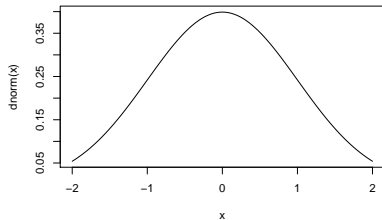


Boxplot

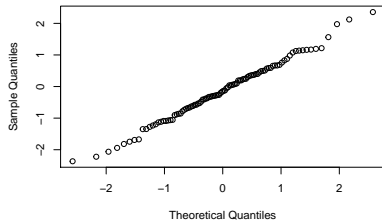


Examples: characteristics of a series

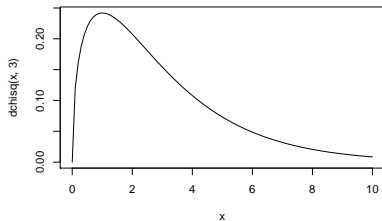
Density plot



Normal Q-Q Plot



Density plot



Normal Q-Q Plot

