Report Biomarkers

Benedetta Corso, Letizia Rossato, Dalila Dattoli June 11, 2024

Abstract

Abstract: introduce rationale of investing brain dopamine function and potential brain asymmetry in PD.

This study, which focuses on Parkinson's disease, aims to investigate the significance of left-right lateralization based on dopamine values using DAT-SPECT imaging data and their correlation with clinical symptoms in Parkinson's disease patient (PD) and healthy controls (HC). DAT-SPECT (Dopamine Transporter Single Photon Emission Computed Tomography) is an imaging technique primarily used to visualize and measure the density of dopamine transporters in the brain. Indeed, this technique is particularly useful in the diagnosis and evaluation of neurodegenerative diseases, such as Parkinson's disease and other forms of parkinsonism. The dataset, used to perform the analysis, consists of variables relating to demographic data, DAT-SPECT scores and finally the Neuropsychological (NP) tests which are a combination of clinical, neurological, and imaging test since there is no specific test for the diagnosis of Parkinson's disease.

ADD RESULTS

1 Introduction

Parkinson's disease is a chronic neurodegenerative disorder of the central nervous system primarily affecting motor control. It is characterized by the progressive loss of nerve cells in the brain region called the substantia nigra, which is responsible for producing a neurotransmitter called dopamine. Dopamine deficiency leads to motor symptoms such as tremors, muscle rigidity, bradykinesia (slowness of movement), and postural instability. In addition to motor symptoms, Parkinson's disease can cause a wide range of non-motor symptoms including sleep problems, depression, anxiety, fatigue, and cognitive difficulties. While the exact cause of Parkinson's disease is not fully understood, it is believed to result from a combination of genetic and environmental factors [1]. Currently, there is no cure for Parkinson's disease, but there are treatments available to manage symptoms and improve the patients' quality of life. However, studies conducted in recent years have shown that the lateralization of brain dopamine in Parkinson's disease (PD) is a significant and distinctive aspect of the pathology. In particular, during its development, the degeneration of dopaminergic neurons in the substantia nigra is not uniform. Generally, one side of the brain is more affected than the other, leading to marked asymmetry. This marked asymmetry is visible not only in motor symptoms, such as more pronounced tremors on one side of the body and greater use of one hand over the other, but it can also be evident in non-motor disturbances, including cognitive issues for example in the sleep behavior and or sense of smell [3]. In previous studies, the connection between dopamine lateralization and symptoms has already been examined. For example, in [2], it resulted that the motor symptoms on the less affected side were more correlated to striatal DAT

binding.

Therefore, the aim of this study is to further investigate this idea, analyzing a dataset containing information about patients' motor and cognitive symptoms and also their lateralization data obtained from DAT-SPECT scans. The dopamine degeneration is explored in three different Regions of Interests (ROIs) in the brain, divided in right and left: Caudate, Putamen and Putamen Anterior. This analysis aims to investigate the relationship (if present) between the lateralization of dopamine function in these brain areas and the symptoms showed by Parkinson's patients, and relationship (if present) between the lateralization of dopamine function in these brain areas and possible covariates showed by healthy controls.

Agreeing with what found in [2], the group expected to find a strong relation between the dopamine lateralization and the motor symptoms. Also the lateralization in different ROIs was expected to be related: a high lateralization in the Caudate was expected if both Putamen and Putamen Anterior were highly lateralized.

2 Material and Methods

2.1 Dataset description

The dataset used in this study was composed of 1556 subjects, of which 256 healthy controls and 1300 PD patients with different levels of symptoms' severity. Not all the variables given were interesting for the study. For example, all the data regarding the MRI acquisitions were discarded, because the analysis focused on the data from the DAT-SPECT scans, which were Striatal Binding Ratios (SBR) of the three ROIs (right and left sides), whether the scan was completed and its quality based on visual interpretation, as well as information regardind the date of the scan and the injection.

The dataset included a part of demographics data, like age, ethnicity, gender, family members affected by PD, height, weight and dominant hand. A total of 967 and 589 patients were males and females, respectively. The average age in the baseline was 62.69 ± 10.11 years (range [29.3–86.5]).

There was then the data regarding the neuropsychological assessments of the patients using the Movement Disorder Society (MDS)sponsored revision of the Unified Parkinson's Disease Rating Scale (MDS-UPDRS). The test is divided in 4 parts, concerning different classes of motor and cognitive symptoms, as specified in the PPMI Program Protocols (THE PARKIN-SON'S PROGRESSION...). Each answer could range from 0 (normal) to 4 (severe), and for each part there was a summing-up variable, that contained the sum of all the scores of that section. Some motor symptoms, especially in the third part, were divided in right and left, for example, the severity of tremor of the right or left upper limb. This was useful to investigate the different relation with the dopamine function lateralization.

Più aggiungere forse grafici con distribuzioni variabili iniziali, tipo grafici lateralizzazione

2.2 Cleaning e preprocessing

The first step was dividing the dataset in patients (PD, SWEDD, Prodromal) and HC.

In the preprocessing of the data, the patterns of missing values were analyzed, to determine the best course of action, through the VIM package in R.

In the HC, as shown in figure 1, GENETICS and FAMILIARITY are mostly empty fields, due to the nature of these variables: for healthy subjects, the chance of having relatives with a PD diagnosis and of having genetics history related to PD are low. On the opposite, for PD patients (figure 2), NP4TOT is missing at the 73.9%, following the same pattern of the PD patients (figure 2). This suggests that the fourth part of the test was taken by a few subjects. For this reason, the NP4 variables were not taken into account in the statistical analysis.

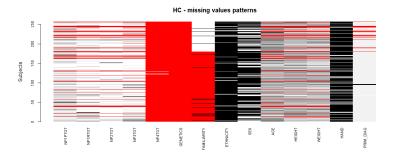


Figure 1: Pattern of missing values in HC

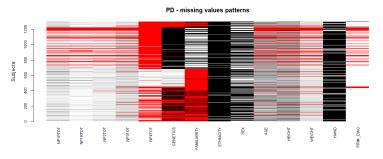


Figure 2: Pattern of missing values in PD

The missing data in the ENROLL_AGE variable was computed using the DAT-SCAN date and the birth date of the subject.

Che dati abbiamo escluso, perchè + MISS-ING DATA e come li abbiamo trattati ++ GRAFICO missing data (spiegazione parte di R) Abbiamo controllato che i hc e pd fossero simili (stessa pop di m/f età etc)

2.3 Statistical methods

2- Che statistiche abbiamo usato, perchè Anova per vedere se c'è diff tra hc e pd [forse è meglio sostituire con wilxcon?]

2.4 Feature extraction

- Come abbiamo trovato il lateralization index [REFERENCE] - Matrice di correlazione in base alla matrice di correlazione - $\dot{\iota}$ scelto le variabili più significative

2.5 Linear regression

 $\begin{array}{lll} {\rm IMMAGINE} & {\rm linear} & {\rm regression} & + & {\rm descrizione} \\ {\rm statistiche} & & & & \\ \end{array}$

3 Results

- A clear and concise description of the statistical results providing answers to the research questions
- A sensitivity analysis of the results to covariates, group matching and data quality (e.g. missing data, data miss balance)

4 Discussion

- Direct answers to the research questions
- An overview of the limitations of the study
- A list of possible suggestions to improve the study in case someone will repeat it in future

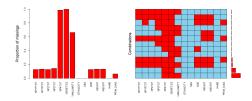


Figure 3: Missing values in HC

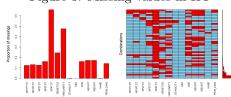


Figure 4: Missing values in PD

References

- [1] Janice M. Beitz. "Parkinson's disease: a review." eng. In: Frontiers in bioscience (Scholar edition) 6.1 (Jan. 2014). Place: Singapore, pp. 65–74. ISSN: 1945-0524 1945-0516. DOI: 10.2741/s415.
- [2] Walter Pirker. "Correlation of dopamine transporter imaging with parkinsonian motor handicap: How close is it?" en. In: *Movement Disorders* 18.S7 (Oct. 2003), S43–S51. ISSN: 0885-3185, 1531-8257. DOI: 10.1002/mds.10579. URL: https://onlinelibrary.wiley.com/doi/10.1002/mds.10579 (visited on 05/31/2024).
- [3] P. Riederer et al. "Lateralisation in Parkinson disease". en. In: Cell and Tissue Research 373.1 (July 2018), pp. 297-312. ISSN: 0302-766X, 1432-0878. DOI: 10.1007/s00441-018-2832-z. URL: http://link.springer.com/10.1007/s00441-018-2832-z (visited on 05/31/2024).