Application of Graph Theory on Neurological Pathways and Cognitive Processes

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Abstract—This study looks at how brain areas can be represented as nodes and their connections as edges in order to understand neural pathways in cognitive processes using graph theory. Investigated are the theoretical foundations of brain network representation, analyzing weighted, directed, and undirected graph forms. In order to characterize the organization of brain networks, important graph-theoretic metrics such as degree centrality, clustering coefficient, and small-world features are investigated. Preprocessing, graph creation, and network analysis techniques are included in implementation approaches utilizing fMRI and DTI neuroimaging data. Applications show how to use changes in network properties to recognize neurological conditions like schizophrenia and Alzheimer's disease.

Keywords—Neural networks, Graph theory, brain connectivity

I. INTRODUCTION

One of the most complex networks in nature is the human brain. Consisting of approximately 86 billion neurons connected through trillions of synaptic connections, it is the central organ of the nervous system which orchestrates every cognitive process that regulates our body. Understanding the intricate mechanisms of how these neural pathways enable cognitive processes such as learning, feeling and memory formation is a central question in neuroscience. A conventional approach in neuroscience to study neural connectivity has been through relying on biological and statistical methods.

Complex networks can be well represented and analyzed using graph theory, an essential area of discrete mathematics. Graph theory is a strong technique in neuroscience that may be used to measure network features, predict patterns of brain connectivity, and detect changes linked to various cognitive states and neurological illnesses. Our understanding of how brain networks support cognitive activities and how these networks are damaged in various pathological situations has been completely transformed by the application of graph-theoretic techniques to neuroscience.

For a while, neuroscience research focused on understanding the functions of individual brain regions. But

it's becoming more and more clear that intricate cognitive processes like memory, attention, language, and awareness result from the coordinated activity and dynamic interaction of various brain regions rather than from the isolated activities of individual parts. This realization spurred a fundamental shift in perspective: from merely identifying "where" brain activity occurs to understanding "how" different brain regions communicate and integrate information. Graph theory offered the precise language to formalize this inherent network structure, allowing researchers to move beyond simple activation maps and quantitatively analyze the complex relationships, information flow, and organizational principles that underpin all aspects of brain function. This capability became crucial for unraveling the brain's remarkable computational power and its vulnerabilities in disease.

This paper aims to provide a comprehensive examination of how graph theory can be applied to model neural pathways in cognitive processes by investigating the theoretical foundations that allow brain networks to be represented as graphs.

II. THEORETICAL FRAMEWORKS

A. Graph

Graph is a fundamental mathematical structure used to model relationships between objects. It is a visual representation which consists of vertices or nodes and edges or lines (citation). Vertices are individual objects or points while edges represent the relationships or connections between the vertices. Hence, an edge connects two vertices. A graph G is defined as an ordered pair G=(V, E) where V is a set of vertices and E is a set of edges.

Undirected Graph

Undirected graph has an edge between two vertices which has a symmetrical relationship. If vertice u is connected to vertice , then v is also connected to u. Undirected graphs are typically used when the connection or relationship between two neural elements is believed to be symmetrical.



Figure 2.1 Undirected Graph (a) and Directed Graph (b)

source : researchgate.net

Directed Graph

In a directed graph (diagraph), the edges have a specific direction. Directed graphs are essential for modeling the directed flow of information in neural pathways or influences between brain regions. Directed graphs are a direct representation of synaptic connections. Directed edges give understanding on feedback in cognitive processing.

Weighted Graph

In a weighted graph, each edge has an assigned numerical value called a weight w: $E \rightarrow \mathbb{R}$ which can represent numerous of things. A weighted graph is defined as G = (V, E, w) where

- V is a set of vertices (nodes)
- E is a set of edges
- w is a weight function



Figure 2.2 Weighted Graph

source : stackoverflow.com

The brain itself is not merely a collection of binary connections. The strength of these connections varies and are occasionally crucial for modulating information processing. Weighted graphs fit to represent this quantitative aspect. The assigned weights to the edges connecting neurons represent the strength of the synaptic connections and can be measured in different ways. These weights can coincide with the number of receptors or the amount of neurotransmitters that are released. In DTI-derived structural networks, the weights on the edges represent the number of white matter streamlines connecting two regions.

B. Brain as a Network

The complexity of the human brain naturally allows networks-based analysis, where graph theory gives a mathematical framework for comprehending brain organization and function, to be a natural fit for this complex structure. Early neuroanatomical research that demonstrated the interconnection of brain areas led to the idea of the brain as a network. Ramon y Cajal laid the foundation for contemporary network neuroscience with his neuron hypothesis in the late 19th century, which postulated that the nervous system is made up of distinct cellular units (neurons) connected by synapses. Building upon this foundational understanding of neurons as discrete, interconnected units, the introduction of modern neuroimaging techniques, such as functional Magnetic Resonance Imaging (fMRI) and Diffusion Tensor Imaging (DTI), enabled researchers to effectively map brain connections at a macroscopic scale. This technological advancement sparked the realization that higher-order cognitive functions do not arise from isolated brain regions but rather from the intricate and dynamic interaction among distributed neural populations. Therefore, graph theory became an essential instrument, offering the challenging mathematical language to translate these complex biological connections into measurable nodes and edges. This allowed for the methodical examination of the topology, efficiency, and modularity of brain networks as well as the functions of particular areas as network hubs, going beyond simple localization to a comprehensive comprehension of brain function.



Figure 2.3 Brain Network

source : researchgate.net

C. Graph theory in Neuroscience

Several graph-theoretic metrics have proven particularly valuable for characterizing brain networks:

1. Degree Centrality

The degree of a node represents the number of connections it has with other nodes. In brain networks, nodes with high degree centrality $C_d(i)$ are considered "hubs" that have important roles in integrating information and network communication.

$$C_d(i) = k_i = \sum_{j \neq i} a_{ij}$$

2. Clustering Coefficient

This indicator reflects local network organization and functional specialization within brain areas and predicts the tendency of a node's neighbors to be connected to one another.

3. Path Length

The fewest number of edges needed to connect two nodes is represented by their shortest path length L. The average path length throughout the network shows how well information is transferred throughout the brain.

4. Betweenness Centrality

This measure identifies nodes that frequently appear on shortest paths between nodes j and h that pass through i, which highlights brain regions that act as critical bridges in network communication.

$$C_{b}(i) = \frac{2}{(N-1)(N-2)} \sum_{i \neq h \neq i} \frac{n_{hj}(i)}{n_{hj}}$$

5. Small-World Properties

Brain networks typically exhibit small-world characteristics, combining high local clustering with short global path lengths. This organization enables both specialized local processing and efficient global integration. A network has small-world properties if it possesses:

- 1. High Clustering Coefficient: Similar to a regular (lattice-like) network, indicating local specialization and efficient processing within modules.
- 2. Short Characteristic Path Length: Similar to that of a random network, indicating efficient global communication and integration of information across distant regions.

Three criterias which are used to determine if a network has small-world properties :

$$\gamma = \frac{C}{C_{rand}}$$
 $\lambda = \frac{L}{L_{rand}}$ $\sigma = \frac{\gamma}{\lambda}$

6. Modularity

This parameter measures how well a network can be separated into distinct communities or modules, reflecting the brain's hierarchical structure into functional systems.

In the brain, modularity reflects the existence of functionally specialized processing units (e.g., visual processing module, auditory processing module) that operate relatively independently but are still able to communicate when needed. Dynamic changes in modularity can reflect cognitive state changes.

III. IMPLEMENTATION

A. Data Acquisition and Preprocessing

1. Functional Magnetic Resonance Imaging (fMRI)

fMRI is primarily used to deduce functional connectivity. The use of fMRI started in the mid-1990s which increased the discovery of bases in neurological disorders. fMRI measures Blood Oxygen level dependant (BOLD) signal, which is a mark of neural activity. BOLD time series are extracted from predefined brain regions (nodes) and are acquired using echo-planar imaging sequences with repetition times of 1-3 seconds. Nodes represent brain regions and edges represent relations or connections. In order to form a complex networks from fMRI in graph, preprocessing such as realignment, slice timing correction and normalization are needed.

There are two categories of computational methods for brain connectivity. Effective connection and functional connectivity. While effective connection focuses on the directed influence of brain regions on one another, functional connectivity gives information about the statistical dependency or temporal correlations between separate neurophysiological processes.

2. MRI

Diffusion MRI (dMRI) data, which measures the diffusion of water molecules in brain tissue, is the main source of information used to build structural brain networks. This technique leverages the anisotropic diffusion of water, meaning its movement is restricted and directed along the organized axonal bundles of white matter, thereby providing a unique window into the brain's physical pathways. Before network construction, the raw dMRI data undergoes crucial preprocessing steps, including correction for head motion, eddy current distortions, and removal of non-brain tissue, ensuring the accuracy and reliability of subsequent fiber tracking algorithms that will reconstruct the white matter tracts.

B. Graph Construction Representing Brain Networks

In order to apply graph theory to neuroscience, it is essential to first convert unstructured neuroimaging data into a structured graph (G=(V,E)). In this procedure, the network's nodes and edges are defined according to the kind of connectivity under examination(structural, functional, or effective).

Node Definition and Brain Parcellation

An important first step is to define the nodes of a brain network, which is usually accomplished by a procedure known as brain parcellation. To create a single node in the network, the constant brain volume or surface should be divided into a collection of unique and non-overlapping sections.

1. Anatomical Parcellation

This common technique sections the brain using standardized anatomical atlases. These atlases, like the Harvard-Oxford or Automated Anatomical Labeling (AAL) atlases, provide a systematic and repeatable method of defining areas of interest (ROIs) by segmenting the brain according to macro-anatomical markers (gyri, sulci, and subcortical structures). Then, each anatomically defined ROI turns into a node in the graph, making sure that analyses from various people and research may be compared.

2. Functional Parcellation

Nodes can be defined by homogenous regions derived from resting-state fMRI data. Methods used for functional parcellation include Independent Component Analysis (ICA) to identify functionally coherent networks, clustering algorithms applied to activation patterns and template-based approaches using existing functional atlases.

Edges Definition and Weighting

Functional network construction quantifies statistical dependencies, known as functional connectivity (FC), between time series of nodes. Usually, the correlation coefficient between the time series of node pairs is used to weight the edges in these networks. Pearson's correlation coefficient is frequently used for fMRI data, but functional interactions are measured for EEG/MEG using techniques like spectral coherence, phase synchronization, or lagged phase synchronization. In order to remove weak or false connections, an unweighted graph can be created by applying a threshold to the correlation matrix, where connections that exceed this threshold are viewed to be present (edge = 1) and others are regarded as absent (edge = 0). Functional networks are typically represented as undirected graphs since correlation does not always imply directionality.



Figure 3.1 Brain Network Construction

source : frontiersin.org

Figure 3.1 above is Schematic representation of a typical pipeline for functional brain network construction and subsequent graph theoretical analysis using fMRI data. Time courses are taken from fMRI data and turned into a binary correlation matrix and a functional brain network. Then, graph analysis is done on the connection network of the brain.

The goal of structural network construction is to create the "wiring diagram" of the brain by mapping the anatomical connections between various brain regions. The main tool used in this procedure is Diffusion Tensor Imaging (DTI), a specialized magnetic resonance imaging (MRI) method that gauges how water molecules diffuse throughout brain tissue. DTI enables researchers to infer the orientation and integrity of these brain networks because water diffuses more readily along the highly structured axons within white matter fiber tracts than across them. The first step in the construction process is tractography, which reconstructs the pathways of the main white matter fiber bundles that connect various brain regions using a computational algorithm applied to the raw DTI data. Once brain regions have been defined as nodes through a parcellation scheme, an edge is established between two nodes if a significant number of these reconstructed streamlines are found to connect them. These structural edges are typically weighted to reflect the strength or density of the anatomical connection. Since the DTI tractography techniques frequently detect the presence of fiber bundles between regions without conclusively calculating the typical direction of information flow along those collected pathways at the macro-scale, the resulting structural brain networks are usually represented as undirected graphs.

In order to better understand how connectivity patterns in the brain change and adapt over time, network neuroscience is shifting its attention from static representations to dynamic brain networks. With neuronal interactions varying on time frames ranging from milliseconds to minutes, this method recognizes that brain function is essentially dynamic and can reflect fleeting cognitive states, attentional shifts, or continuing mental activities. Separating relatively large neuroimaging time series (from fMRI, EEG, or MEG, for example) into several shorter, sometimes overlapping time periods is usually the first step in building dynamic brain networks. Through these defined windows, a functional connectivity matrix is computed using methods such as Pearson's correlation coefficient for fMRI data, or spectral coherence for EEG/MEG. Instead of a single brain graph, the outcome is a series of connectivity matrices that each show the network state of the brain at a certain point in time.

Constructing Conceptual Brain Networks

To better show how graph theory can be used to build brain networks, the Python networkx tool can be used to make a simplified conceptual network. This code describes a collection of nodes that are divided into functional groupings like motor, cognitive, and sensory regions. In order to simulate stronger connections within functional groups (which stand for modularity and functional segregation) and sparser but essential connections across groups for global integration, it then creates edges between these nodes with set weights. To illustrate these theoretical linkages, the network is built as an undirected graph.

import networkx as nx

import matplotlib.pyplot as plt

regions = {

'sensory': ["Visual Cortex", "Auditory Cortex", "Somatosensory Cortex"],

'cognitive': ["Prefrontal Cortex", "Parietal Lobe", "Temporal Lobe", "Hippocampus", "Amygdala"],

'motor': ["Motor Cortex", "Basal Ganglia", "Cerebellum", "Thalamus"] } colors = {'sensory': '#9dd69b', 'cognitive': '#87b6de', 'motor': '#e85e60'}

G = nx.Graph()

for group, nodes in regions.items():

G.add_nodes_from(nodes,group=group,color=colors[group])

for group in regions.values():

[G.add_edge(u, v, weight=0.8) for i, u in enumerate(group) for v in group[i+1:]]

cross_conn = [

("Visual Cortex", "Parietal Lobe", 0.9), ("Auditory Cortex", "Temporal Lobe", 0.85),

("Somatosensory Cortex", "Parietal Lobe", 0.9), ("Prefrontal Cortex", "Motor Cortex", 0.7),

("Parietal Lobe", "Motor Cortex", 0.75), ("Amygdala", "Thalamus", 0.8),

("Thalamus", "Visual Cortex", 0.6), ("Thalamus", "Auditory Cortex", 0.6),

("Thalamus", "Somatosensory Cortex", 0.6), ("Hippocampus", "Prefrontal Cortex", 0.7),

("Cerebellum", "Parietal Lobe", 0.65)

]

[G.add_edge(u, v, weight=w) for u, v, w in cross_conn]

if not nx.is_connected(G):

[G.add_edge(list(c)[0], list(c)[0], weight=0.5) for c in nx.connected_components(G)]

plt.figure(figsize=(12, 10))

pos = nx.spring_layout(G, k=0.3, seed=42)

nx.draw_networkx_nodes(G, pos, node_color=[d['color'] for n,d in G.nodes(data=True)],

node_size=2500, alpha=0.9)

nx.draw_networkx_labels(G, pos, font_size=10, font_weight="bold")

nx.draw_networkx_edge_labels(G, pos,

 $edge_labels=\{(u,v): f''\{d['weight']:.2f\}'' for u,v,d in G.edges(data=True)\},$

font_size=8)

plt.axis('off')

plt.tight_layout()

plt.show()

Based on the idea of functional parcellation, each node in this network directly reflects a conceptual brain region, color-coded to represent its larger functional grouping. The connections or communication channels between these brain regions are illustrated by the edges connecting these nodes. The numerical weights given to these edges-the numbers shown on the lines-are also very important since they conceptually show the different levels of interaction effectiveness and strength, modeling how actual brain connections can be stronger or weaker. For example, a more strong information transmission line may be indicated by a thicker or higher-weighted edge. Each node's degree, or the number of direct connections it has (counting the lines that connect each colored circle), shows how connected it is to the rest of the network. High degree nodes are generally seen as more central and can act as local "hubs" for information processing inside their functional group or for making connections across various groups.

C. Graph theory analysis implementation

1. Global Network Measures

The complete structure and effectiveness of the entire brain network are described by global network metrics. They measure the efficiency of information integration and communication between various brain areas throughout the entire system.





source : <u>frontiersin.org</u>

Segregation measures such as modularity and clustering coefficient evaluate the degree of local interconnection among nodes, facilitating effective communication inside modules characteristic path length assesses integration and reflects the effectiveness of information flow globally. Small-world networks provide for both local clustering and short global pathways by balancing regular and random structures. Furthermore, assortativity assesses the durability of networks; assortative networks are more resistant to hub failures than disassortative ones.

2. Local Network Measures

Local network measures concentrate on the traits and significance of specific nodes or their nearby communities within the larger network. They provide information on the distinct functions and contributions of various brain areas. These measures allow researchers to identify crucial individual brain regions and understand their specific roles in information processing. For example, Degree Centrality quantifies the number of direct edges (connections) a specific node (brain region) possesses. The Clustering Coefficient of a node, on the other hand, measures the density of connections among its immediate neighbors, indicating the extent to which a region is part of a closely integrated local processing cluster or "clique". Betweenness Centrality highlights nodes that are frequently on the shortest paths between other pairs of nodes, signifying their importance as "bridges" or "bottlenecks" for information flow across different parts of the brain network. Lastly, Eigenvector Centrality assigns a score to a node based on the influence of its connections, giving higher values to nodes connected to other high-scoring nodes, by then identifying highly influential regions that may not have the most direct connections but are well-integrated into important network pathways. By identifying specific processing capacities and crucial integration or vulnerable spots within neurological networks, these local measurements collectively provide a precise insight of how individual brain regions contribute to overall network performance.



Figure 3.3 Local Network Measures

source : frontiersin.org

D. Disorder and Mental/emotional States Identification

Characterizing different mental and emotional states and quantitatively identifying neurological and psychiatric illnesses are two significant and effective uses of graph theory in neuroscience. This is done by examining the ways in which the topological characteristics of brain networks, which are accurately described by graph theory metrics, differ from a normal basis or display distinctive patterns. The fundamental idea is that abnormal brain activity or distinct cognitive experiences are frequently represented by measurable changes in the complex network structure of the brain.



Figure 3.4 Brain Network of Neurological and Psychiatric Diseases

Source : medium.com

Neurological and Psychiatric Disorders

By measuring the effects of Alzheimer's disease (AD) on the network organization of the brain, particularly the disruption of nodes (which represent different parts of the brain) and their edges (which represent connections between them), graph theory becomes a useful framework for identifying AD. Long-distance information transfer becomes difficult in AD, as shown by the brain's network generally exhibiting a decreased global efficiency and an increased characteristic path length (measures of ideal paths for information flow between nodes via edges). This frequently shows up as a disturbance of the ideal small-world property, where the important equilibrium between global integration and local specialized processing-represented by the coefficient between each node's clustering neighbors-becomes compromised. Moreover, highly linked "hub" nodes-which are measured by their degree or betweenness centrality-are more vulnerable as their edges degrade or vanish, significantly impairing network communication as a whole. A breakdown or change in the structure of functional brain communities may also be indicated by changes in modularity, or the way nodes create communities through their edges. Researchers can find unique network fingerprints linked to AD by carefully measuring these particular graph theoretical metrics that describe the arrangement of nodes and edges.

Graph theory provides an essential tool for comprehending schizophrenia by measuring the disorder's influence on functional brain networks. When graph analysis is applied to fMRI data, it consistently shows that schizophrenia disrupts the healthy brain's ideal small-world features. In particular, it shows decreased global efficiency, which suggests that long-distance communication between brain nodes is restricted. There are noticeable changes in the distribution of edges, such as less short-range and more long-range functional connections, while the local clustering coefficient, which describes regional network patterns, is mostly maintained. Additionally, shifts in nodal metrics such as betweenness centrality indicate a change in the function of certain nodes in regulating the flow of information. The complex network dysfunction that underlies schizophrenia can be objectively detected by graph theory by identifying these particular graph theoretical abnormalities in nodes and edges.

Identifying Cognitive Processes and Mental States

Graph theory reveals how brain networks dynamically reconfigure to serve higher-order tasks, making it a helpful tool for studying a variety of cognitive processes and mental states outside of pathological settings. In contrast to diminished integration and higher modularity in unconscious states, graph analysis of consciousness reveals excellent global integration (efficiency) during waking. Through quantifiable shifts in edge weights (connectivity strength) and the rearranging of modules or hub node functions, which reflect ideal information flow, graph theory captures brain plasticity for learning and skill acquisition. Similar to this, dynamic network reconfigurations involving temporary shifts in modularity and adaptive changes in the centrality of critical nodes are a sign of a variety of mental states, such as high cognitive load or attention, demonstrating the brain's adaptability in resource allocation.

IV. CONCLUSION

The immense complexity of the human brain continues to be a primary area of scientific research. This study demonstrates how graph theory, an essential element of discrete mathematics, may be used to effectively describe and comprehend the complex network of connections found in the brain. Graph theory offers an organized method of analyzing brain organization by illustrating brain regions as nodes and their anatomic alongside all functional links as edges, derived from methods such as MRI, DTI, and fMRI data. This powerful framework allows for the quantification of key network features through global measures like efficiency and local measures such as centrality, analyses of properties like small-worldness and modularity.

Importantly, these graph-theoretic approaches have significantly improved our comprehension of neurological disease as well as normal brain function. The distinctive small-world architecture of the brain supports a variety of cognitive functions, including learning, memory formation, and consciousness, by striking a balance between specialized local processing and effective global integration. Furthermore, neurological disorders have been directly connected to measurable changes in certain network qualities, such as altered efficiency, disturbed hub structure, or modified modularity. Graph theoretical analysis will continue to provide a deeper understanding of the structure, function, and dysfunction of the brain through the potent lens of discrete mathematics.

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PERNYATAAN

Dengan ini saya menyatakan bahwa makalah yang saya tulis ini adalah tulisan saya sendiri, bukan saduran, atau terjemahan dari makalah orang lain, dan bukan plagiasi.

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