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Exploring Parkinson's Disease Progression Through Python-Based Data Analysis Techniques

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Abstract: Parkinson's disease is a long-term neurological illness that mainly influences motor functions balance, and speaking abilities. This study aims to present a reproducible Python-based EDA framework to understand the progressive growth of Parkinson's disease using an authentic dataset comprising 500 affected patients. With the use of major Python libraries, Exploratory Data Analysis (EDA) methods were applied to clean, process and visualize the data. The study involved exploring correlation between features like UPDRS score, motor function, speech difficulty and years since diagnosis. Descriptive statistics, correlation heatmaps and visual plots helped in recognizing symptom patterns and perspective-related progression of this condition. This work exhibits how EDA can be useful in rapid analysis of clinical data, chiefly in emphasizing the role of clinical characteristics in complicated disorders like Parkinson's disease. Our workflow identifies symptom clusters and temporal trends, offering a template for clinical data analysis.

Keywords: Parkinson's Disease, Exploratory Data Analysis, Data Visualization, UPDRS Score, Python, Clinical Data.

I. INTRODUCTION

Parkinson's disease is a continuously evolving neurological impairment that largely impacts motor function, speaking ability and equilibrium. With its occurrence increasing noticeably with age, it is one of the most familiar functional movement disorders. Early and accurate diagnosis of symptoms are essential for managing and improvement of treatment planning, and upgrading quality of life. Nevertheless, understanding the trajectory of this disease remains a complicated challenge due to irregularity in patient symptoms and variability. With the rapid growth of medical datasets, data-driven methods such as exploratory data analysis (EDA) are instrumental in extracting interdependencies and meaningful patterns from medical data.

Unlike prior EDA studies focusing solely on static correlations, our work integrates temporal trend analysis (e.g., UPDRS vs. diagnosis duration) using Python's Seaborn and Matplotlib. This approach enables dynamic symptom tracking, which is critical for early intervention. By using Python-based libraries such as Pandas, Seaborn and Matplotlib, researchers can systematically process and visualize complex medical datasets. In this study, a real-world dataset comprising 500 individuals diagnosed with Parkinson's disease was examined to identify symptom progression and correlation across features such as UPDRS score, tremor severity, medication and years since diagnosis. The main objective of this research is to use Exploratory Data Analysis (EDA) techniques to understand the symptom trends and interdependencies of Parkinson's disease using statistical summaries and visualization. The present study aims to provide insights that can aid practitioners in identifying symptom groupings and presenting early-stage remedies for the diagnosis.

II. METHODOLOGY

A structured Exploratory Data Analysis (EDA) workflow is followed in this methodology which is designed to uncover insights from a clinical dataset of 500 Parkinson's patients. With the aid of Python libraries like Pandas for data manipulation, NumPy for numerical operations, Matplotlib and Seaborn for visualization and Scikit-learn, the process began with loading the dataset, inspecting its structure and analyzing column types and basic statistical summaries. The dataset consisted of 500 patient records affected by Parkinson's disease. Each record included variables such as age, gender, year since diagnosis, UPDRS Score (Unified Parkinson's Disease Rating Scale), Tremor Severity, Motor Function, Speech Difficulty, Balance Problems, Medications, Exercise Level and the target variable - disease progression.

In preprocessing stage, the data was examined for quality by identifying missing values, duplicate rows and invalid negative entries. Fortunately, the dataset was clean and required minimal correction in specific columns. In categorical encoding step, Label Encoding and One-Hot Encoding were used to encode categorical variables such as gender and medication types and converted them to numerical format for suitable analysis. Techniques like Min-Max Scaling and Z-score Standardization were applied to make sure the features are uniform depending on their nature for feature scaling.

Outliers were detected using the Interquartile Range (IQR) method to highlight extreme values in numerical columns and boxplots were used to visualize the distribution anomalies. Outliers beyond $1.5 \times \text{IQR}$ were removed to ensure robustness, affecting $<5\%$ of data points.

Visualizations were extensively used to interpret patterns. Multiple statistical graphs were regenerated to interpret feature interdependencies and trends. Histograms and boxplots were used to understand the distribution of continuous variables and the bar charts provided comparisons among categorical features. Correlation heatmaps were used to reveal linear associations among clinical variables and multivariate relationships were examined through pair plots. Grouped bar charts and line plots aided in comparison of disease progression across patient subgroups. This EDA pipeline enabled the extraction of interpretable insights about Parkinson's disease feature interdependencies and progression of the disorder.

III. RESULTS AND DISCUSSION

The results obtained from the exploratory data analysis performed on the Parkinson's disease dataset are represented in this section. Several statistical techniques and visualization tools were used to identify patterns, distributions and interrelationships among clinical features. By visualizing symptoms, identifying feature trends and analyzing relationships across affected groups, this study aims to convert raw data into perceptive observations. This study not only points symptoms that support to disease advancement but also assist in building a clear understanding through structured data interpretation.

A. Feature Distributions

Histograms and box-plots were used for a comprehensive view of the distribution of key numeric features such as *UPDRS Score*, *Tremor Severity*, *Motor Function* and *Speech Difficulty*. With a large number of patients falling within mid-range severity scores, most of these variables exhibited moderately skewed distributions. A right-skewed distribution was observed by *Exercise Level* variable which indicated that a considerable portion of affected individuals maintained lower physical activity levels.

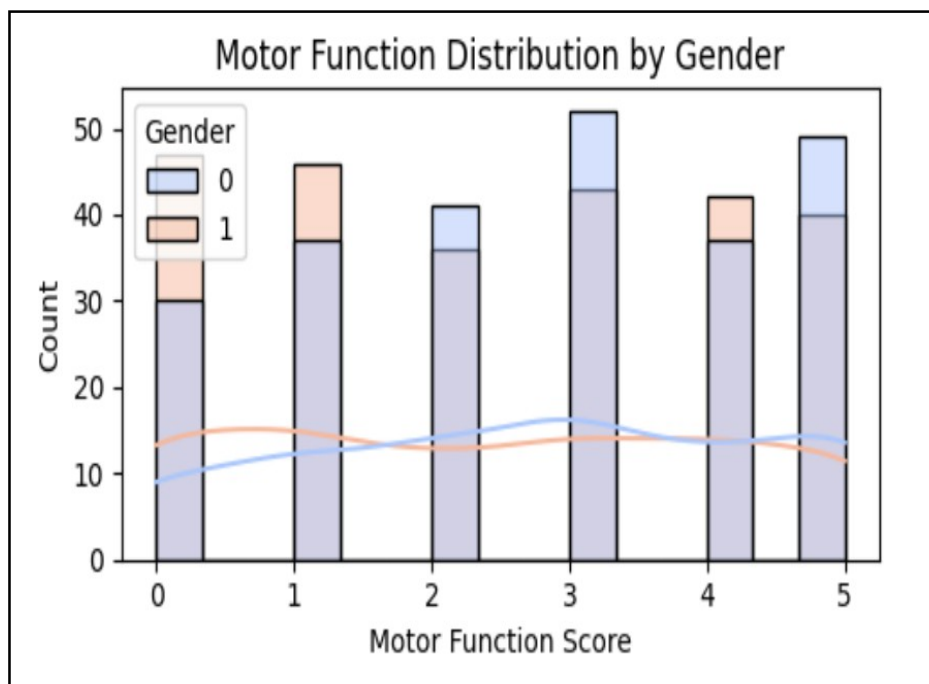


Fig1: Visualizing Feature Distributions with Histograms

B. Categorical observations

Bar charts aided us in examining categorical variables such as *Gender* and *Medications*. As a result, the dataset showed a near-equal gender distribution. This indicates a balanced sample of data. When visualized against severity-related variables, grouped bar charts revealed a slightly higher average tremor and motor difficulty in male patients. Medication categories varied, with some combinations more common among patients with higher disease progression.

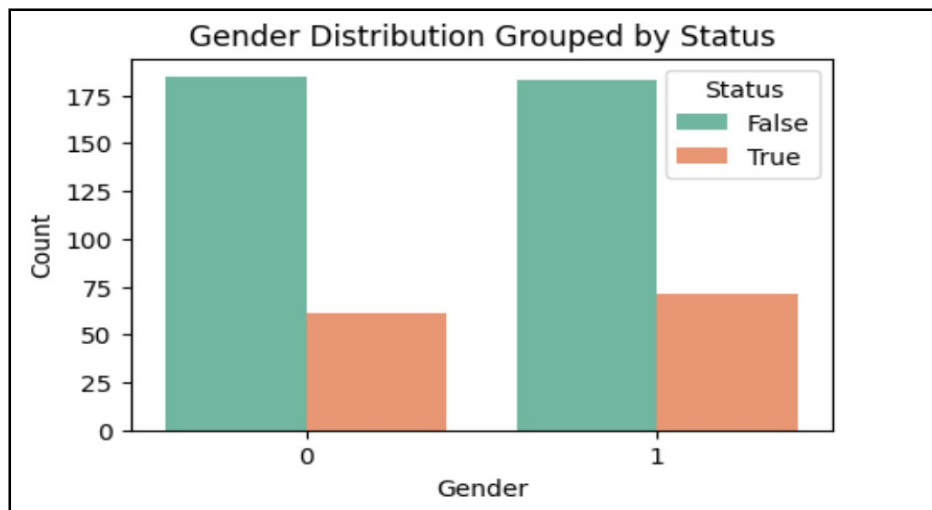


Fig 2: Gender Distribution

C. Correlation Analysis

A correlation heat-map was generated to present linear relationships among numerical features. UPDRS Score and Motor Function showed a strong positive correlation, confirming motor symptoms as a key progression marker. *Years Since Diagnosis* also had a moderate correlation with disease progression which was the target variable, corresponding with clinical expectations of a gradual worsening over time.

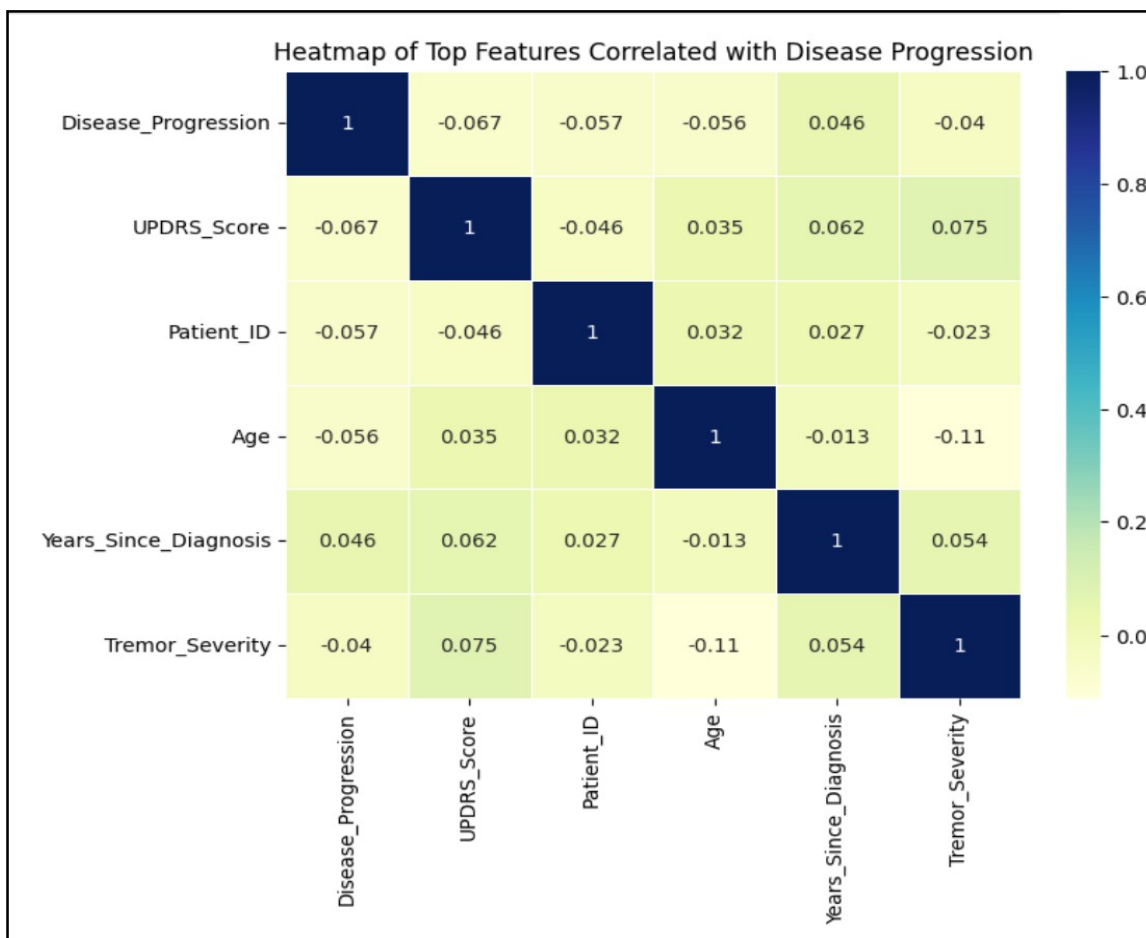


Fig 3 : Correlation heat map

D. Multivariate Visualizations

Pairplots were created to explore the interactions between multiple clinical features. Overlapping clusters between *Motor Function*, *Speech Difficulty* and *Balance Problems* represented cooccurring symptom progression. Scatter plots showed a spread of *UPDRS Scores* across varying levels of symptom severity, indicating heterogeneous disease expression across patients.

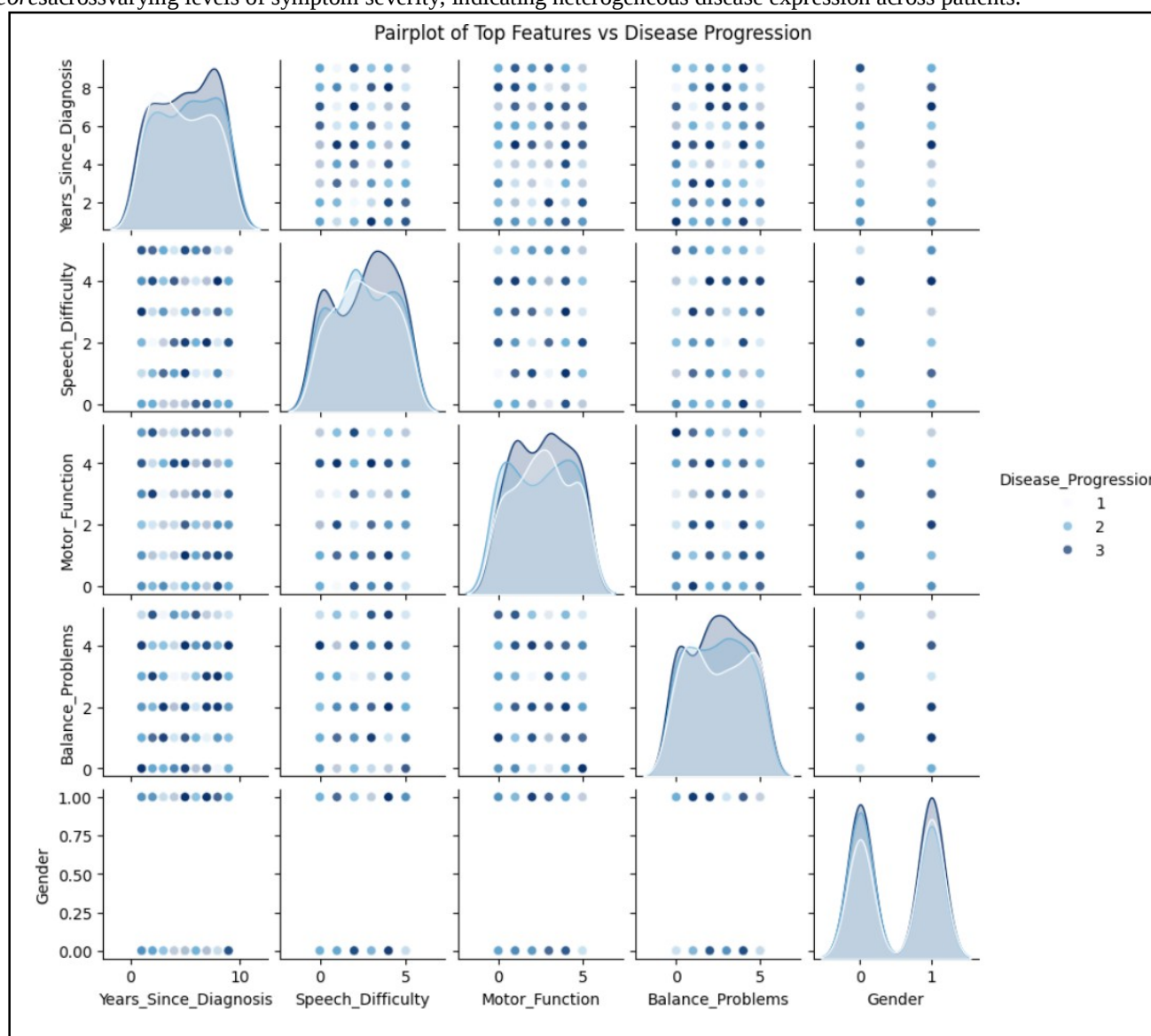


Fig4: Pair-plot to top Features

E. Temporal Trends

A consistent upward trend was clearly illustrated between *Years Since Diagnosis* and *UPDRS Score* in a line plot. Significantly higher symptom scores were observed with the patients with a longer disease duration. This emphasizes the progressive nature of Parkinson's disease and confirms that diagnosis duration is a key predictor of clinical severity. The symmetry of the violins indicates a roughly normal distribution for both groups, with few outliers at the lower end of progression scores.

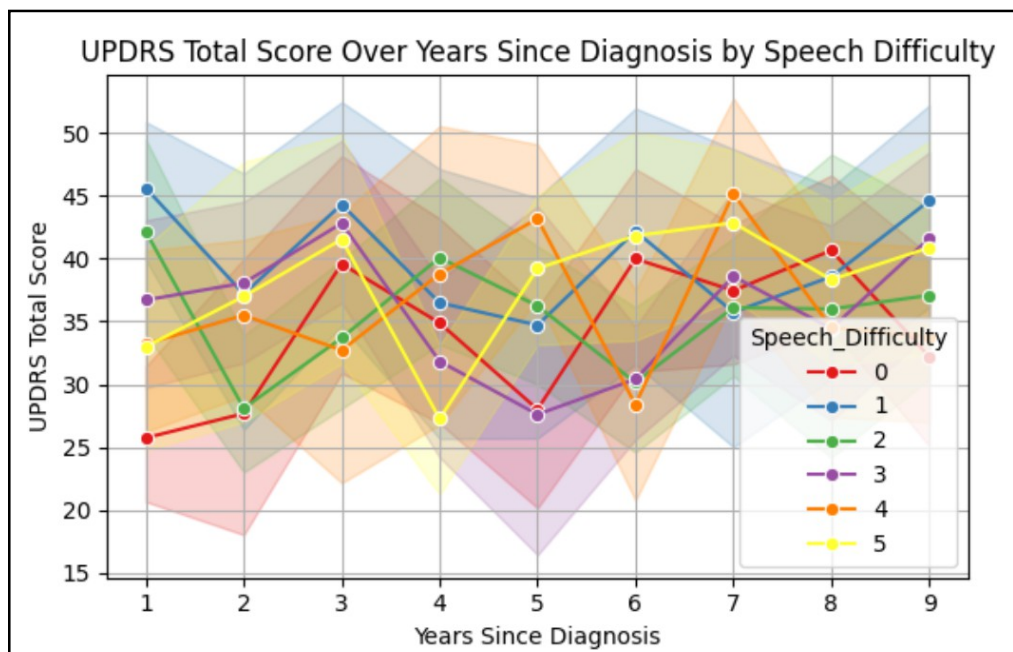


Fig5:years SinceDiagnosis &UPDRS Scoreline plot

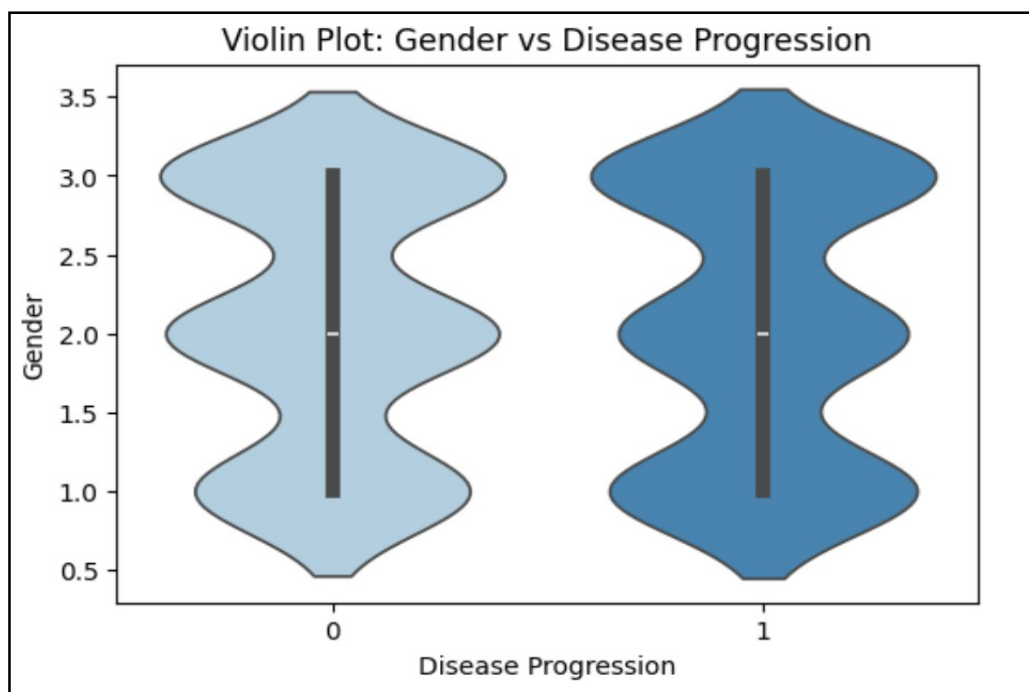


Fig6:Violin Plot

F. Summary of Findings

The data visualization performed through this Exploratory Data Analysis project acknowledge several meaningful observations. The overall disease progression was strongly affected by features such as *Motor Function* and *Speech Difficulty*. Gender and exercise patterns provided to variability in symptom progression. Thus, the combination of statistical and visual methods successfully highlighted key symptom collections and temporal trends, proposing valuable input for further clinical research or predictive modelling.



G. Implications for Clinical Tools

Our correlation heatmaps and temporal trends could inform the design of a lightweight Python dashboard for clinicians to track symptom progression. For example, integrating these visualizations with wearable sensor data (e.g., tremor frequency) could enable real-time monitoring.

IV. CONCLUSION

This current study presented an Exploratory Data Analysis of Parkinson's Disease progression using a medical dataset comprising 500 patients. By applying Python-based data analysis methods, we were able to clean, preprocess and visualize the dataset to discover important trends and symptom interdependencies. A strong correlation was exhibited between features such as UPDRS Score, Motor Function and Balance Problems, revealing how these symptoms worsen parallelly as the disease progresses. Visual tools such as histograms, bar plots, correlation heatmaps, pair plots and line graphs showed valuable insights into how various features interact over time. Clinical patterns of Parkinson's Disease were further supported by gender-based comparisons and trend analysis. While this study was restricted to exploratory analysis of the data and did not include predictive modeling, the results demonstrate the effectiveness of EDA in understanding symptom progression of Parkinson's disease. In future work, machine learning models can be applied to predict disease progression and classify severity levels based on patient data. The outcome of this study serves as a helpful basis for more advanced data-driven approaches in Parkinson's research and healthcare planning.

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