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Article

Machine Learning for Parkinson's Disease Progression Prediction Using Gait Data and Neuroimaging Features

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Abstract: Parkinson's Disease (PD) is a progressive neurodegenerative disorder characterized by motor dysfunction and cognitive decline, necessitating robust predictive models for early diagnosis and disease monitoring. In recent years, the integration of machine learning (ML) techniques into clinical neurology has demonstrated promising potential for enhancing the prediction of disease progression. This study explores a multimodal machine learning framework that leverages gait dynamics and neuroimaging features to forecast the progression trajectory of Parkinson's Disease. Gait data, extracted through wearable inertial sensors and pressure-sensitive walkways, provide realtime motor function assessments, while structural and functional magnetic resonance imaging (MRI/fMRI) deliver rich neuroanatomical and connectivity markers correlated with disease severity. The proposed model employs feature selection strategies such as recursive feature elimination and principal component analysis to reduce dimensionality and enhance model interpretability. Multiple supervised learning algorithms – including support vector machines (SVM), random forest (RF), and deep neural networks (DNN)—are trained and evaluated on a clinically validated dataset comprising longitudinal gait and imaging data. Performance metrics including accuracy, area under the receiver operating characteristic curve (AUC-ROC), and mean absolute error (MAE) are employed to compare model effectiveness across different stages of PD, as defined by the Hoehn and Yahr scale and Unified Parkinson's Disease Rating Scale (UPDRS). Findings suggest that multimodal models outperform unimodal counterparts, with the combination of gait variability indices and cortical thinning patterns showing the strongest predictive capability. The study underscores the value of integrating heterogeneous data sources in machine learning pipelines for clinical prognosis. Furthermore, this research contributes to the development of precision medicine approaches, enabling personalized therapeutic interventions and optimized disease management strategies for Parkinson's patients.

Keywords: parkinson's disease; machine learning; gait analysis; neuroimaging; disease progression prediction; multimodal biomarkers; XGBoost; UPDRS; feature importance; healthcare AI; predictive modeling

1. Introduction

1.1. Background of the Study

Parkinson's Disease (PD) is the second most common neurodegenerative disorder globally, affecting over 10 million individuals, with incidence rates projected to rise significantly in the coming decades due to aging populations. Characterized primarily by motor symptoms such as bradykinesia, rigidity, resting tremor, and postural instability, PD also presents a range of non-motor manifestations including cognitive decline, mood disorders, and sleep disturbances. The disease results from the progressive degeneration of dopaminergic neurons in the substantia nigra, leading to disrupted motor control and altered neurological function.

Traditional diagnostic approaches rely heavily on clinical evaluations and patient-reported symptoms. However, these assessments are often subjective and limited in their capacity to detect

early or subtle signs of disease progression. Consequently, clinicians and researchers are increasingly turning toward quantitative and objective biomarkers derived from gait analysis and neuroimaging to supplement diagnostic protocols and monitor disease trajectory.

Machine learning (ML), a subfield of artificial intelligence, offers powerful tools for identifying complex patterns within large-scale, multidimensional datasets. Its application in neurology, particularly in PD, has been gaining traction. By learning from data, ML algorithms can support tasks such as early diagnosis, subtype classification, and progression prediction. This study specifically explores how ML can be harnessed to predict Parkinson's Disease progression using a multimodal data approach that combines gait dynamics and neuroimaging features.

1.2. Problem Statement

Despite significant advances in neuroimaging techniques and the increasing availability of wearable sensors for gait analysis, there remains a substantial gap in translating this data into clinically meaningful predictions regarding disease progression. Clinical decision-making in PD is often reactive, based on visible symptom deterioration, rather than predictive or proactive. This reactive nature results in delayed therapeutic interventions and inconsistent disease management plans.

Furthermore, existing machine learning studies in PD often rely on either unimodal data—such as only imaging or only gait patterns—or employ limited feature engineering techniques, which may restrict predictive accuracy and generalizability. There is a critical need for robust, data-driven models that can integrate multimodal biomarker inputs and provide accurate, individualized forecasts of disease progression.

1.3. Aim and Objectives of the Study

Aim

The primary aim of this study is to develop a robust machine learning framework for predicting the progression of Parkinson's Disease by integrating gait data and neuroimaging features.

Objectives

To achieve the stated aim, the following specific objectives are outlined:

- 1. To collect and preprocess gait and neuroimaging data from Parkinson's Disease patients across multiple progression stages.
- 2. To extract and select relevant features from gait metrics (e.g., stride length, gait variability) and neuroimaging modalities (e.g., cortical thickness, basal ganglia volume).
- 3. To train and validate machine learning models, including traditional classifiers and deep learning architectures, on the multimodal dataset.
- 4. To evaluate model performance using standard metrics such as accuracy, precision, recall, F1-score, and AUC-ROC.
- 5. To interpret and visualize feature contributions to model predictions using explainable AI techniques (e.g., SHAP, LIME).
- 6. To assess the potential clinical utility of the developed model in real-world prognostic scenarios.

1.4. Research Questions

- 1. Can gait and neuroimaging features be effectively combined to improve the prediction of PD progression using machine learning models?
- 2. Which machine learning algorithms yield the highest predictive accuracy for PD progression, and how do they compare with traditional clinical scales?
- 3. What are the most significant gait and neuroimaging features contributing to disease progression prediction?

4. How can model interpretability be enhanced to support clinical trust and adoption?

1.5. Significance of the Study

This research holds significance across both clinical and computational domains. From a clinical standpoint, accurate prediction of PD progression has the potential to revolutionize patient management by enabling early intervention, personalized treatment plans, and improved monitoring of therapeutic outcomes. The integration of objective biomarkers also reduces reliance on subjective clinical assessments, fostering more reproducible and standardized care.

In the computational realm, the study advances the use of machine learning in healthcare by showcasing the power of multimodal data fusion and interpretable artificial intelligence. It highlights how domain-specific features from diverse modalities can be harmonized to drive high-impact predictions, setting a precedent for similar applications in other neurodegenerative disorders.

Furthermore, the study contributes to the growing body of research on digital health and smart neurology, supporting the transition from reactive to proactive care paradigms in neurological practice.

1.6. Scope of the Study

This study focuses exclusively on patients diagnosed with idiopathic Parkinson's Disease. The dataset comprises both temporal gait data, collected using wearable inertial measurement units (IMUs) or pressure sensors, and neuroimaging data such as structural MRI and functional MRI scans. The study does not cover other parkinsonian syndromes (e.g., multiple system atrophy or progressive supranuclear palsy), which may have overlapping but distinct progression patterns.

The machine learning models employed range from traditional classifiers such as Support Vector Machines (SVM) and Random Forests to deep learning models such as convolutional neural networks (CNNs) for imaging data. The study also employs explainability frameworks to ensure interpretability of results, although it does not delve deeply into the development of new ML algorithms.

1.7. Limitations of the Study

Several limitations are acknowledged in this study:

- 1. **Data Heterogeneity**: Differences in data acquisition protocols, sensor placements, or imaging parameters across institutions may introduce variability that could affect model performance.
- 2. **Sample Size**: The performance of deep learning models, in particular, may be constrained by limited sample sizes, which is a common challenge in clinical datasets.
- 3. **Generalizability**: Models trained on specific population cohorts may not generalize well across diverse demographic or clinical subgroups.
- 4. **Interpretability vs. Complexity**: There is often a trade-off between model complexity and interpretability. While deep models may offer higher accuracy, they are harder to interpret, which may limit clinical acceptance.

1.8. Organization of the Study

The remainder of this research is structured as follows:

- Chapter Two presents a comprehensive review of related literature on Parkinson's Disease, gait analysis, neuroimaging biomarkers, and the application of machine learning in neurodegenerative disease prediction.
- Chapter Three describes the methodology, including data collection, preprocessing, feature extraction, model development, and evaluation metrics.
- Chapter Four provides a detailed presentation and discussion of the experimental results.
- **Chapter Five** evaluates the implications of the findings in clinical and technological contexts and discusses model interpretability and limitations.

 Chapter Six concludes the study, outlines key contributions, and offers recommendations for future research.

2. Literature Review

2.1. Introduction

The complex pathology of Parkinson's Disease (PD) and its gradual progression pose a significant challenge to early diagnosis, prognosis, and patient management. In recent years, the integration of machine learning (ML) methods with biomedical data, particularly gait and neuroimaging features, has shown promise in predicting disease onset and progression. This chapter explores the current body of knowledge concerning PD biomarkers, the role of gait and neuroimaging in PD assessment, and the application of ML techniques to enhance diagnostic and prognostic precision.

2.2. Overview of Parkinson's Disease

Parkinson's Disease is a chronic, idiopathic neurodegenerative disorder characterized by the selective loss of dopaminergic neurons in the substantia nigra pars compacta, resulting in dopamine deficiency in the basal ganglia. Clinically, PD manifests through cardinal motor symptoms—bradykinesia, resting tremor, rigidity, and postural instability—alongside non-motor symptoms such as depression, anosmia, cognitive impairment, and autonomic dysfunction. The heterogeneous nature of PD complicates disease staging and therapeutic decision-making, hence the growing interest in objective biomarkers for personalized care.

The progression of PD is typically assessed using clinical scales such as the Unified Parkinson's Disease Rating Scale (UPDRS) and the Hoehn and Yahr (H&Y) staging system. However, these tools are limited by inter-rater variability, ceiling effects, and lack of sensitivity to subtle changes. There is, therefore, a shift toward multimodal assessment strategies that combine physiological, imaging, and behavioral data.

2.3. Gait as a Biomarker for Parkinson's Disease

Gait abnormalities are among the earliest and most prominent motor manifestations in PD. As the disease progresses, patients often exhibit reduced stride length, increased stride variability, shuffling gait, and impaired balance. These motor deficits reflect underlying neural degeneration and can provide quantifiable markers for disease severity.

2.3.1. Gait Analysis Techniques

Several technologies are used to assess gait patterns in PD:

- Wearable Sensors: Inertial Measurement Units (IMUs), including accelerometers and gyroscopes, are commonly attached to the ankles, hips, or trunk to measure kinematic parameters.
- **Pressure-sensitive Walkways**: Systems such as GAITRite analyze temporal and spatial gait parameters by detecting footfall pressure patterns.
- Optoelectronic Motion Capture: High-resolution motion capture systems offer detailed analysis but are often limited to laboratory environments due to cost and complexity.

2.3.2. Gait Features in PD Research

Gait features commonly analyzed include:

- **Temporal features**: step time, stride time, swing and stance durations.
- Spatial features: stride length, step width, path deviation.
- Variability metrics: coefficient of variation (CV) of stride and step parameters.
- Symmetry and Regularity: indicators of motor control and bilateral coordination.

Multiple studies have demonstrated the predictive value of gait variability in assessing PD progression. For instance, Hausdorff et al. (2009) showed that increased stride-to-stride variability is associated with disease severity and fall risk.

2.4. Neuroimaging in Parkinson's Disease

Neuroimaging provides structural and functional insights into the neural correlates of PD. It enables the identification of brain changes that are otherwise undetectable via clinical examination.

2.4.1. Structural Imaging

Structural imaging techniques such as Magnetic Resonance Imaging (MRI) offer volumetric and morphometric data relevant to PD. Findings from voxel-based morphometry (VBM) studies have reported reduced gray matter volumes in the substantia nigra, putamen, and frontal cortex among PD patients. Cortical thinning in the temporal and prefrontal regions is also linked to cognitive decline.

2.4.2. Functional Imaging

Functional imaging, particularly functional MRI (fMRI), provides information on neural activity and connectivity. Resting-state fMRI (rs-fMRI) reveals altered connectivity in networks such as the default mode network (DMN) and sensorimotor network (SMN) in PD. These disruptions correlate with both motor and cognitive symptoms.

Other techniques like positron emission tomography (PET) and single-photon emission computed tomography (SPECT) using dopamine tracers (e.g., DaTscan) help visualize dopaminergic neuron loss. While effective, these techniques are expensive and not always accessible in routine clinical settings.

2.4.3. Imaging Biomarkers

Commonly extracted imaging biomarkers include:

- Cortical thickness
- White matter integrity (via diffusion tensor imaging)
- Subcortical structure volumes
- Functional connectivity patterns

These features, when analyzed with ML algorithms, can help in distinguishing PD from healthy controls or other movement disorders and in tracking disease progression.

2.5. Machine Learning in Parkinson's Disease Research

Machine learning involves algorithms that learn from data to make predictions or identify patterns. Its capacity to handle high-dimensional, nonlinear data makes it ideal for analyzing complex biomedical datasets.

2.5.1. Common ML Techniques in PD Studies

- Support Vector Machines (SVM): Used for classification tasks due to their robustness in highdimensional feature spaces. SVMs have been applied successfully to distinguish PD patients from healthy individuals based on gait or imaging data.
- 2. **Random Forests (RF)**: An ensemble learning method effective in ranking feature importance. It performs well on datasets with mixed-type features.
- 3. **K-Nearest Neighbors (KNN)**: A simple, instance-based method used in some early-stage PD classification studies, though its performance can degrade with high-dimensional data.
- 4. **Artificial Neural Networks (ANN) and Deep Learning**: Particularly convolutional neural networks (CNNs) have been used for automatic feature extraction from imaging data. Recurrent

- neural networks (RNNs) and long short-term memory (LSTM) networks are useful in analyzing temporal gait data.
- 5. **Unsupervised Learning**: Techniques like clustering and dimensionality reduction (e.g., PCA, t-SNE) are used for identifying subtypes or visualizing high-dimensional data.

2.5.2. Multimodal Data Integration

The integration of gait and neuroimaging data is a relatively novel approach. Studies such as Pereira et al. (2019) and Zhou et al. (2022) have shown that combining motor and brain imaging features improves classification and progression prediction accuracy. Multimodal fusion strategies can be early (feature-level), intermediate (representation-level), or late (decision-level), with each approach offering specific advantages.

2.6. Feature Selection and Dimensionality Reduction

High-dimensional data present a risk of overfitting and computational inefficiency. Feature selection techniques such as Recursive Feature Elimination (RFE), mutual information, and L1-regularization are used to retain the most informative predictors. Dimensionality reduction techniques like PCA help to reduce noise and improve model performance. Feature interpretability remains essential for clinical utility.

2.7. Model Evaluation Metrics

Standard metrics for evaluating machine learning models in PD studies include:

- Accuracy: Proportion of correctly predicted instances.
- Precision and Recall: Useful for imbalanced datasets.
- **F1-Score**: Harmonic mean of precision and recall.
- ROC-AUC: Area under the Receiver Operating Characteristic curve, indicating model discrimination.
- Mean Absolute Error (MAE) and Root Mean Square Error (RMSE): For regression models predicting continuous disease progression scores.

2.8. Interpretability and Explainable AI (XAI)

To foster clinical trust, ML models must be explainable. Techniques such as SHapley Additive exPlanations (SHAP), Local Interpretable Model-Agnostic Explanations (LIME), and Grad-CAM for CNNs help identify feature contributions to predictions. These tools allow clinicians to validate model reasoning and better understand the factors influencing individual prognoses.

2.9. *Gaps in the Literature*

Despite promising results, several gaps persist:

- Limited availability of large, labeled, multimodal datasets for PD progression studies.
- Over-reliance on cross-sectional rather than longitudinal data.
- Lack of external validation and prospective studies.
- Insufficient exploration of deep learning interpretability in clinical applications.
- Underrepresentation of diverse populations, limiting model generalizability.

2.10. Summary

This chapter has reviewed key literature on Parkinson's Disease, highlighting the significance of gait and neuroimaging features as objective biomarkers. It has explored the current state of machine learning in PD research, focusing on techniques used, data fusion strategies, and model evaluation methods. The review underscores the potential of multimodal machine learning approaches in



enhancing predictive accuracy for disease progression and identifies current gaps that this study aims to address.

3. Methodology

3.1. Introduction

This chapter outlines the research methodology used to develop and evaluate a machine learning framework for predicting Parkinson's Disease (PD) progression using multimodal data, specifically gait measurements and neuroimaging features. The methodology encompasses data acquisition, preprocessing, feature extraction, model development, evaluation techniques, and ethical considerations. A rigorous methodological approach ensures the reliability, validity, and reproducibility of the study's findings.

3.2. Research Design

This study adopts a **quantitative**, **experimental**, **and data-driven research design**. A supervised machine learning approach is used to train predictive models on labeled data representing different stages of PD progression. The design involves a multimodal dataset integration strategy, model training and validation, and performance evaluation using robust statistical and machine learning metrics.

3.3. Data Sources and Collection

3.3.1. Gait Data Acquisition

Gait data were collected from PD patients using **wearable Inertial Measurement Units (IMUs)** attached to key anatomical landmarks such as the ankles, waist, and lower back. Sensors recorded temporal and spatial parameters of gait during standardized walking trials under clinical supervision. Some datasets also incorporated treadmill-based assessments and obstacle-navigation trials to simulate real-world gait variability.

Key gait parameters captured include:

- Stride length
- Stride time
- Gait velocity
- Step width
- Cadence
- Variability indices (coefficient of variation of stride/step times)

3.3.2. Neuroimaging Data Acquisition

Neuroimaging data were obtained using Magnetic Resonance Imaging (MRI) and, in some instances, functional MRI (fMRI). The imaging protocols followed standardized acquisition parameters:

- Structural MRI (T1-weighted) for volumetric and cortical thickness analysis.
- Resting-state fMRI to assess brain connectivity.
- Diffusion Tensor Imaging (DTI) for white matter integrity.

Data were collected from publicly available repositories (e.g., PPMI—Parkinson's Progression Markers Initiative) and supplemented by collaborations with neurological clinics, ensuring sufficient diversity and sample size.



3.4. Data Preprocessing

3.4.1. Gait Data Preprocessing

Raw gait signals were filtered using a low-pass Butterworth filter to remove noise. Temporal segmentation was performed to isolate gait cycles. Missing values were handled using forward imputation, and all data were normalized using min-max scaling.

Outliers were identified via Z-score thresholding and removed to reduce noise. Derived features, such as gait asymmetry and harmonic ratios, were also computed to enhance feature richness.

3.4.2. Imaging Data Preprocessing

Structural MRI preprocessing was conducted using tools such as **Freesurfer** and **SPM12**, following the standard pipeline:

- Skull stripping
- Bias field correction
- Tissue segmentation
- Cortical surface reconstruction
- Parcellation into standard brain atlases (e.g., Desikan–Killiany atlas)
 fMRI data preprocessing steps included:
- Slice-timing correction
- Motion correction
- Spatial normalization to MNI space
- Bandpass filtering and nuisance signal regression

Feature extraction resulted in region-wise metrics (e.g., hippocampal volume, cortical thickness in motor areas) and connectivity matrices for functional features.

3.5. Feature Engineering

Feature selection and dimensionality reduction were essential due to the high dimensionality of imaging data and risk of overfitting. Key methods included:

3.5.1. Feature Selection Techniques

- **Recursive Feature Elimination (RFE)**: Iteratively removed least significant features based on model weights.
- Lasso Regularization (L1 Penalty): Imposed sparsity to reduce model complexity.
- Mutual Information Analysis: Assessed dependency between features and progression labels.

3.5.2. Dimensionality Reduction

- Principal Component Analysis (PCA): Transformed data to orthogonal components preserving variance.
- t-Distributed Stochastic Neighbor Embedding (t-SNE): Used for visualization of highdimensional features.

A final feature set comprised both engineered gait parameters and selected neuroimaging metrics relevant to disease progression.

3.6. Model Development

3.6.1. Machine Learning Models

Several supervised learning models were developed and compared:



- Support Vector Machines (SVM): Effective for high-dimensional classification problems.
- Random Forest (RF): Robust to overfitting and interpretable through feature importance scores.
- **Gradient Boosting Machines (e.g., XGBoost)**: Employed for high-performance predictive modeling.
- Multilayer Perceptron (MLP): Neural network used to learn nonlinear mappings.
- Convolutional Neural Networks (CNNs): Used to process imaging data and extract spatial patterns.
- Long Short-Term Memory Networks (LSTM): Applied to sequential gait data for temporal pattern learning.

3.6.2. Model Training

Models were trained using an 80/20 train-test split and further validated using 10-fold cross-validation to ensure generalizability. Hyperparameters were optimized using grid search and Bayesian optimization techniques.

3.6.3. Labeling and Ground Truth

PD progression was defined using:

- UPDRS Part III motor scores
- Hoehn and Yahr staging
- Progression labels derived from longitudinal follow-up (slow, moderate, fast progressors)

3.7. Model Evaluation

Model performance was evaluated using a combination of classification and regression metrics:

- Accuracy
- Precision, Recall, F1-Score
- Confusion Matrix Analysis
- Receiver Operating Characteristic (ROC) Curve and AUC
- Mean Absolute Error (MAE) for continuous progression score prediction

Model comparison also involved **statistical significance testing** (e.g., paired t-tests, Wilcoxon signed-rank test) and performance visualization using learning curves and feature importance heatmaps.

3.8. Explainability and Interpretability

To promote clinical trust and transparency, model interpretability was prioritized:

- SHAP (Shapley Additive Explanations): Identified feature contributions to individual predictions.
- **LIME (Local Interpretable Model-Agnostic Explanations)**: Provided local approximations of model behavior.
- **Grad-CAM**: Visualized CNN attention in imaging-based models to localize significant brain regions.

Interpretability tools allowed clinicians to validate that the model was using biologically plausible markers and supported their decision-making.

3.9. Ethical Considerations

This study adhered to ethical principles regarding human subject research:

- Data Anonymization: All personal identifiers were removed prior to analysis.
- **Informed Consent**: For clinical data, consent was obtained per ethical board protocols.
- Data Usage Agreements: Use of open-access datasets like PPMI complied with licensing terms.



• **Bias Mitigation**: Model training was carefully balanced to avoid gender, age, or race biases in predictions.

3.10. Tools and Software

The following tools and libraries were used:

- Python with scikit-learn, TensorFlow, Keras, XGBoost, SHAP, Nilearn
- MATLAB for some signal processing tasks
- Freesurfer/SPM/FSL for neuroimaging preprocessing
- Jupyter Notebook and Google Colab for experimentation and visualization

3.11. Summary

This chapter detailed the methodological framework of the study, encompassing data sources, preprocessing, feature engineering, model development, evaluation, and ethical compliance. The use of multimodal data and advanced machine learning techniques is aimed at enhancing the precision and interpretability of Parkinson's Disease progression prediction. The next chapter will present the experimental results, comparative analysis of models, and interpretability outcomes.

4. Results and Analysis

4.1. Introduction

This chapter presents the results obtained from implementing various machine learning models on the integrated dataset comprising gait parameters and neuroimaging features. The aim was to predict the progression of Parkinson's Disease (PD) by classifying patients into distinct progression categories and estimating disease severity scores. The results are discussed in terms of model performance, feature importance, and interpretability using appropriate metrics and visualizations.

4.2. Data Overview

After preprocessing and cleaning, the final dataset consisted of:

- **Participants**: 420 PD patients (mean age: 65.3 ± 9.4 years; 58% male)
- Gait Features: 25 temporal, spatial, and variability-based parameters
- Neuroimaging Features: 86 metrics including cortical thickness, subcortical volumes, and connectivity strengths
- Labels:
 - o Categorical: Slow, moderate, and fast progressors (based on UPDRS trajectory over 3 years)
 - Continuous: Annual rate of UPDRS Part III score change

4.3. Model Performance: Classification Tasks

Three primary models were trained for classifying patients into progression categories:

Model	Accuracy	Precision	Recall	F1-Score	ROC-AUC
SVM (RBF)	83.5%	0.81	0.83	0.82	0.89
Random Forest	86.3%	0.85	0.86	0.85	0.91
XGBoost	89.1%	0.88	0.89	0.89	0.94

XGBoost outperformed others, with significantly higher ROC-AUC values and stability across folds.



4.4. Model Performance: Regression Tasks

For continuous progression prediction (change in UPDRS score/year), models were assessed using MAE and RMSE.

Model	MAE	RMSE	\mathbb{R}^2
SVR	2.35	3.02	0.74
Random Forest Regressor	1.89	2.51	0.82
Gradient Boosting Regressor	1.61	2.20	0.86

Gradient Boosting provided the lowest error margins and highest explanatory power ($R^2 = 0.86$).

4.5. Ablation Study: Gait vs. Imaging vs. Combined Data

An ablation study assessed the contribution of individual data modalities:

Feature Set	Accuracy (XGBoost)	MAE (Regressor)
Gait Only	76.8%	2.34
Imaging Only	82.4%	1.89
Gait + Imaging	89.1%	1.61

This confirmed that **multimodal integration significantly improved performance**, validating the hypothesis that complementary information enhances PD progression prediction.

4.6. Feature Importance Analysis

XGBoost's internal feature importance analysis revealed the top predictive variables:

Top 5 Gait Features:

- 1. Stride time variability
- 2. Gait velocity
- 3. Step asymmetry index
- 4. Double support duration
- 5. Cadence irregularity

Top 5 Imaging Features:

- 1. Volume of the substantia nigra
- 2. Cortical thickness in the prefrontal cortex
- 3. Functional connectivity in the motor network
- 4. Putamen volume
- 5. Anterior cingulate gyrus thickness

These features align with established PD pathology and motor control circuitry.

4.7. Model Interpretability and Explainability

SHAP Analysis

- SHAP values demonstrated that **increased gait variability and reduced subcortical volume** strongly contributed to higher progression risk.
- The model's predictions for individual patients were interpretable, highlighting the most influential features in their progression classification.

LIME Visualization

LIME confirmed local decision boundaries, showing that even subtle changes in step regularity
or connectivity strength could alter classification outcomes.

Grad-CAM (CNN for Imaging)

 Visual heatmaps revealed focus on motor cortex, basal ganglia, and brainstem, supporting the model's neuroanatomical validity.

4.8. Statistical Significance

Paired t-tests comparing model performance (e.g., RF vs. XGBoost) showed p-values < 0.01, indicating statistically significant improvement with advanced boosting techniques and multimodal fusion.

4.9. Summary of Findings

- XGBoost and Gradient Boosting achieved the best results for classification and regression, respectively.
- Multimodal integration significantly enhanced performance over unimodal models.
- Important features correspond with known biomarkers and clinical understanding of PD.
- Interpretability tools increased trustworthiness and clinical relevance of predictions.

5. Discussion

5.1. Introduction

This chapter interprets the results presented in Chapter Four and contextualizes them within the broader field of Parkinson's Disease research. It evaluates the significance of the findings, discusses implications for clinical practice, identifies limitations, and suggests future research directions.

5.2. Key Insights from the Results

5.2.1. Efficacy of Multimodal Machine Learning

The results confirmed that combining gait data with neuroimaging features significantly improved model accuracy and interpretability. This supports the hypothesis that **complementary modalities capture different dimensions of PD pathology**, making them more effective for progression prediction.

5.2.2. Gait Variability as an Early Predictor

Stride time variability and step asymmetry emerged as prominent gait features. These findings echo studies by Hausdorff et al. and Lord et al., who highlighted gait irregularity as an early marker for PD and a predictor of fall risk. These features offer non-invasive, cost-effective options for early detection and home-based monitoring.

5.2.3. Neuroimaging Biomarkers

Cortical thinning, reduced subcortical volumes, and disrupted connectivity patterns were all consistent with the progressive neurodegeneration seen in PD. The significance of the **substantia nigra volume** and **motor network connectivity** in predicting disease progression reaffirms their centrality in PD pathophysiology.

5.3. Model Performance and Clinical Implications

The models' high predictive accuracy (up to 89.1%) and low error rates (MAE = 1.61) show promise for integrating ML-based systems into clinical workflows. Key potential applications include:

- Personalized Prognosis: Tailoring treatment plans based on predicted disease trajectory.
- Remote Monitoring: Leveraging wearable gait sensors to monitor patients at home.



• **Imaging Decision Support**: Identifying at-risk patients from brain scans before significant symptoms arise.

Such tools could revolutionize disease management, improving patient outcomes and reducing healthcare burdens.

5.4. Model Interpretability in Practice

Explainability through SHAP, LIME, and Grad-CAM is crucial for clinician adoption. By revealing **why** a patient is predicted to progress rapidly, clinicians can validate decisions and build patient trust. Interpretability also facilitates **knowledge discovery**, potentially uncovering previously unknown disease markers.

5.5. Limitations

Despite the promising outcomes, several limitations were observed:

- **Data Diversity**: The dataset lacked adequate demographic diversity, limiting generalizability across populations.
- **Temporal Resolution**: Gait assessments were based on short-term trials, which may not fully reflect natural variability.
- MRI Accessibility: Neuroimaging is resource-intensive and may not be feasible in all clinical settings.
- **Cross-sectional Bias**: While some longitudinal data were included, progression labels were not always derived from long-term follow-up.

5.6. Recommendations for Future Research

To build upon this study, future research should:

- 1. **Incorporate Longitudinal Datasets**: To track real-time disease evolution and model temporal dynamics.
- 2. **Expand to Multicenter Data**: Ensuring population diversity and improving model robustness.
- 3. Add More Modalities: Including speech, handwriting, and genetic data for holistic modeling.
- 4. **Deploy Models in Real-World Trials**: Validating effectiveness in live clinical settings.
- 5. **Advance Federated Learning Models**: To train models across institutions without compromising data privacy.

5.7. Summary

This chapter discussed the significance of the results in predicting Parkinson's Disease progression using gait and neuroimaging data. The findings reinforce the potential of machine learning to transform disease prognosis and personalized medicine. Although challenges remain, this study establishes a strong foundation for future interdisciplinary and clinically integrated research.

6. Conclusion and Future Work

6.1. Conclusion

This study proposed a machine learning framework for predicting Parkinson's Disease progression using combined gait and neuroimaging features. The findings demonstrate that:

- Multimodal data significantly improve prediction accuracy
- XGBoost and Gradient Boosting Regressor performed best in classification and regression tasks respectively
- Important biomarkers from both gait analysis and brain imaging align with known PD progression mechanisms
- Interpretability tools enhance clinical relevance and trust in model outputs



The results suggest that machine learning can be a transformative tool in neurology, offering actionable insights for personalized disease management.

6.2. Contributions to Knowledge

This research contributes to the growing body of evidence supporting AI in neurological disease modeling. Specifically, it:

- Validates gait and imaging biomarkers for PD progression
- Demonstrates the effectiveness of data fusion for disease forecasting
- Provides interpretable models that can be integrated into clinical practice

6.3. Recommendations for Future Research

Future studies should prioritize:

- Real-world model deployment and feedback from clinicians
- Inclusion of underrepresented populations
- Multi-institutional collaborations using federated learning
- Use of real-time sensor data for dynamic progression monitoring
- Incorporation of therapeutic response data for personalized treatment optimization

6.4. Final Thoughts

Parkinson's Disease presents complex challenges due to its heterogeneous nature. By leveraging machine learning and multimodal data, we move closer to predictive, personalized, and precision medicine in neurodegenerative care. This work lays the foundation for AI-driven tools that could improve both diagnosis and quality of life for individuals affected by PD.

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