## The Mean: Takeaways 🖻

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## **Syntax**

• Generating randomly a distribution of integers:

```
set.seed(1)
distribution <- sample.int(value_max, size_of_distribution)</pre>
```

• Computing the mean of any numerical vector using R base functions sum() and length():

```
mean <- sum(vector) / length(vector)</pre>
```

• Computing the mean of any numerical vector using R base function mean():

```
mean <- mean(vector)</pre>
```

• Creating your own compute\_mean() function from algebraical definition of mean:

```
compute_mean <- function(distribution) {
    N <- length(distribution)
    sum_of_the_distribution = 0
    for ( i in 1:N) {
        sum_of_the_distribution <- sum_of_the_distribution + distribution[i]
    }
    sum_of_the_distribution / N
}</pre>
```

• Checking if the mean is equidistant from below and above values:

```
round(sum(distribution - mean)) == 0
```

• Reading a TSV (tab-separated value) dataset:

```
library(readr)

data <- read_tsv(file_name)</pre>
```

• Generating a scatter plot to represent visually how the sampling error changes as the sample size increases:

• Generating a histogram to represent visually the distribution of sample means:

## **Concepts**

- We can summarize the distribution of a numerical variable by computing its **mean**.
- The mean is a single value and is the result of taking into account **equally** each value in the distribution.
- The mean is **the balance point** of a distribution the total distance of the values below the mean is equal to the total distance of the values above the mean.
- The mean of a population can be defined algebraically in several equivalent ways:

• The mean of a sample can be defined algebraically in several equivalent ways:

 $\bullet\,$  The sample mean  $\,$  is an unbiased estimator for the population mean  $\,$  .

## Resources

- The Wikipedia entry on the mean.
- Useful documentation:
  - sample.int()
  - <u>mean()</u>
  - replicate()



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