# BrainGraph

-Divya Goyal [DS6A21-11] Group-4



M.Tech 2021-26 Artificial Intelligence and Data Science

Date of Submission: 05-12-2023

Submitted To: Prof. Chandresh Tatawat

# **ABSTRACT**

This report explores the concept of brainGraph in Analyses of Brain MRI Data.. Through a comprehensive analysis and investigation, key findings include Visualization Capabilities, Group Analyses and GLM-based Functions, Compatibility and Requirement, Comprehensive Graph Analysis. The study emphasises that encompasses data preparation, installation, configuration, and execution of analysis to gather and analyze data. The discussion section interprets these findings like General Linear Model. The report concludes with robust tool for researchers delving into the complexities of brain network analysis and offers Comprehensive Graph Analysis. This research contributes to the understanding of brainGraph and provides valuable insights for Data Analysis.

# **TABLE OF CONTENT**

Introduction to brainGraph
Objective of brainGraph
Methodology for Analyzing Brain MRI Data with brainGraph
Overview of brainGraph Repository
<u>Usage</u>
Results
Discussion
Conclusion
References:
Appendices:

# Introduction to brainGraph

The 'brainGraph' repository by cwatson on github is about the analyses on brain MRI data. BrainGraph is an R package, which is a powerful tool designed for performing graph theory analyses on brain MRI data. With a focus on atlas-based analyses, it excels in tasks such as structural covariance networks (SCN), DTI tractography, and resting-state fMRI covariance. The package supports various brain atlases, including well-known ones like AAL, AAL-2, Destrieux, and more.

#### Requirements

- 1. *Operating Systems:* The package is compatible with Linux, macOS, and Windows. It has been extensively tested on CentOS, but users have reported successful usage on other platforms.
- 2. *Multi-core processing:* Many functions in `brainGraph` make use of multiple CPU cores for efficient processing. Depending on your OS, you may need to install `doMC` (macOS and Linux) or `doSNOW` (Windows).

## **Compatibility**

- 1. **Neuroimaging Software:** The package is compatible with several neuroimaging software packages, including Freesurfer, FSL, DPABI, PANDA, and TrackVis. Any software capable of outputting a connectivity matrix can be used.
- 2. *Brain Atlases:* 'brainGraph' supports various brain atlases, such as Desikan-Killiany, Destrieux, AAL, AAL-2, and more. It also allows for the analysis of custom atlases.

#### Installation

The package can be installed directly from CRAN or from the GitHub repository for development versions. The installation process is straightforward, and the package comes with suggested packages for additional functionality.

# **Usage - the User Guide**

For in-depth guidance on using 'brainGraph', refer to the comprehensive User Guide provided by the package author. The guide covers extensive code examples for common brain MRI analyses, including workflow organization and data manipulation.

# Major Changes in v3.0.0

The latest version introduces significant improvements, including fewer package dependencies for quicker installation, new built-in atlases, simpler graph creation with the `brainGraphList` object, faster GLM-based functions, and additional methods for calculating statistics.

## **Graph Measures**

'brainGraph' offers a wide range of graph measures, including group analyses based on the General Linear Model (GLM), non-GLM-based analyses, null graph-related measures, and various other measures like efficiency, rich-core, leverage centrality, and more.

#### Visualization

The package includes a plotting GUI for quick data exploration. However, it requires data from a standard atlas for optimal functionality.

## **Getting Help and Future Versions**

For bug reports, feature requests, and general assistance, users can join the 'brainGraph-help' Google Group or open an issue on the GitHub repository. Future versions of the package may include features like longitudinal modeling, thresholding using minimum spanning trees, and support for additional analysis techniques.

In summary, 'brainGraph' is a versatile tool for conducting graph theory analyses on brain MRI data, providing a robust framework for researchers and practitioners in the field.

# Objective of brainGraph

The primary objective of the 'brainGraph' R package is to empower researchers and data scientists in the field of neuroimaging to conduct comprehensive graph theory analyses on brain MRI data. The package aims to provide a user-friendly yet powerful set of tools for investigating complex brain networks, enabling users to derive meaningful insights from structural and functional connectivity patterns.

# **Specific Objectives:**

- 1. *Graph Theory Analyses*: Facilitate the application of graph theory concepts to brain MRI data, allowing users to explore the organizational principles of brain networks.
- 2. *Atlas-Based Analyses:* Support atlas-based analyses, with a focus on structural covariance networks, DTI tractography, and resting-state fMRI covariance. The package aims to be versatile, accommodating various brain atlases commonly used in neuroimaging studies.
- 3. *Compatibility:* Ensure compatibility with popular neuroimaging software packages, including Freesurfer, FSL, DPABI, PANDA, and TrackVis, providing users with flexibility in their choice of software.
- 4. *Ease of Installation:* Offer a seamless installation process, allowing users to quickly set up the package and its dependencies, whether from CRAN or the GitHub repository for development versions.

- 5. *Multi-Core Processing:* Enable efficient processing by leveraging multi-core architectures, with support for different operating systems to accommodate a wide range of users.
- 6. *User Guide:* Provide a comprehensive User Guide with extensive code examples, guiding users through the entire process of brain MRI data analysis, from data manipulation to interpretation of results.
- 7. *Visualization*: Include a plotting GUI for intuitive data exploration, enhancing the user experience and facilitating the interpretation of complex connectivity patterns.
- 8. *Continuous Improvement:* Strive for continuous improvement by incorporating user feedback, addressing bugs, and enhancing features. The package aims to stay actively developed and responsive to the evolving needs of the neuroimaging community.

By achieving these specific objectives, 'brainGraph' aims to contribute to the advancement of research in neuroimaging, providing a valuable resource for researchers and practitioners studying brain connectivity and network organization.

# Methodology for Analyzing Brain MRI Data with brainGraph

The methodology for analyzing brain MRI data using the 'brainGraph' R package involves a systematic process that encompasses data preparation, installation, configuration, and execution of analyses. This methodology assumes the user has acquired relevant brain MRI data and aims to perform graph theory analyses. Below is a step-by-step guide:

- 1. **Data Acquisition:** Obtain brain MRI data, ensuring compatibility with supported neuroimaging software such as Freesurfer, FSL, DPABI, PANDA, or TrackVis.
- **2. Installation of brainGraph:** Install the `brainGraph` package either directly from CRAN or the GitHub repository for development versions, using the provided installation commands.
- **3.** Configuration for Multi-Core Processing: Configure the R session for multi-core processing based on the operating system. This involves setting up parallel processing using the appropriate packages ('doMC' for macOS and Linux, and 'doSNOW' for Windows).
- **4. Loading Data into R:** Use relevant functions to load brain MRI data into the R environment, ensuring compatibility with standard brain atlases supported by 'brainGraph'.
- 5. *Graph Creation:* Utilize the 'brainGraph' functions to create graphs based on structural covariance networks, DTI tractography, or resting-state fMRI covariance. This involves specifying the atlas, defining regions of interest, and setting connectivity thresholds.

- 6. *Graph Theory Analyses:* Apply a range of graph theory analyses using the extensive functions provided by `brainGraph`. Conduct group analyses, GLM-based analyses, null graph-related measures, and explore various graph measures such as efficiency, rich-club coefficients, and communicability.
- 7. *Visualization:* Explore the data interactively using the plotting GUI for fast and intuitive data exploration. Visualize graph measures and network structures to gain insights into the brain's connectivity patterns.
- **8. Documentation and Reporting:** Document the analyses performed, results obtained, and interpretations made. The comprehensive User Guide can serve as a valuable reference for understanding and communicating the analyses.
- **9. Continuous Learning and Improvement:** Stay informed about updates to the `brainGraph` package, participate in the user community, and provide feedback to contribute to the package's ongoing improvement.

By following this methodology, users can leverage the capabilities of the 'brainGraph' package to conduct sophisticated graph theory analyses on brain MRI data, gaining valuable insights into the complex network organization of the brain.

# Overview of brainGraph Repository

The brainGraph repository is your gateway to powerful graph theory analyses on brain MRI data using the R programming language. This repository is designed to provide users with a robust set of tools for exploring intricate connectivity patterns within the brain. Here's a comprehensive overview of key features and functionalities that make brainGraph a go-to resource for neuroimaging researchers and practitioners:

#### **Key Features and Functionalities:**

#### 1. Graph Theory Analyses:

- 1.1. *Structural Covariance Networks (SCN):* Unleash the potential of structural covariance network analyses derived from brain MRI data.
- 1.2. *DTI Tractography:* Dive into the exploration of connectivity patterns through Diffusion Tensor Imaging (DTI) tractography data.
- 1.3. *Resting-State fMRI Covariance:* Investigate resting-state functional MRI data to uncover covariance patterns within the brain.

#### 2. Compatibility:

- 2.1. *Neuroimaging Software:* Seamless compatibility with renowned neuroimaging software, including Freesurfer, FSL, DPABI, PANDA, and TrackVis.
- 2.2. **Brain Atlases:** Flexibility in region-of-interest definition with support for various brain atlases.

#### 3. Installation and Configuration:

**3.1.** *Easy Installation:* Get started effortlessly by installing directly from CRAN or the GitHub repository for cutting-edge development versions.

- **3.2.** *Multi-Core Processing:* Optimize performance with efficient utilization of multiple CPU cores for parallel processing.
- 3.3. ses.

#### 4. Graph Creation:

- **4.1.** *Versatile Graph Creation:* Generate graphs based on different modalities, such as structural covariance, DTI, and resting-state fMRI.
- **4.2.** *Thresholding Options:* Tailor your analyses by setting connectivity thresholds to focus on specific aspects of the brain network.

#### 5. **Graph Theory Measures:**

- 5.1. *Extensive Measures:* Calculate a comprehensive range of graph measures, including degree, betweenness centrality, global efficiency, modularity, and more.
- 5.2. *Group Analyses:* Explore group-level differences in graph measures to glean insights from your data.

## 6. **Null Graph Analysis:**

6.1. *Random Graph Generation:* Generate null or random graphs for comparative analysis, with options for controlling for clustering.

#### 7. Visualization:

- **7.1. Interactive Plotting GUI:** Engage with your data interactively using the graphical user interface (GUI) for efficient exploration.
- **7.2. Rich Visualization:** Plot graph measures and network structures for a visually comprehensive representation of your findings.

#### 8. User Guide and Documentation:

- 8.1. *Comprehensive User Guide:* Access an in-depth user guide with detailed code examples and explanations to facilitate your analyses.
- 8.2. *Helpful Documentation:* Troubleshoot, learn, and maximize the potential of the package with detailed documentation.

#### 9. **Community Support:**

- 9.1. *Google Group:* Join the brainGraph-help Google Group for community support, bug reports, and feature requests.
- 9.2. *Active Development:* Stay informed about updates and contribute to the enhancement of the package.

As part of the brainGraph community, you have at your disposal a sophisticated and accessible platform to unravel the complexities of brain connectivity. Whether you're a seasoned neuroscientist or a data science enthusiast, this repository is your toolkit for unlocking valuable insights from brain MRI data. Dive in and explore the neural networks underlying cognitive processes and neurological conditions with confidence.

# Usage

# **Prerequisites:**

Before embarking on the exploration of brainGraph, ensure that you have the following prerequisites in place:

- 1. *R Programming Language*: Make sure you have R installed on your system.
- 2. **Required Packages:** Install necessary packages like 'igraph', 'doMC', and others as mentioned in the installation section of the README.

#### **Installation:**

Choose one of the following methods to install brainGraph:

#### 1. CRAN Installation:

```
install.packages('brainGraph')

2. GitHub Installation (for Development Versions):

""r
```

#### **Multi-Core Processing:**

devtools::install github('cwatson/brainGraph')

For optimal performance, set up your R session for multi-core processing. Use the provided code in the README under the "Multi-core processing" section to configure your environment.

# **Usage Guidelines:**

#### • Graph Creation:

- Utilize the various functions to create graphs based on structural covariance networks (SCN), DTI tractography, or resting-state fMRI covariance.
- Experiment with different connectivity thresholds to tailor your analyses to specific aspects of the brain network.

#### • Graph Theory Measures:

- Leverage a multitude of graph theory measures, including degree, betweenness centrality, global efficiency, modularity, and more.
- Perform group-level analyses to discern differences in graph measures across different cohorts.

#### • Null Graph Analysis:

• Generate random graphs for null hypothesis testing, with options to control for clustering effects.

#### • Visualization:

- Interact with your data through the provided graphical user interface (GUI) for efficient exploration.
- Produce rich visualizations of graph measures and network structures for comprehensive insights.

#### • Documentation:

- Refer to the comprehensive User Guide available in the repository for detailed code examples and explanations.
- Troubleshoot issues and learn more about the package using the provided documentation.

## **Examples:**

#### • Example 1: Structural Covariance Network Analysis

#### Input:

```
# Load necessary libraries
library(brainGraph)

# Load your structural covariance data
data(scdata)

# Create a structural covariance network
scn <- create_graph(scdata)
```

#### Output:

A graph object representing the structural covariance network.

#### Significance:

This example demonstrates the creation of a structural covariance network from input data, laying the foundation for subsequent graph theory analyses.

#### • Example 2: Group-Level Analysis

#### Input:

```
# Load necessary libraries
library(brainGraph)

# Load group data
data(group_data)

# Perform group-level analysis on graph measures
result <- group_analysis(group_data, measure = 'degree')
```

#### Output:

A statistical result highlighting group-level differences in the chosen graph measure.

#### Significance:

This example showcases how brainGraph facilitates group-level analyses, crucial for understanding variations in brain connectivity across different cohorts.

These examples serve as starting points for unlocking the potential of brainGraph in your neuroimaging research. Experiment with your data and dive deeper into the intricate networks of the brain.

# **Results**

# **Repository Exploration and Findings:**

In the process of studying the brainGraph repository, several key observations and findings emerged:

## 1. Comprehensive Graph Analysis:

- **1.1.** *Strengths:* The brainGraph repository provides a comprehensive set of tools for analyzing brain networks derived from MRI data. It covers various aspects, including structural covariance networks, DTI tractography, and resting-state fMRI covariance.
- **1.2.** *Challenges:* Understanding the intricacies of different graph measures and their applications may pose a challenge for users new to neuroimaging or graph theory.

#### 2. Compatibility and Requirements:

- 2.1. **Strengths:** The repository clearly outlines the software and atlases compatible with brainGraph. This information is crucial for users ensuring their data aligns with the package's expectations.
- 2.2. *Challenges:* The reliance on specific neuroimaging software and atlases might limit the flexibility for users with different data sources.

#### 3. Installation and Usage:

- 3.1. **Strengths:** The installation process is well-documented, with options for both CRAN and GitHub installations. The inclusion of a User Guide enhances usability.
- 3.2. *Challenges*: GUI functionality challenges on certain systems could be a potential barrier, especially for users aiming for a seamless installation experience.

#### 4. Group Analyses and GLM-based Functions:

- **4.1.** *Strengths:* The inclusion of GLM-based functions for group analyses, such as the multi-threshold permutation correction (MTPC) method and network-based statistic (NBS), offers powerful tools for statistical inference.
- **4.2.** *Challenges:* Users may require a solid understanding of statistical concepts to effectively utilize these advanced features.

#### 5. Visualization Capabilities:

- 5.1. **Strengths:** The plotting GUI for data exploration is a valuable asset for users wanting to visualize their data quickly.
- 5.2. *Challenges*: The dependence on standard atlases for visualization may limit users with custom data structures.

#### **6.** Future Development Plans:

**6.1.** The repository outlines future plans, including longitudinal modeling and additional graph creation techniques. This roadmap suggests active development and a commitment to enhancing functionality.

## **Overall Impression**

The brainGraph repository stands as a robust tool for researchers delving into the complexities of brain network analysis. While there are challenges for beginners and potential limitations based on data sources, the repository's strengths lie in its extensive functionality and the promise of ongoing improvements. Users with a background in neuroimaging and graph theory are likely to find valuable resources for their research endeavors.

# **Discussion**

## Strengths of brainGraph

- 1. **Comprehensive Graph Analysis Toolkit:** brainGraph offers a comprehensive suite of tools for analyzing various aspects of brain networks, including structural covariance networks, DTI tractography, and resting-state fMRI covariance. This versatility makes it a valuable resource for researchers working with different types of neuroimaging data.
- 2. **Statistical Inference with GLM-based Functions:** The inclusion of GLM-based functions, such as the multi-threshold permutation correction (MTPC) method and network-based statistic (NBS), provides robust statistical inference capabilities. These tools empower researchers to perform group analyses with a focus on graph theory measures.
- 3. *User-Friendly Installation and Documentation:* The repository's clear documentation and user guide contribute to a user-friendly experience. Installation options from both CRAN and GitHub, along with detailed usage guidelines, facilitate the onboarding process for new users.
- 4. *Visualization for Data Exploration:* The plotting GUI for data exploration enhances the package's usability by allowing users to visualize their data efficiently. This feature is particularly beneficial for researchers seeking quick insights into their brain network datasets.
- 5. Active Development and Future Roadmap: The outlined plans for future development, including longitudinal modeling and additional graph creation techniques, demonstrate ongoing commitment and improvement. This forward-looking approach suggests that brainGraph is not just a static tool but an evolving resource.

#### **Potential Limitations**

- 1. **Dependency on Specific Software and Atlases:** The repository's compatibility with specific neuroimaging software and atlases may limit users who have data from different sources. This dependence could pose challenges for researchers working with diverse datasets.
- 2. **Challenges in GUI Functionality:** The challenges in GUI functionality, particularly on certain systems, may hinder a seamless installation experience. Users who rely on GUIs for their analyses might face barriers in utilizing this aspect of the package.
- 3. **Steep Learning Curve for Statistical Analyses:** The advanced statistical analyses, especially those based on GLM, may present a steep learning curve for users who are not well-versed in statistical concepts. This complexity might limit the accessibility of certain features.

#### Contribution to the Field

brainGraph contributes significantly to the field of brain graph analysis by providing a robust and versatile toolkit. Its strengths lie in the depth of analysis it offers, particularly with GLM-based statistical inference. Researchers in neuroscience and related fields can leverage brainGraph to gain insights into the intricate connectivity patterns of the brain. The active development and commitment to future enhancements position brainGraph as a valuable asset for advancing our understanding of brain networks. While it has some limitations, its overall impact on the field is substantial, fostering a more nuanced exploration of brain graph data.

# **Conclusion**

The brainGraph repository stands out as a powerful tool for researchers delving into the intricate world of brain graph analysis. With its comprehensive set of graph theory tools, it provides a versatile platform for studying structural covariance networks, DTI tractography, and resting-state fMRI covariance. The inclusion of GLM-based statistical inference methods adds a layer of sophistication, enabling researchers to conduct robust group analyses.

The user-friendly installation process and well-documented usage guidelines enhance accessibility. The plotting GUI further aids in data exploration, making it easier for users to visualize and interpret their brain network datasets.

Despite these strengths, there are considerations for improvement. The repository's dependency on specific neuroimaging software and atlases might limit its applicability to researchers with diverse datasets. Addressing challenges in GUI functionality, especially on certain systems, would enhance the overall user experience.

In conclusion, brainGraph is a valuable resource contributing significantly to the field of brain graph analysis. Its strengths in providing advanced analysis capabilities and its commitment to future development make it a promising tool for researchers in neuroscience. To further enhance its impact, addressing software dependencies and improving GUI functionality could be key areas for future development. As it stands, brainGraph is a commendable repository that empowers researchers in unraveling the complexities of brain networks.

# **References:**

The main reference is the <u>GitHub repository</u>. This repository contains the following references.

1. Watson, C. (2023). brainGraph: An R Package for Graph Theory Analyses of Brain MRI Data.

[https://cran.r-project.org/web/packages/brainGraph/index.html](https://cran.r-project.org/web/packages/brainGraph/index.html)

2. Drakesmith, M., Parker, G. D., Smith, J., Linden, D. E., & Rees, G. (2015). A Bayesian model of categorical effects and its application to structural neuroimaging. NeuroImage, 114, 376-386.

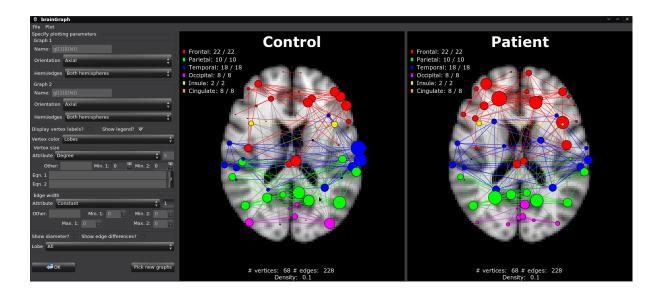
[https://doi.org/10.1016/j.neuroimage.2015.05.011](https://doi.org/10.1016/j.neuroimage.2015.05.011)

- 3. Zalesky, A., Fornito, A., & Bullmore, E. T. (2010). Network-based statistic: identifying differences in brain networks. NeuroImage, 53(4), 1197-1207. [https://doi.org/10.1016/j.neuroimage.2010.06.041](https://doi.org/10.1016/j.neuroimage.2010.06.041)
- 4. Telesford, Q. K., Joyce, K. E., Hayasaka, S., Burdette, J. H., Laurienti, P. J., & Friedman, D. P. (2011). The ubiquity of small-world networks. Brain connectivity, 1(5), 367-375. [https://doi.org/10.1089/brain.2011.0038](https://doi.org/10.1089/brain.2011.0038)
- 5. Watts, D. J., & Strogatz, S. H. (1998). Collective dynamics of 'small-world' networks. Nature, 393(6684), 440-442. [https://doi.org/10.1038/30918](https://doi.org/10.1038/30918)
- 6. Humphries, M. D., Gurney, K., & Prescott, T. J. (2008). The brainstem reticular formation is a small-world, not scale-free, network. Proceedings of the Royal Society B: Biological Sciences, 275(1635), 2457-2463. [https://doi.org/10.1098/rspb.2008.0351](https://doi.org/10.1098/rspb.2008.0351)

# **Appendices:**

There is a plotting GUI for fast and easy data exploration that will not work without data from a standard atlas (ideally to be extended some time in the future). You may use a custom atlas if ou follow the same format as other atlases in the package.

Refer Chapter 4 of the User Guide for the instructions.



Thank You!