



# Network Science applied to Epilepsy

*BioBytes Group*

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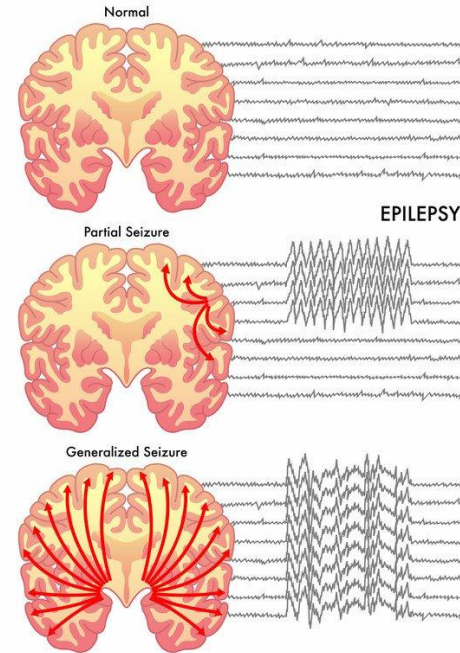
# Introduction



What is **epilepsy**<sup>[1]</sup>?

- Epilepsy is a disease caused by the **synchronous, excessive or abnormal** activity of neuronal cells in the brain.
- Multiple causes.
- 50 million people/Brazil = 2% <sup>[2]</sup>

[2] - World Health Organization



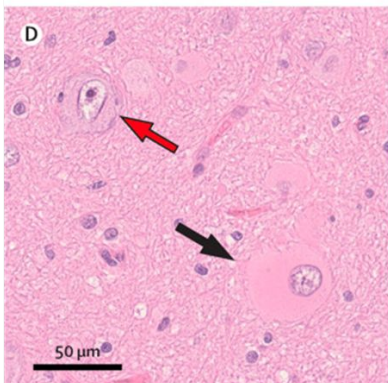
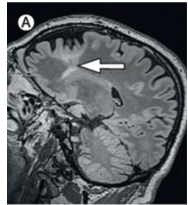
Normal Brain (on top) and Brain with epilepsy (below) <sup>[2]</sup>

# Introduction



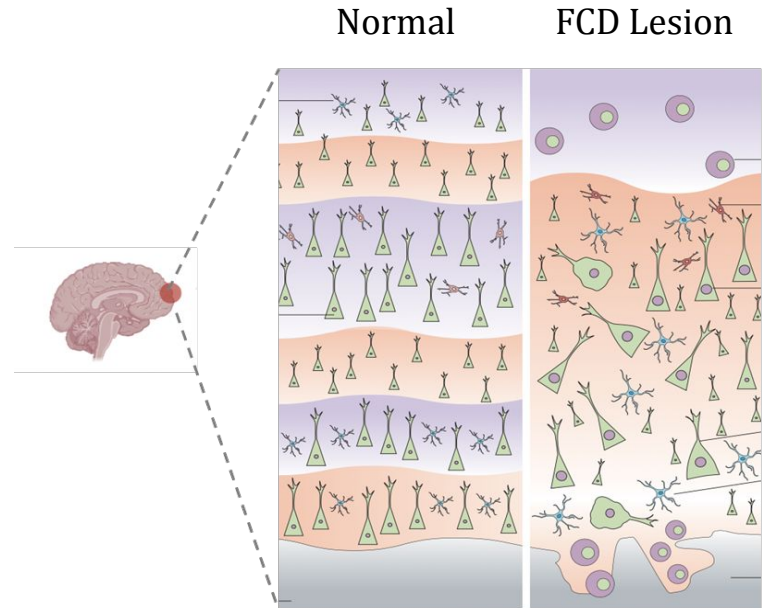
- Focal Cortical Dysplasia

- Pediatric epilepsy
- Malformation of cortical development
- Cortical and cellular abnormalities



Baulac et al, Lancet Neur., 2021

**Dysmorphic  
Neurons  
+  
Balloon Cells**

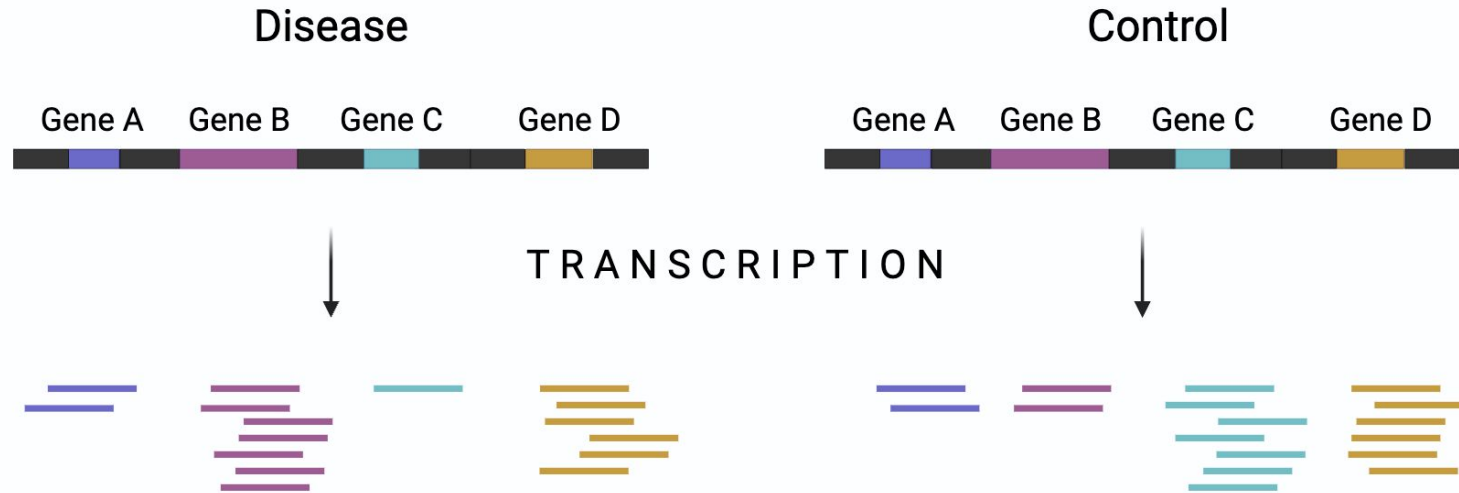


Sisodiya et al., Lancet Neur., 2009

# Transcriptomics



- Quantifies RNA expression by number of transcript counts

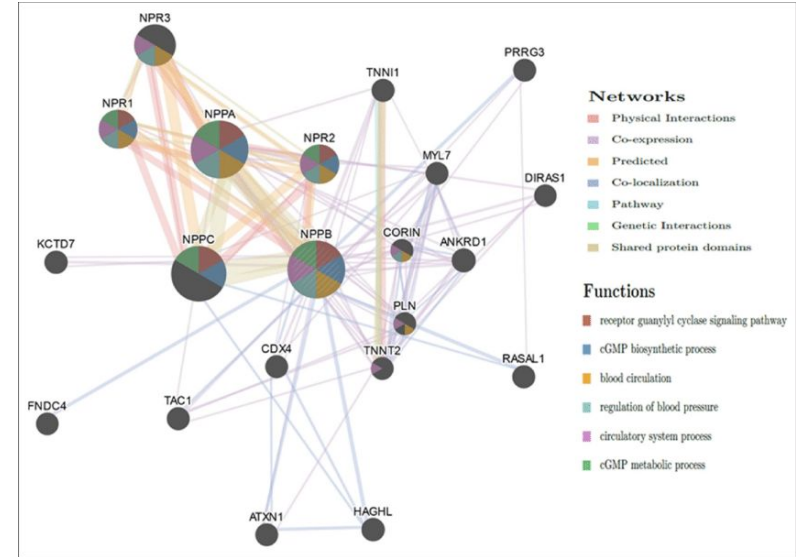


Genes B and C = Differentially Expressed Genes (DEGs)



# Proposal

- Apply **network science methods** in the analysis of differentially expressed genes in patients diagnosed with **epilepsy** caused by Focal Cortical Dysplasia type 2b (FCD 2b)
- Analyze the correlation between **genes** and **biological functions**



Schematic illustration of the gene network and biological functions involving the NPPB gene<sup>[4]</sup>



# Research Questions

- What are the differentially expressed genes between **disease** and **control group**?
- Which **biological functions** are most represented in the disease?  
→ **functional networks**
- How are these genes being regulated? → **regulation networks**

# Methodology



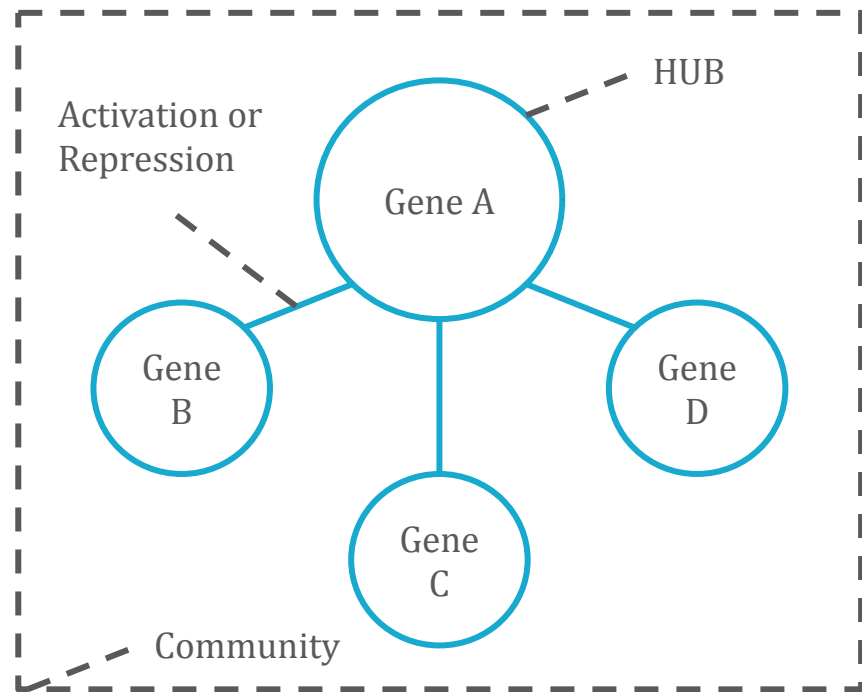
- Determine gene expression difference using statistical methods of **DESeq2**<sup>[5]</sup> package of *Bioconductor Software*
  - Models read counts using a negative binomial distribution
- Create a network of interactions with the identified **differentially expressed genes**
- Integrate the network with functional annotation information, using *Gene Ontology*<sup>[6]</sup>, and relate them to their respective **biological functions** and **signaling pathways** that are involved, resulting in a functional enrichment analysis



# Methodology



- Use Network Science techniques to explore the organization and dynamics of genes in the network:
  - **Centrality analysis:** identify genes that play important roles in the functioning of epilepsy
  - **Community detection:** revealing groups of genes that strongly interact with each other
  - **Perturbation and robustness analysis:** assess how the removal or perturbation of specific genes affects network structure and function





# Softwares and Tools

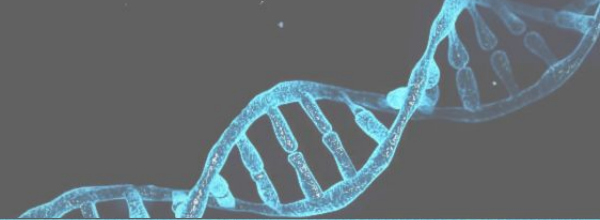
- **DESeq2**<sup>[5]</sup>: Differential expression tests



- **Cytoscape**<sup>[7]</sup>:
  - Generation of a network of differentially expressed genes with associated biological functions
  - Analysis using Network Science techniques

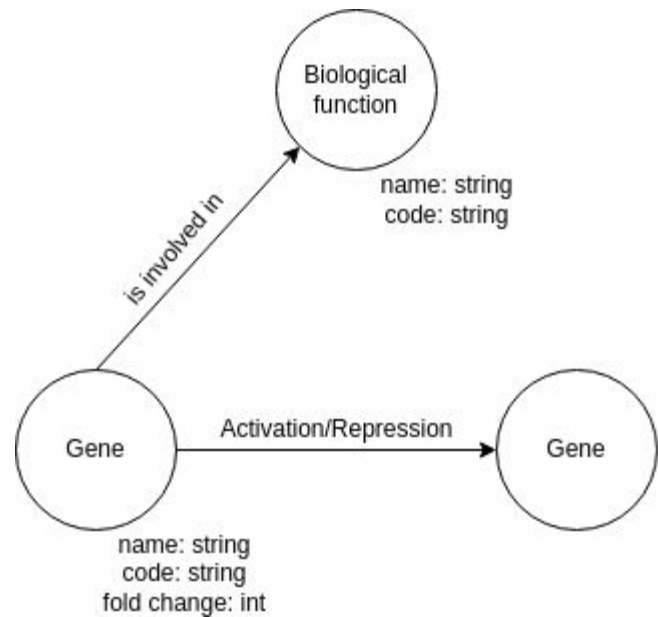


# Data Base



- Transcriptomes distinguish human FCD subtypes (NCBI)<sup>[8]</sup>:
  - Format: CSV (data) and XLSX (metadata)
  - Size: 5.5 Mb (data) and 5 Mb (metadata)
  - Row represents genes and columns represent samples
- Gene Ontology<sup>[6]</sup>:
  - Formats: OBO and OWL
  - Computational representation of the current scientific knowledge about the functions of genes from many different organisms
    - Functions of genes: the protein and non-coding RNA molecules produced by genes

# Logic Model



An abstract graphic in the top-left corner featuring a network of white nodes connected by thin white lines, set against a dark blue background with a subtle bokeh effect.

Thank You!  
Questions?

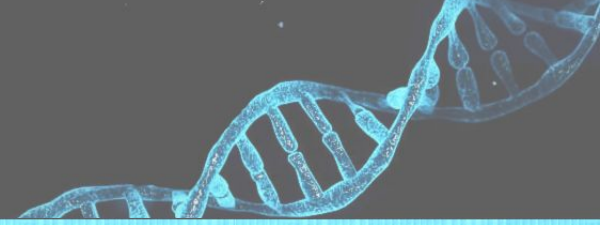
# References



- [1] - Patel P, Moshé SL. The evolution of the concepts of seizures and epilepsy: What's in a name? *Epilepsia Open*. 2020 Jan 10;5(1):22-35. doi: 10.1002/epi4.12375. PMID: 32140641; PMCID: PMC7049807.
- [2] - World Health Organization: <https://www.who.int/news-room/fact-sheets/detail/epilepsy/>
- [3] - Drugs.com: <https://www.drugs.com/health-guide/seizure.html>
- [4] - Associations of B-Type Natriuretic Peptide and Its Coding Gene Promoter Methylation With Functional Outcome of Acute Ischemic Stroke: A Mediation Analysis - DOI:10.1161/JAHA.120.017499
- [5] - DESeq2 (Bioconductor Software Package): <https://bioconductor.org/packages/release/bioc/html/DESeq2.html>



# References



- [6] - Gene Ontology: <https://geneontology.org/>
- [7] - Cytoscape: <https://cytoscape.org/>
- [8] - Transcriptomes distinguish human FCD subtypes:  
<https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE128300>
- [9] - Brain image theme: Getty Images (Yuichiro Chino)
- [10] - DNA image theme: iStock (Shutter2U)