Network Science applied to Epilepsy

BioBytes Group

MO413 1º semester of 2024 • Institute of Computing • UNICAMP

Carnot Luiz B. G. Filho

Laíz Luiza M. de Oliveira ra200744

Isabella C. Galvão

Ana Carolina da Hora ra272499 João Paulo F. da Silva

Vitoria D. M. Pinho

18/06/2024

Introduction

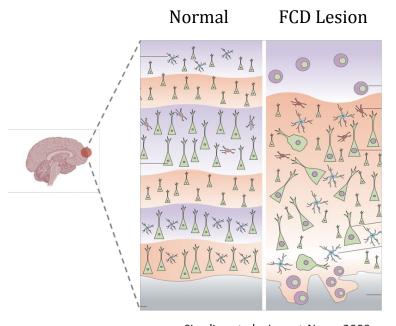


Epilepsy

 Epilepsy is a disease caused by the synchronous, excessive or abnormal activity of neuronal cells in the brain.

Focal Cortical Dysplasia

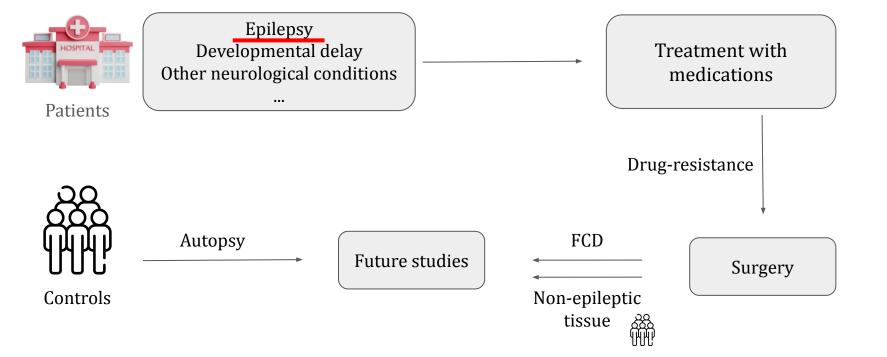
- Pediatric epilepsy
- Malformation of cortical development
- Cortical and cellular abnormalities



Sisodiya et al., Lancet Neur., 2009

Samples



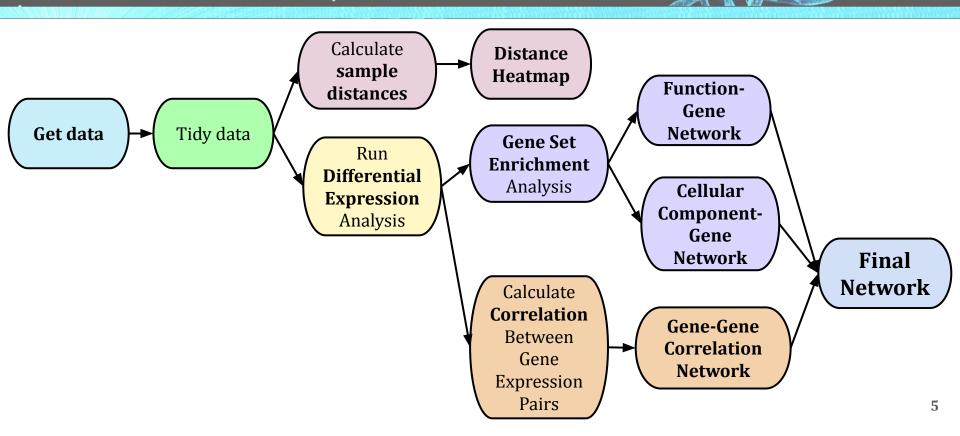


Proposal and Research Questions

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

- What are the differentially expressed genes between disease and control group?
- Which biological functions/cellular components are most represented in the disease? → functional networks
- How are these genes being regulated? → regulation networks
- What are the most important genes, in the network, for FCD IIb, and how do they relate to the change in gene expression between the two groups?

Workflow



Get Data



- RNA-seq data
- Expression matrix
- n = 5 FCD IIb (collected during epilepsy surgery)
- n = 8 control (Autopsy + non-epileptic patients)
- Metadata obtained from the authors (.xlsx)



https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE1283 00

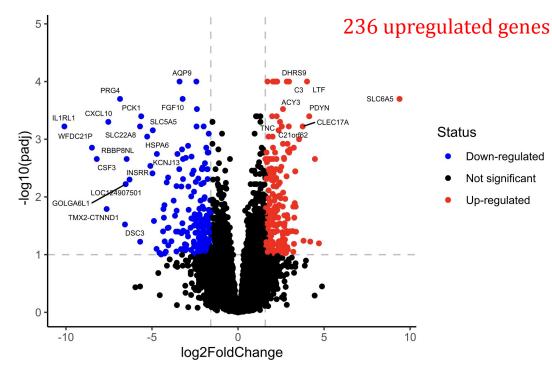
Differential Expression Analysis

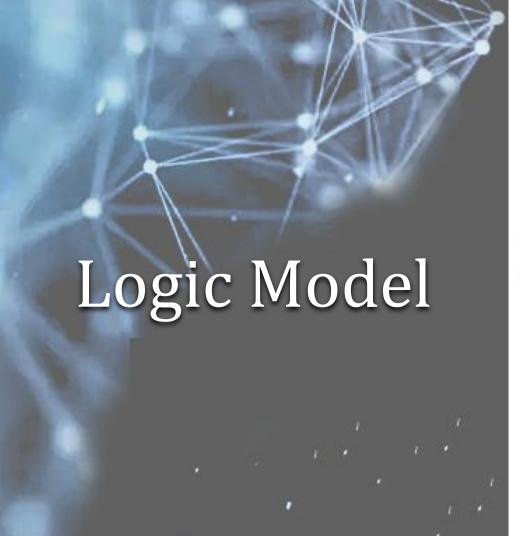


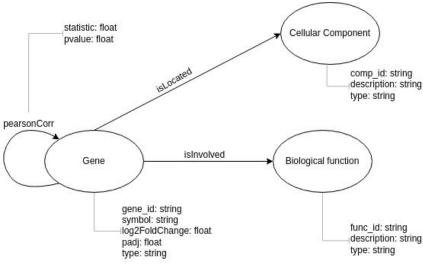


Criteria:

- FC > 3
- padj < 0.1







Gene Set Enrichment Analysis

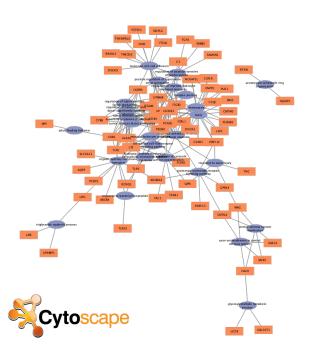


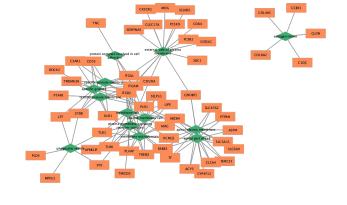




ClusterProfiler

- EnrichGO
 - o ont = "BP"
 - o ont = "CC"





Function-Gene Network

Component-Gene Network

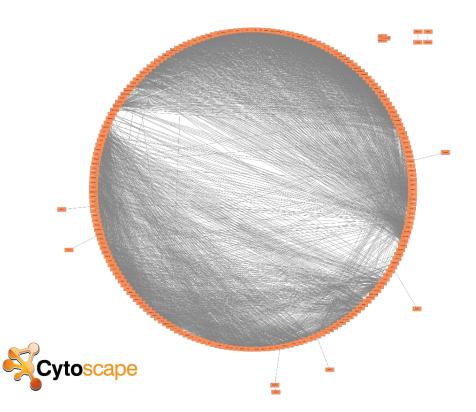
Correlation: Gene-Gene Network



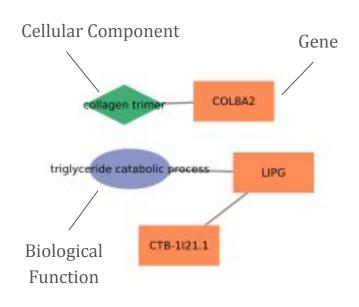


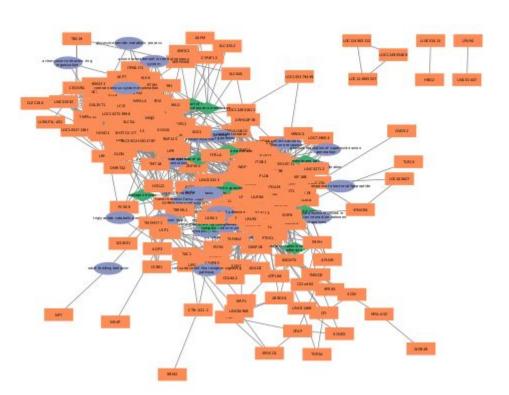
Pearson Correlation

- P < 0.05
- statistic > 0.85



Final Network









Number of gene nodes	236
Number of function nodes	26
Number of cellular component nodes	13
Number of gene-gene edges	3650
Number of gene-function edges	210
Number of gene-cc edges	130

Number of nodes and edges

Av. deg. of genes	32.37
Av. deg. of functions	8.08
Av. deg. of cellular components	10.00

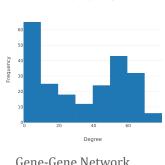
Average degree of genes, functions and cellular components





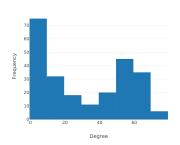
Statistical Analysis





Frequency of Degree

Gene-Gene Network

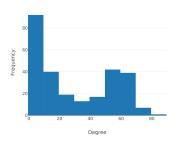


Frequency of Degree

Frequency of Degree Degree

Function-Gene Network

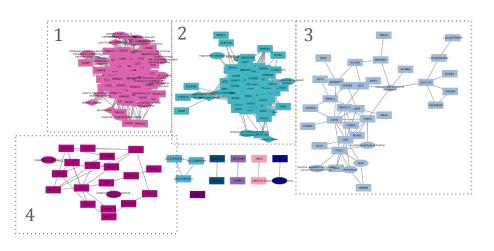
Frequency of Degree



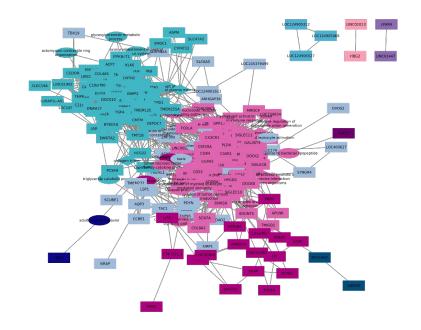
Component-Gene Network

Complete-Gene Network

Communities Analysis: Leading EigenVector



Complete Network 12 communities



Communities Analysis: Leading EigenVector

Name	Degree	Comm
membrane microdomain	12	1
apical part of cell	15	2
external side of plasma membrane	15	4

Components with the highest degrees of each community

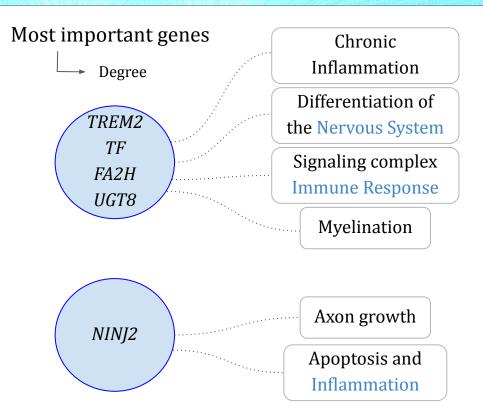
TREM2	82	1
TF	74	2
F2RL1	22	3
LTF	22	4

Name	Degree	Comm
leukocyte cell-cell adhesion	15	1
axon ensheathment in central nervous system	5	2
taxis	18	3
organic hydroxy compound transport	10	4

Functions with the highest degrees of each community

Most important elements



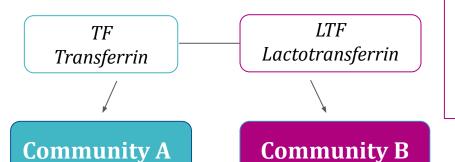


Most important functions	Degree
Chemotaxis	18
Leukocyte cell-cell adhesion	15
Myeloid leukocyte activation	14
Tumor necrosis factor production	12
Organic hydroxy compound transport	10

Most Important Component	Degree
External side of plasma membrane	15
Apical plasma membrane	15
Apical part of cell	15
Membrane raft	12
Membrane microdomain	12

Community Analysis

Iron capture for myelin sheath maintenance



Transferrin family

- Anti-inflammatory activity
- Regulation of cellular growth and differentiation
- protection against reactive oxygen species

- Axon ensheathment in central nervous system
- Central nervous system myelination

- Response to axon injury
- Organic hydroxy compound transport

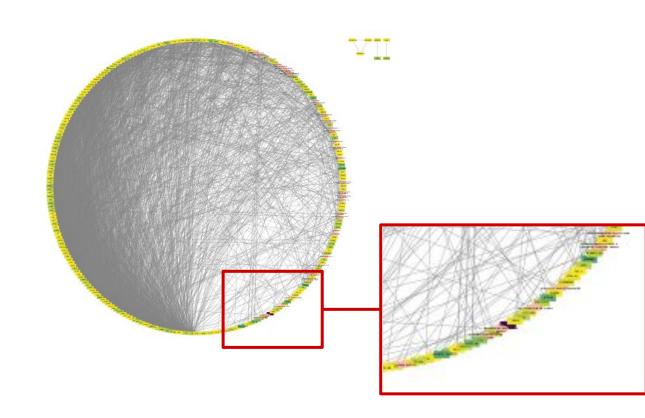
Fold-Change vs Degree

Network important genes x Gene Expression

Highest degree	Highest Fold Change
TREM2	SL6A5
TF	SERPINA5
NINJ2	LOC105379499
FA2H	LOC124901612
UGT8	ATP1B4

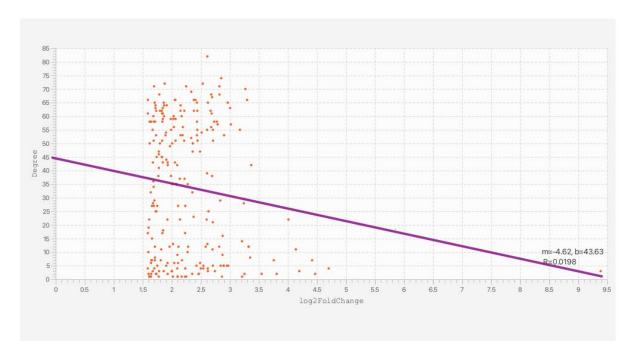
Fold-Change vs Degree

Network important genes x Gene Expression



Fold-Change vs Degree

Network important genes x GeneExpression



Conclusions

Biological findings

- We found 236 up-regulated genes (FC > 3; padj < 0.1), some of which have been classically associated with epilepsy/cortical malformations (e.g. CCBE1, ASPM) and some that had not been reported in the literature for the disease (e.g. NINJ2/LTF).
- Most of the biological functions overrepresented in the disease are linked to immune system activation/inflammation, which is in line with more recent findings in FCD.

Conclusions

Difficulties

- Networks assembling via Cytoscape (genes + functions + components)
 - It was necessary to create a new column to differentiate the type of node
- Robustness Analysis and Link Prediction with no clear biological effect
 - There was no biological interpretation for these kinds of analysis
- Community analysis using Cytoscape

Future Directions

- Experimental validation
- Exploratory study of disease-related genes not found in the literature
- Use of new technologies in this study



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.
- [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)