

# MY STUDY

My name

23/10/2020

# SECTION1

## SUBSECTION 1.1

# SLIDE 1

The objectives of the current study are. . .

# SLIDE 1

The objectives of the current study are. . .

Write a bit more about it. . .

# SLIDE 2

- Item 1

# SLIDE 2

- Item 1
- Item 2

# SLIDE 2

- Item 1
- Item 2
- Item 3



## SUBSECTION 1.2

## SLIDE 3

$$x^n + y^n = z^n$$

## SLIDE 3

$$x^n + y^n = z^n$$

$$E = mc^2$$

## SECTION 2

## SUBSECTION 2.1

# SLIDE 4

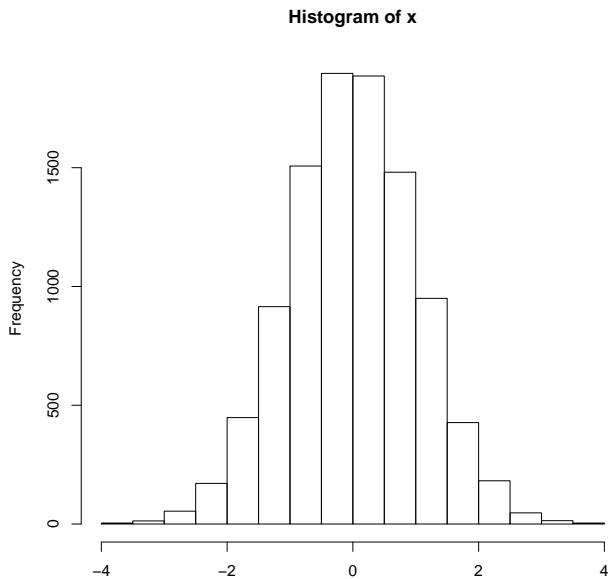
## **Title 1**

This is important, because. . .

## **Title 2**

This is a minor issue, due to. . .

## SLIDE 5



## SUBSECTION 2.2



## SLIDE 6



FIGURE 1: This is the caption

# SLIDE 7

## REMARK

Sample text

## IMPORTANT THEOREM

Sample text in red box

## EXAMPLES

Sample text in green box. The title of the block is "Examples".

## SECTION 3

## SUBSECTION 3.1

# SLIDE 8

Step 1

# SLIDE 8

Step 1



Step 2

# SLIDE 8

Step 1



Step 2



Step 3

## SLIDE 8

Step 1



Step 2



Step 3



Step 4



# SLIDE 9

Seminal work done:

- Study 1 <sup>1</sup>
- Study 2 <sup>2</sup>
- Study 3 <sup>3</sup>

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<sup>1</sup>“Gauss's Derivation of the Normal Distribution and the Method of Least Squares, 1809” (n.d.)

<sup>2</sup>WATSON & CRICK (1953)

<sup>3</sup>Li & Dewey (2011)

## SUBSECTION 3.2

# SLIDE 10



# SLIDE 11

**Thank you for your attention!**

# REFERENCES I

1. in. *A History of Parametric Statistical Inference from Bernoulli to Fisher, 1713–1935* 55–61 (Springer New York).  
[https://doi.org/10.1007%2F978-0-387-46409-1\\_7](https://doi.org/10.1007%2F978-0-387-46409-1_7).
2. Li, B. & Dewey, C. N. RSEM: accurate transcript quantification from RNA-Seq data with or without a reference genome. *BMC Bioinformatics* **12**. <https://doi.org/10.1186%2F1471-2105-12-323> (Aug. 2011).
3. WATSON, J. D. & CRICK, F. H. C. Molecular Structure of Nucleic Acids: A Structure for Deoxyribose Nucleic Acid. *Nature* **171**, 737–738. <https://doi.org/10.1038%2F171737a0> (Apr. 1953).