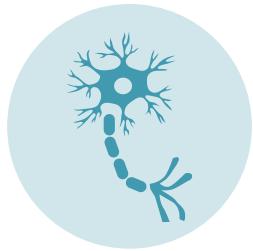


Parkinson's Disease Predictive Machine Learning Model

Luna Pérez Troncoso

Introduction

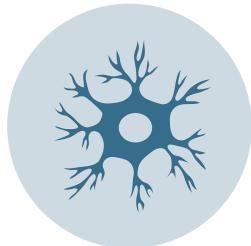


The **early detection of PD** is a **growing priority** within both clinical practice and research.

Introduction



The early detection of PD is a growing priority within both clinical practice and research.

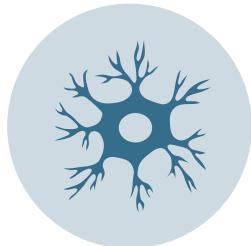


Many individuals remain undiagnosed until the disease has already progressed, limiting the effectiveness of available therapeutic interventions.

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Introduction



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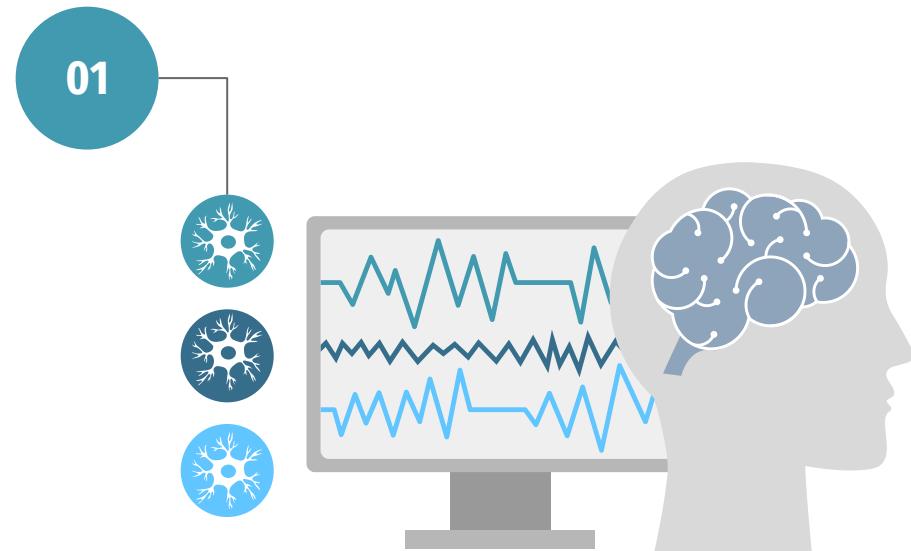
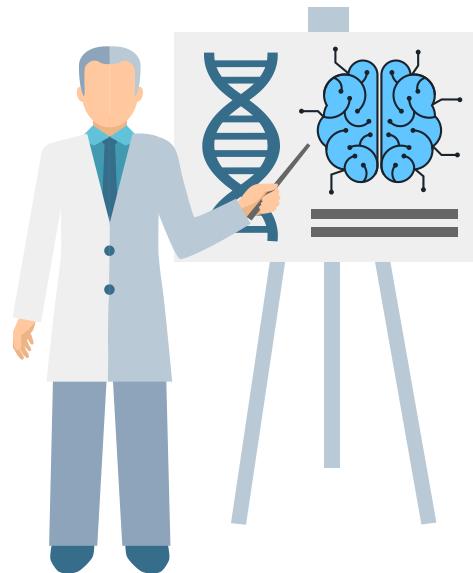
Many individuals remain undiagnosed until the disease has already progressed, limiting the effectiveness of available therapeutic interventions. Consequently, timely diagnosis can significantly influence patient outcomes and long-term quality of life.



Early identification could allow for **timelier monitoring, lifestyle adjustments, and targeted therapeutic strategies** that may slow **disease progression or improve quality of life**.

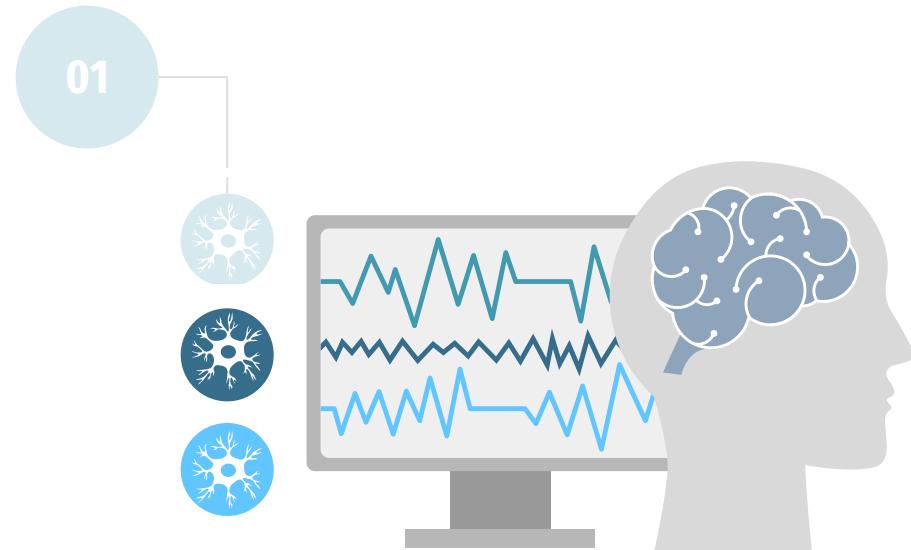
Why develop a predictive model?

A reliable predictive system has the potential to support clinicians in recognizing subtle signs that might otherwise go unnoticed.



Why develop a predictive model?

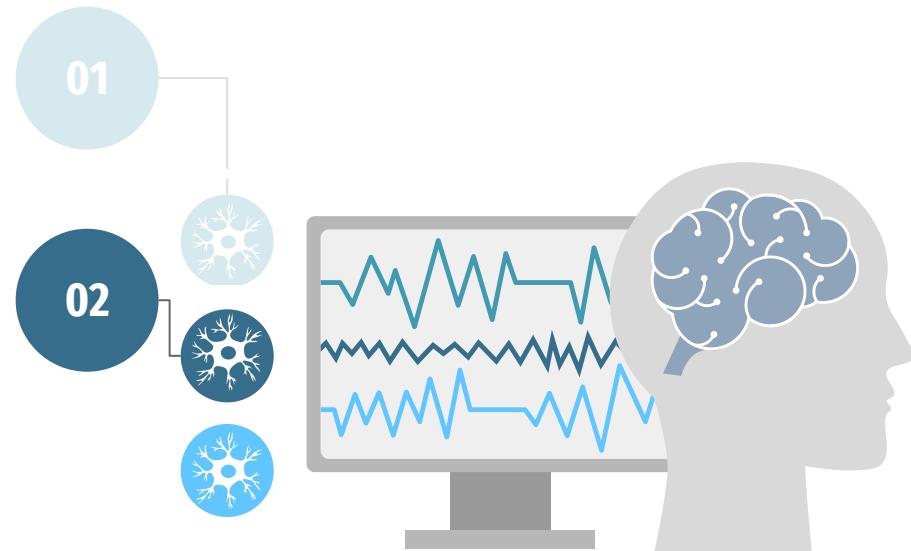
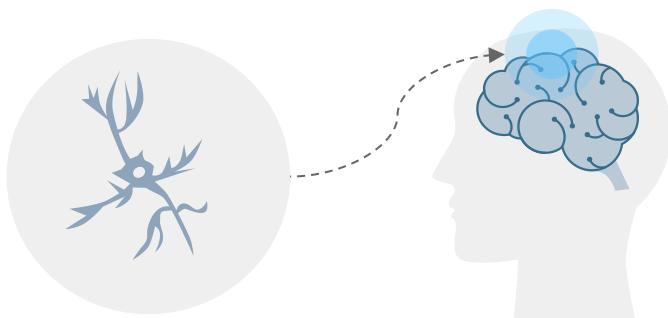
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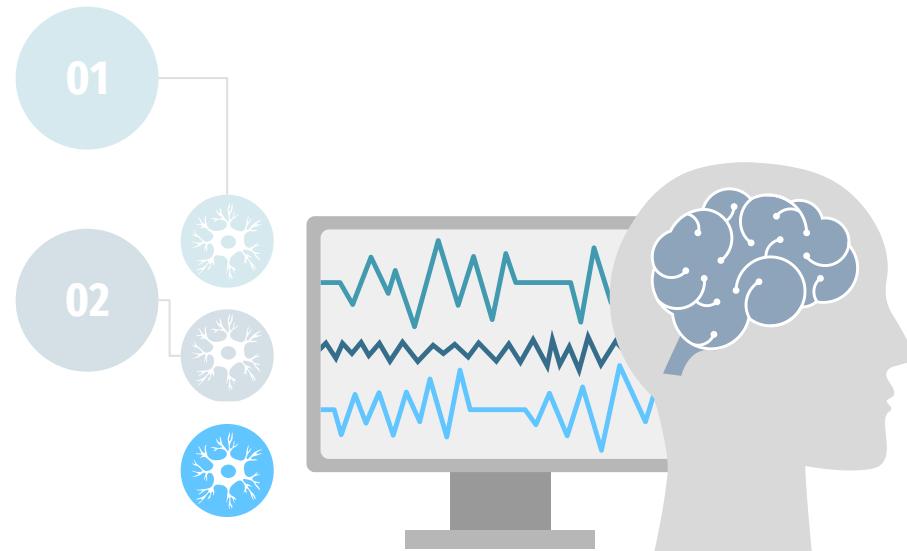
Predictive modeling can help researchers **gain deeper insight into the complex interactions that contribute to the onset of neurodegenerative disorders**



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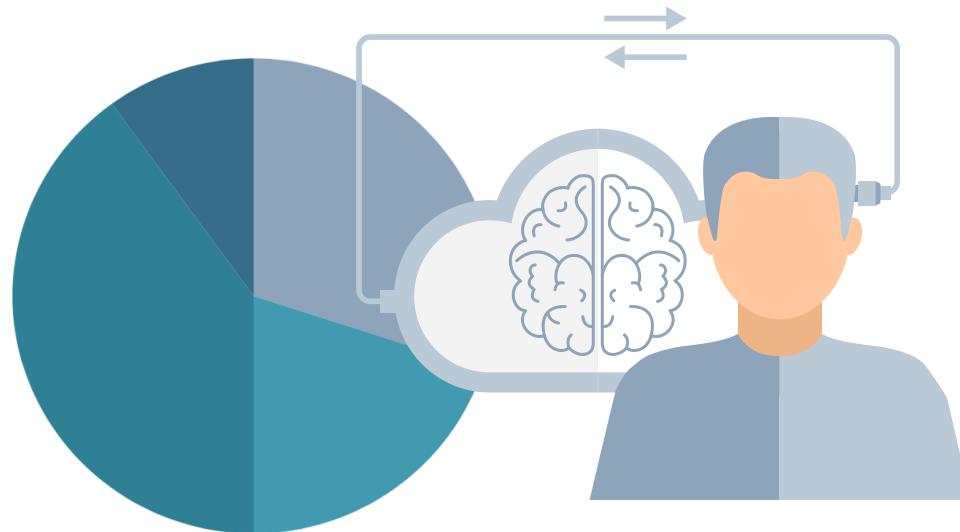
Predictive modeling can help researchers gain deeper insight into the complex interactions that contribute to the onset of neurodegenerative disorders

Beyond clinical impact, creating a predictive model encourages the integration of modern **data-driven approaches** into neurological healthcare, which stands out as a promising path to **more personalized and proactive patient care**.

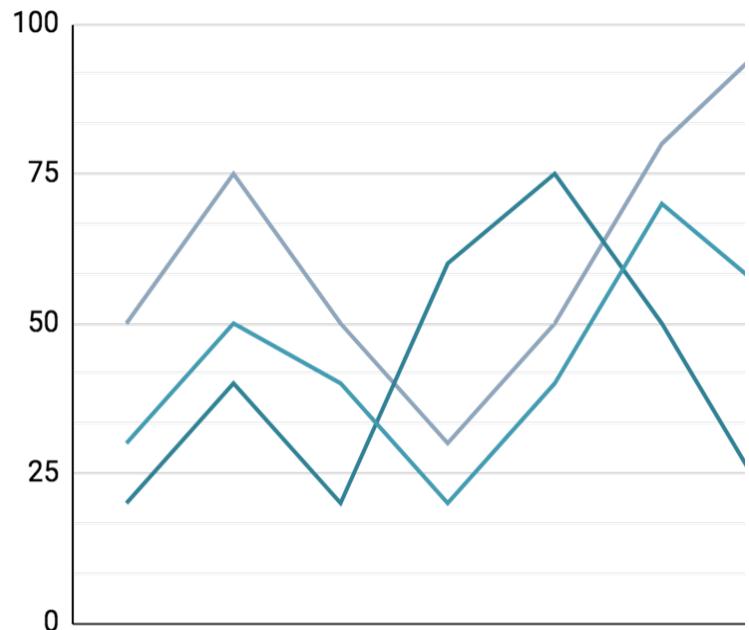


Objective

Integrating diverse variables (demographic, lifestyle, clinical, cognitive, and symptom-related variables) into a unified predictive framework, the project seeks to **evaluate multiple machine learning algorithms and determine their capability to accurately identify patients at risk.**

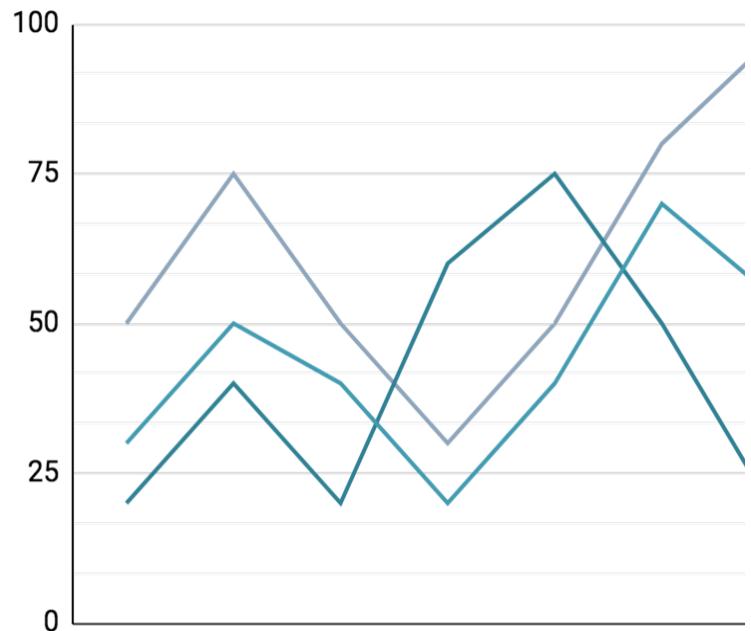


Data Description and Sources



As part of this project, I selected a synthetic dataset from Kaggle generated by Mr. Rabie El Kharoua, to support the development of a predictive model for Parkinson's disease.

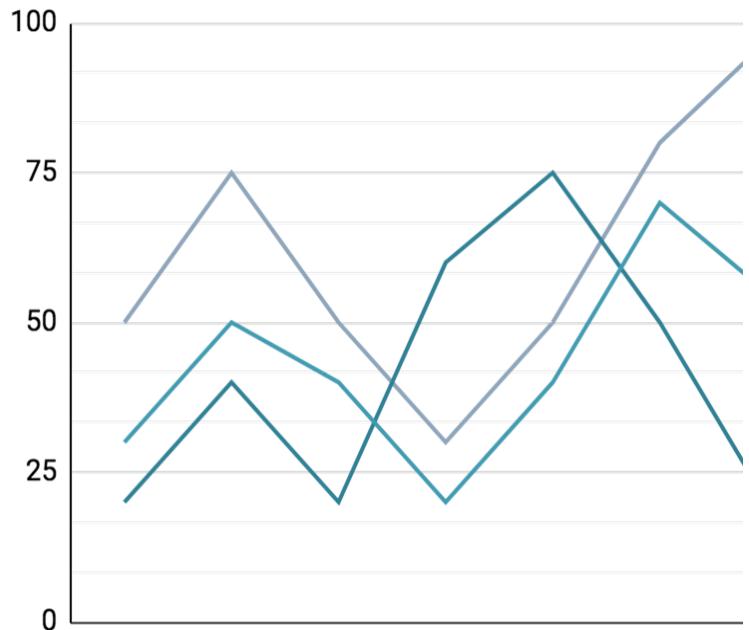
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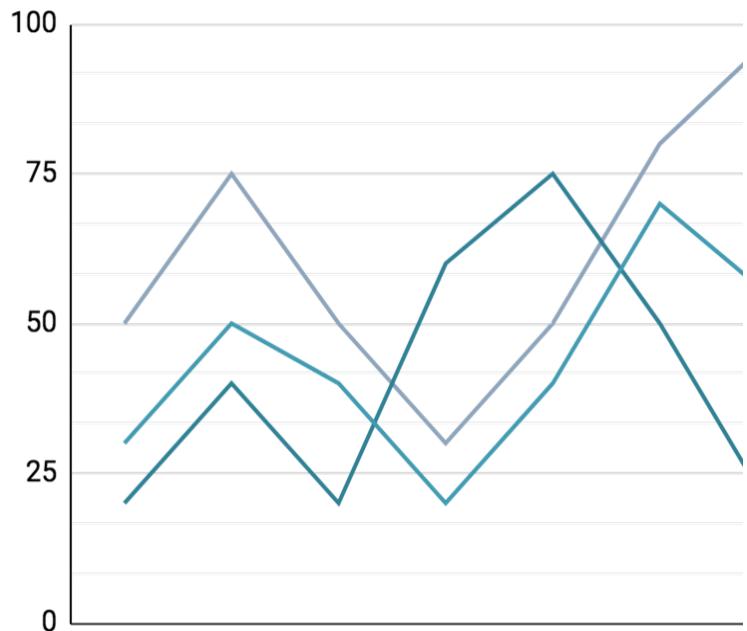


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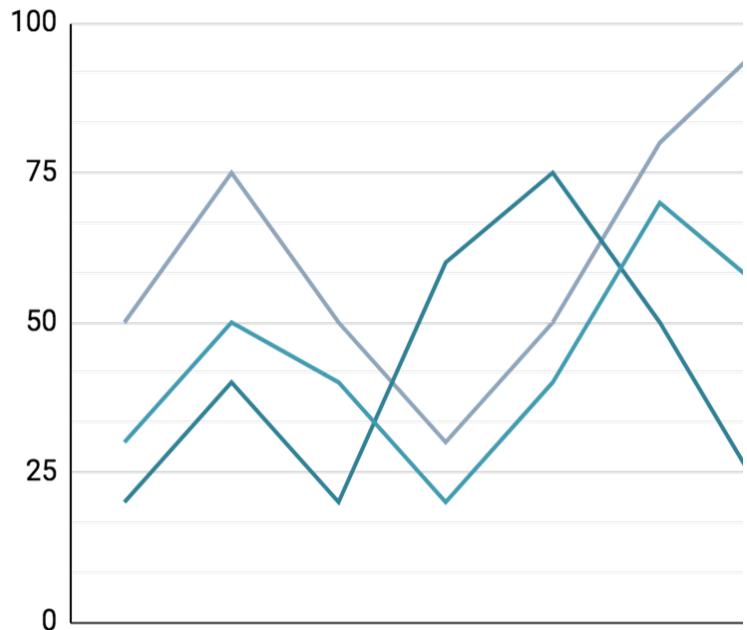


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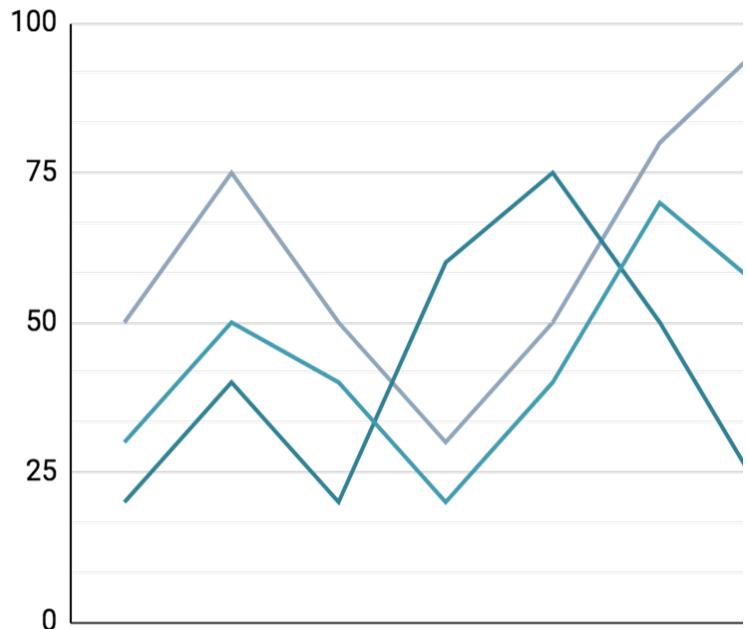


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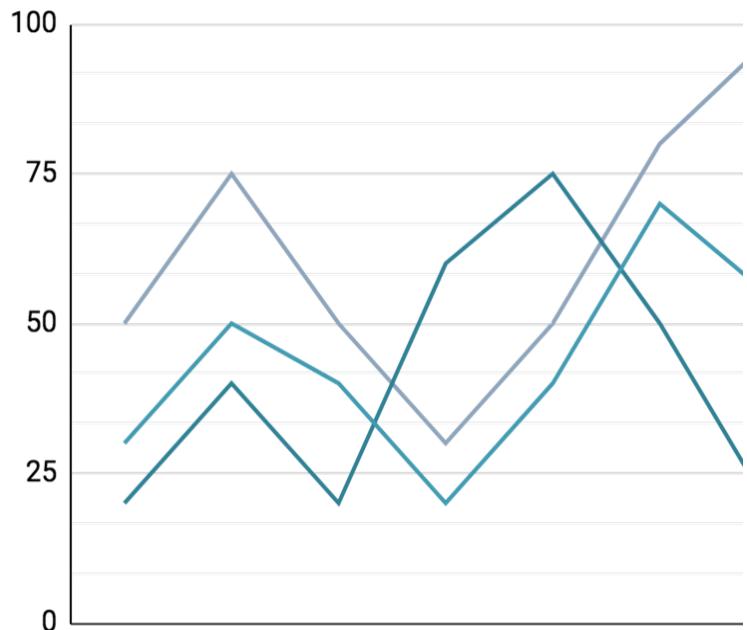


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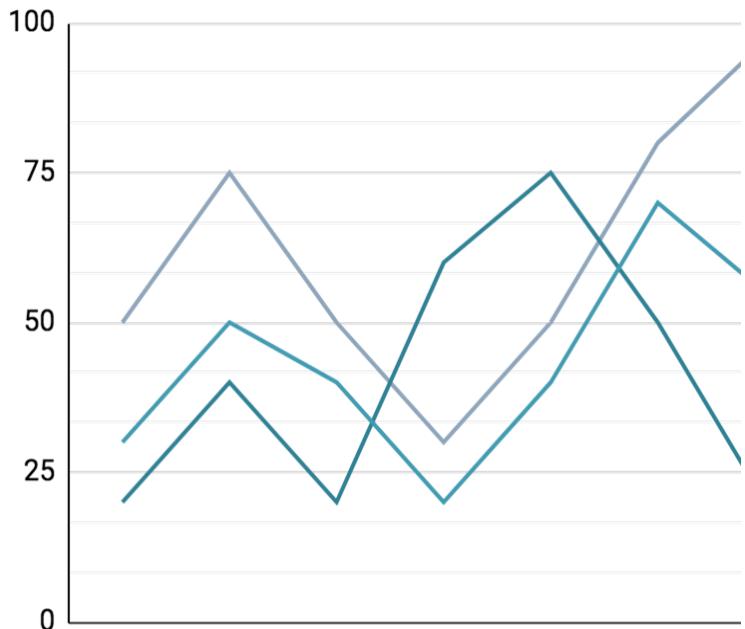


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- **Symptom indicators** (Tremor, Constipation, Rigidity).

Methodology

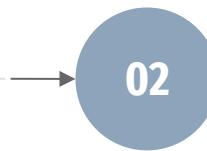
01

Standarization/Scaling

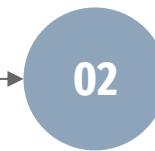
StandarScaler

MinMaxScaler or None

Methodology

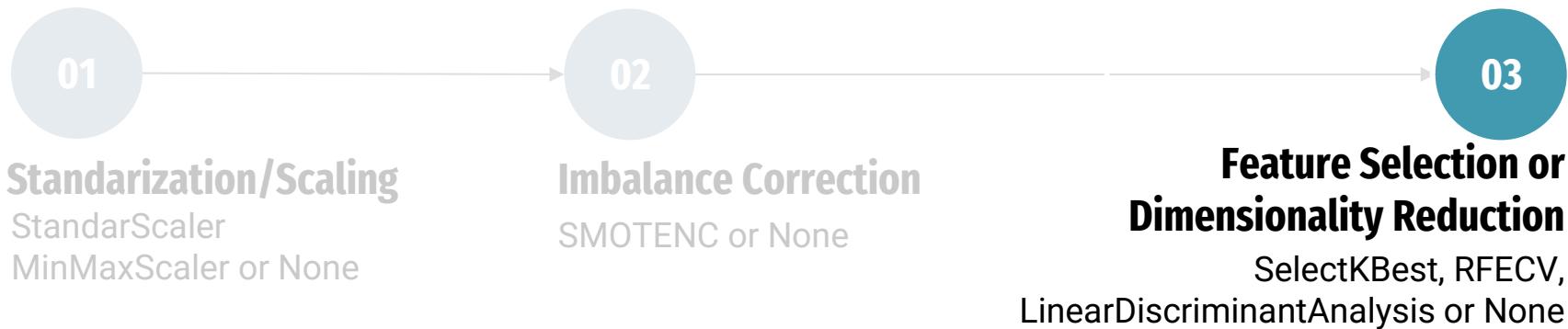


Standardization/Scaling
StandarScaler
MinMaxScaler or None

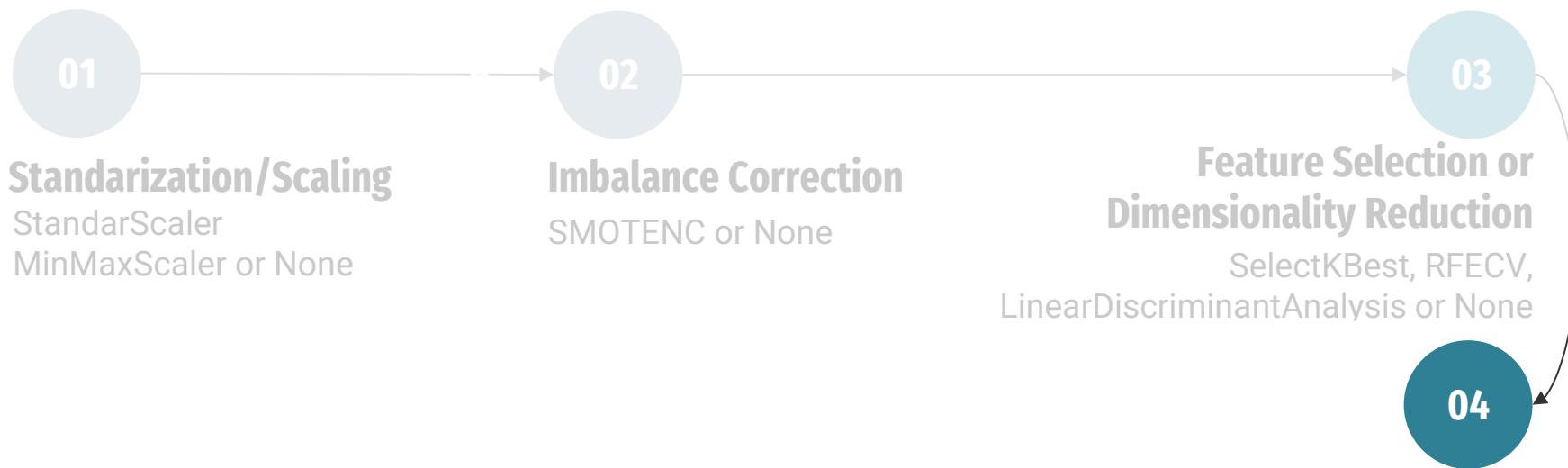


Imbalance Correction
SMOTENC or None

Methodology

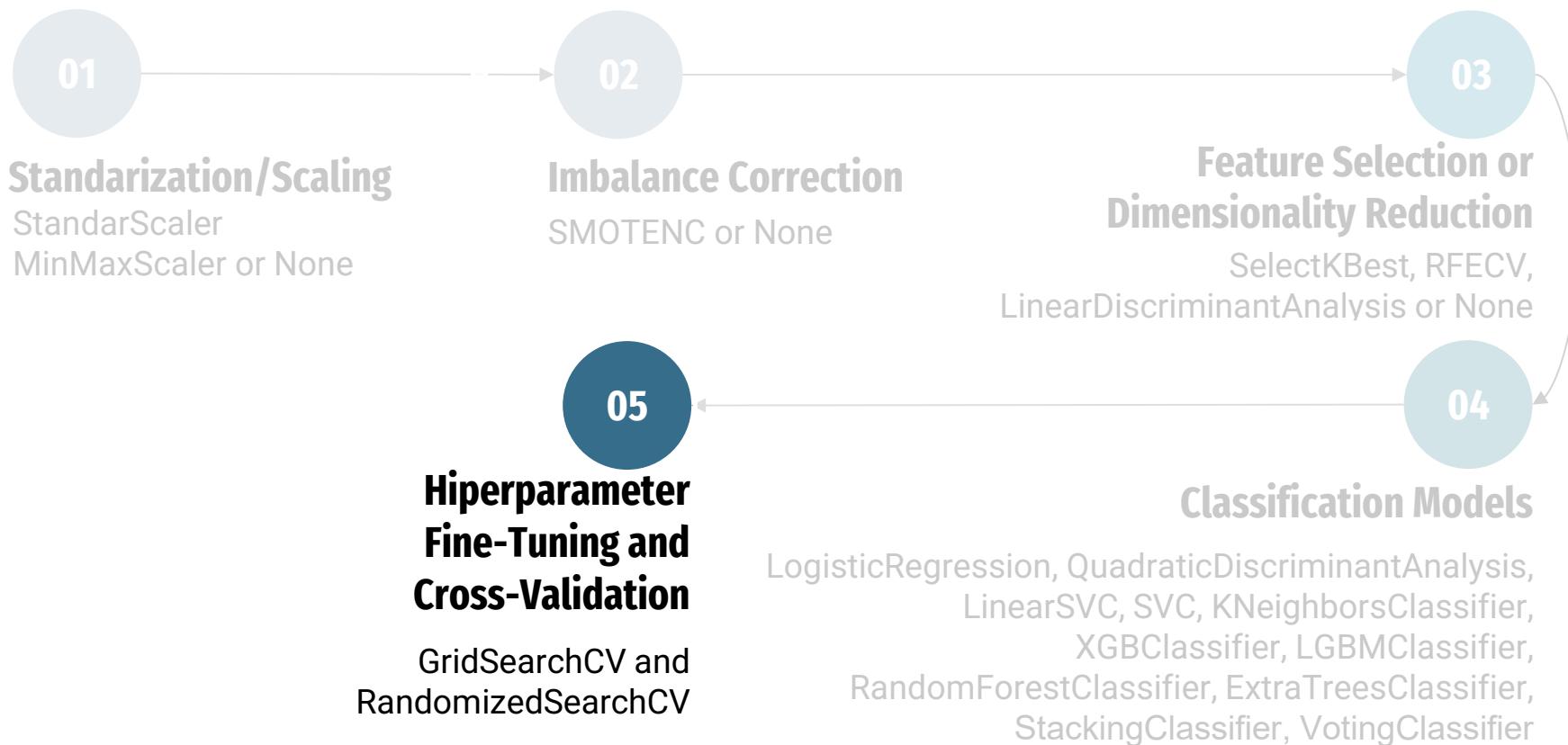


Methodology

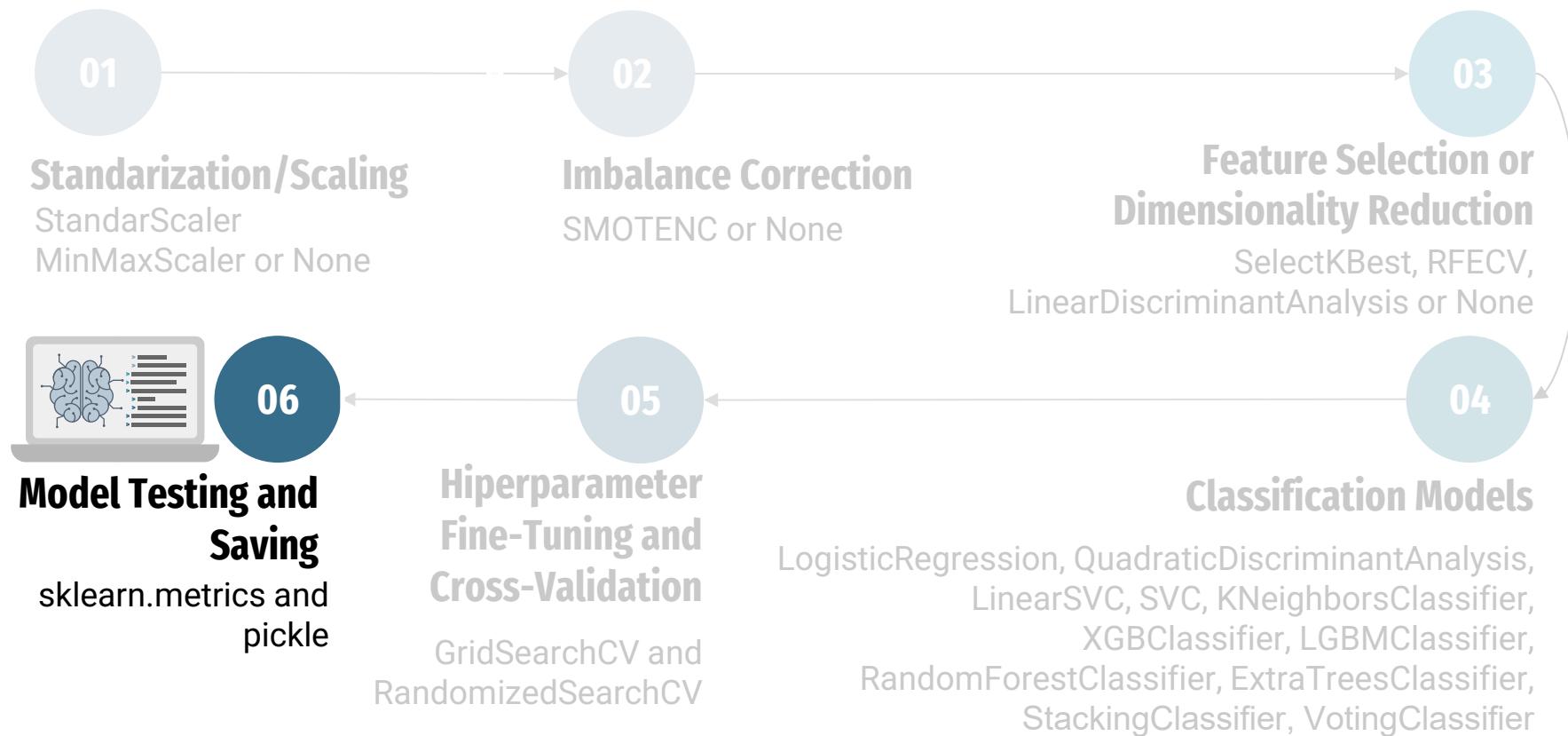


LogisticRegression, QuadraticDiscriminantAnalysis,
LinearSVC, SVC, KNeighborsClassifier,
XGBClassifier, LGBMClassifier,
RandomForestClassifier, ExtraTreesClassifier,
StackingClassifier, VotingClassifier

Methodology

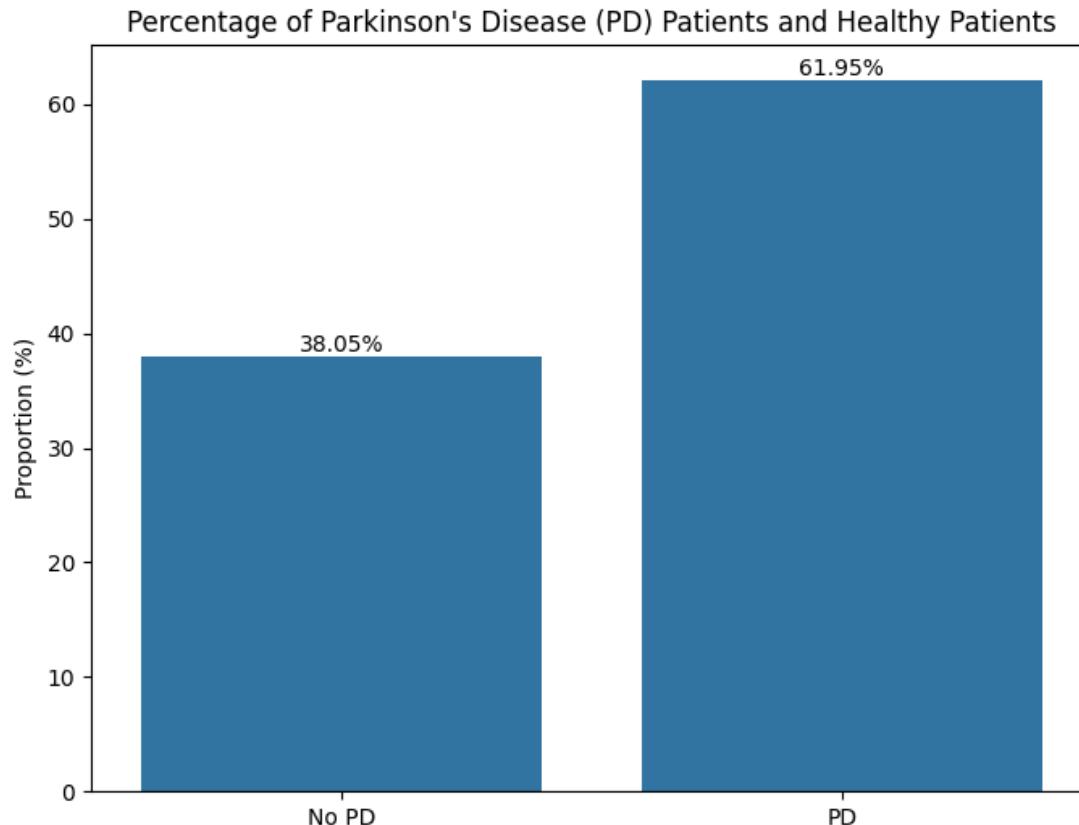


Methodology



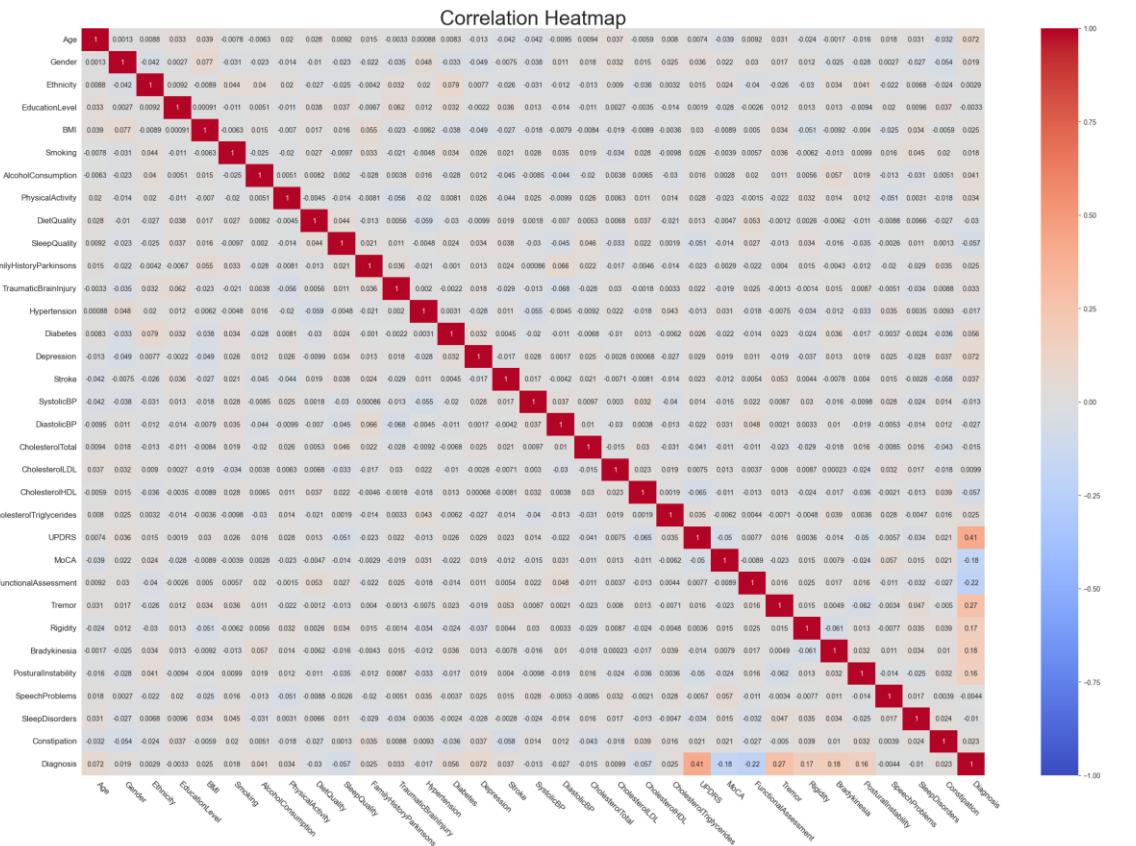
Exploratory Data Analysis

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Exploratory Data Analysis

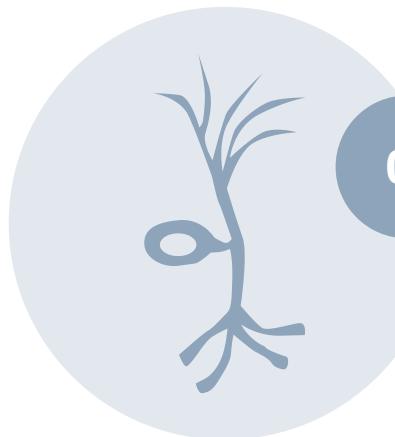
| | | Diagnosis |
|--|--------------------------|-----------|
| | UPDRS | 0.411858 |
| | Tremor | 0.271641 |
| | Bradykinesia | 0.183083 |
| | Rigidity | 0.167933 |
| | PosturalInstability | 0.159615 |
| | Depression | 0.072315 |
| | Age | 0.072114 |
| | Diabetes | 0.056443 |
| | AlcoholConsumption | 0.041170 |
| | Stroke | 0.036873 |
| | PhysicalActivity | 0.034081 |
| | TraumaticBrainInjury | 0.033186 |
| | CholesterolTriglycerides | 0.025269 |
| | FamilyHistoryParkinsons | 0.024888 |
| | BMI | 0.024799 |
| | Gender | 0.019451 |
| | Smoking | 0.017677 |
| | CholesterolLDL | 0.009858 |
| | Ethnicity | 0.002865 |
| | EducationLevel | -0.003334 |
| | SpeechProblems | -0.004429 |
| | SleepDisorders | -0.010265 |
| | SystolicBP | -0.013242 |
| | CholesterolTotal | -0.015405 |
| | Hypertension | -0.017471 |
| | DiastolicBP | -0.026635 |
| | DietQuality | -0.029933 |
| | SleepQuality | -0.057237 |
| | CholesterolHDL | -0.057320 |
| | MoCA | -0.179093 |
| | FunctionalAssessment | -0.217524 |

RESULTS: Final Models

In this project **two reliable predictive models with subtle differences** were developed

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01

MODEL 1

Maximizes the
accuracy score

RESULTS: Final Models

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01

MODEL 1

Maximizes the
accuracy score



02

MODEL 2

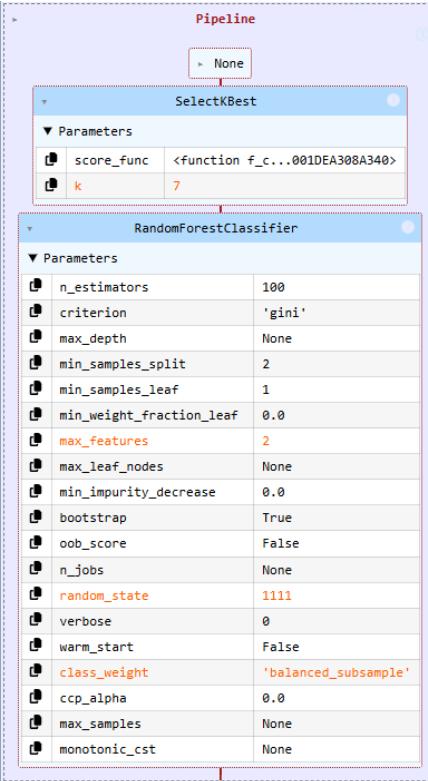
Maximizes the sensitivity
score

01

MODEL 1: Pipeline and Hyperparameters

01

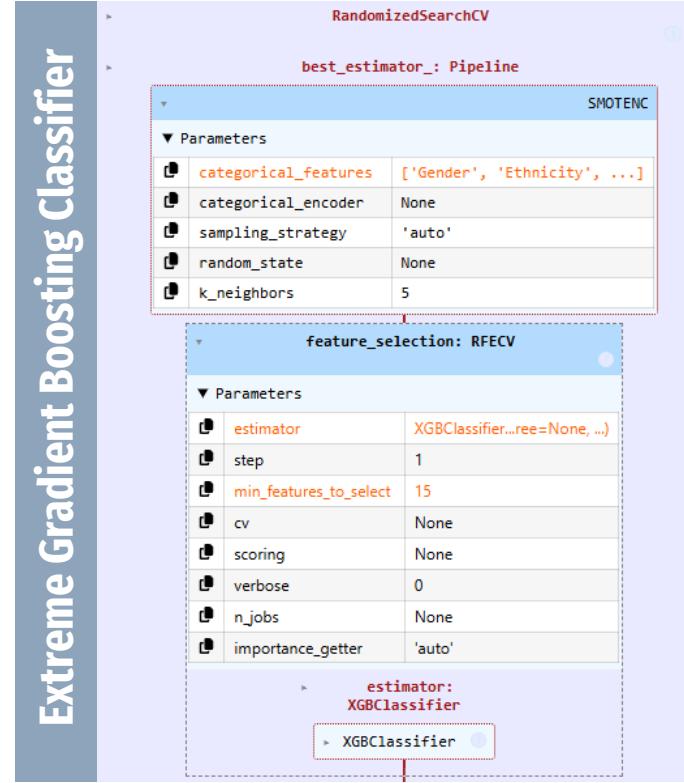
MODEL 1: Pipeline and Hyperparameters



Random Forest Classifier



Extreme Gradient Boosting Classifier



Random Forest Classifier



Extreme Gradient Boosting Classifier



01

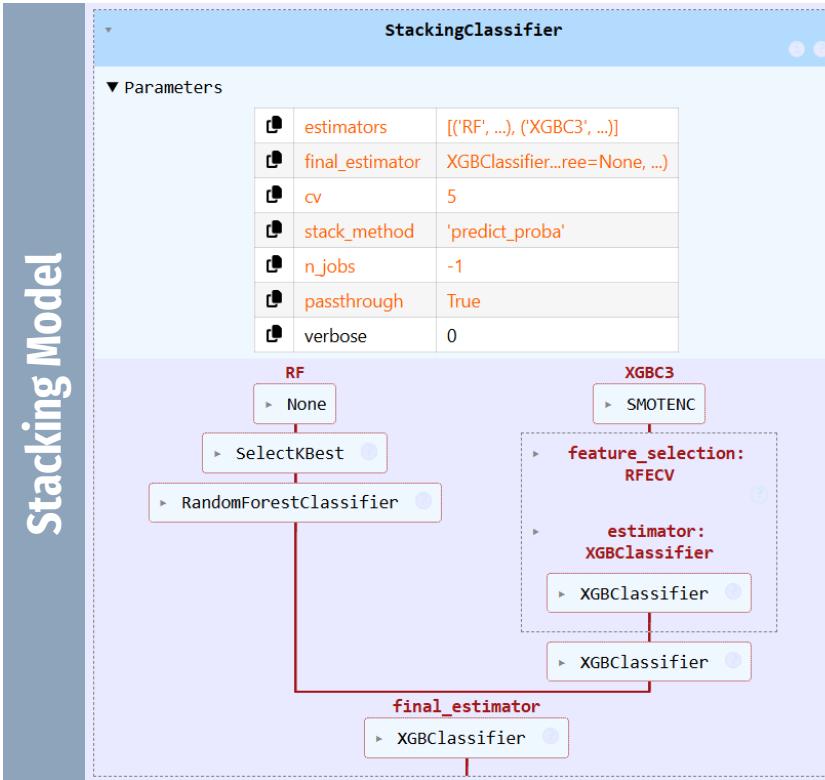
MODEL 1: Pipeline and Hyperparameters

01

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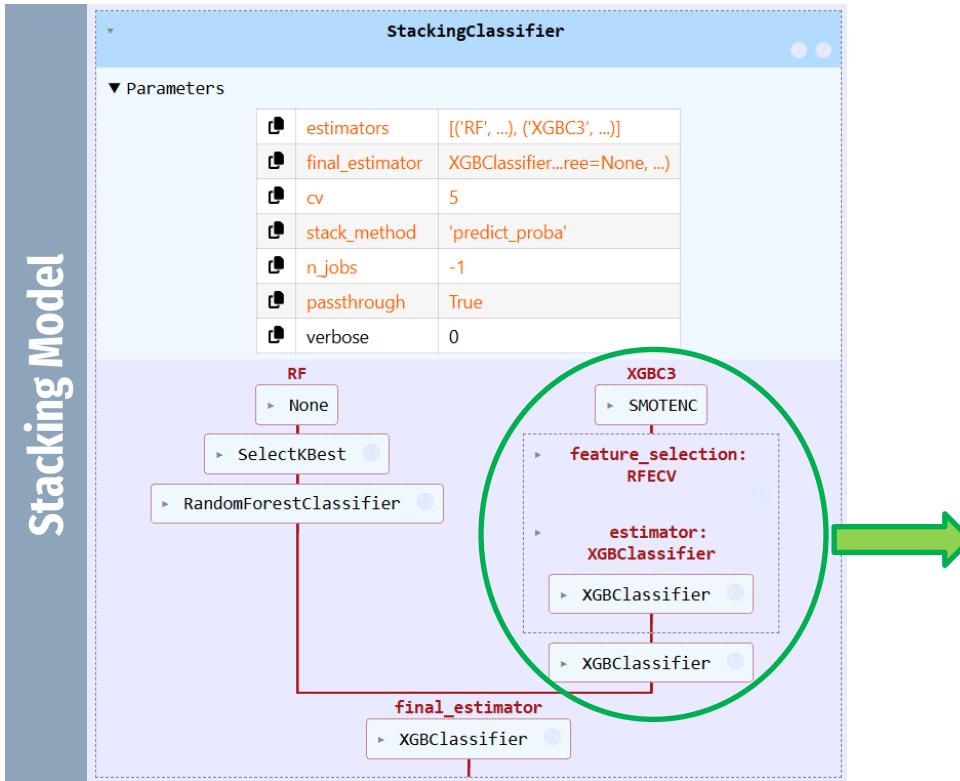
01

MODEL 1: Pipeline and Hyperparameters



01

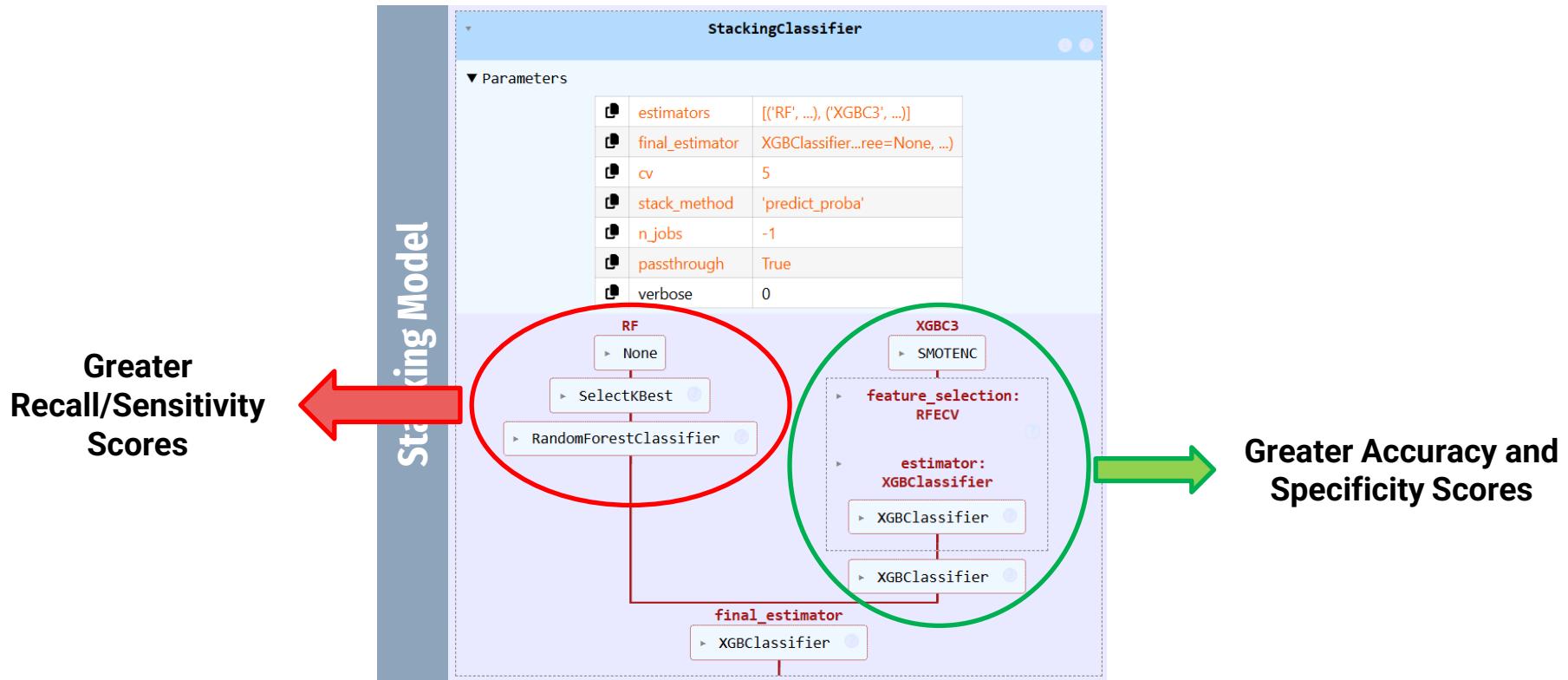
MODEL 1: Pipeline and Hyperparameters



Greater Accuracy and Specificity Scores

01

MODEL 1: Pipeline and Hyperparameters

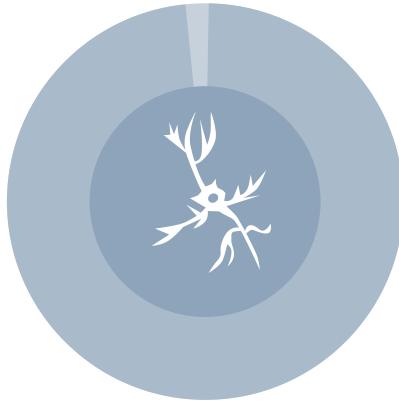


01

MODEL 1: Test Results

01

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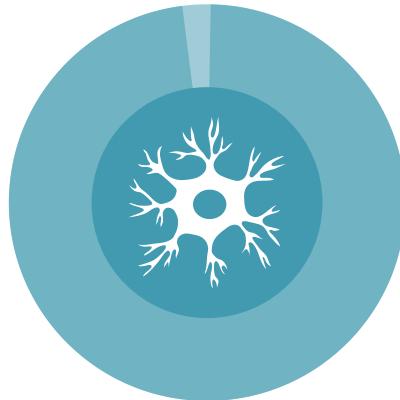
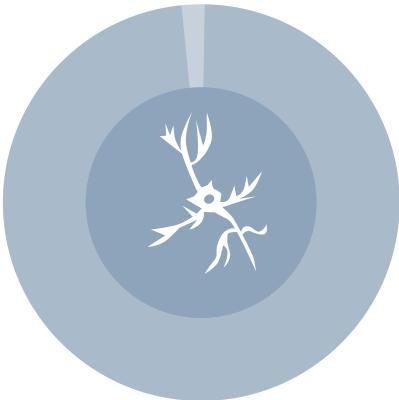


96.9% Accuracy

This model predicted correctly 408 of 421 patients' diagnosis

01

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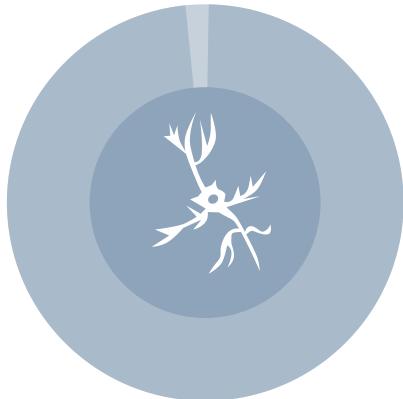
This model predicted correctly 408 of 421 patients' diagnosis

95.7% Specificity

155 of 162 of the healthy patients were diagnosed as healthy.

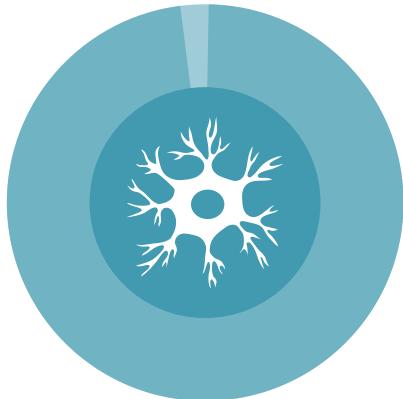
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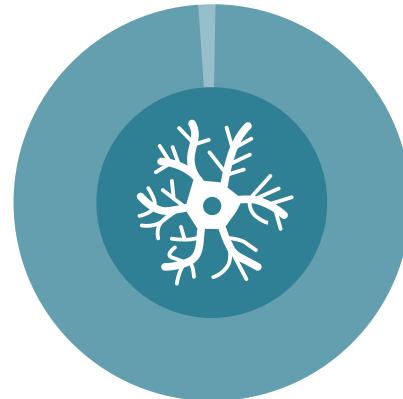
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97.7% Sensitivity

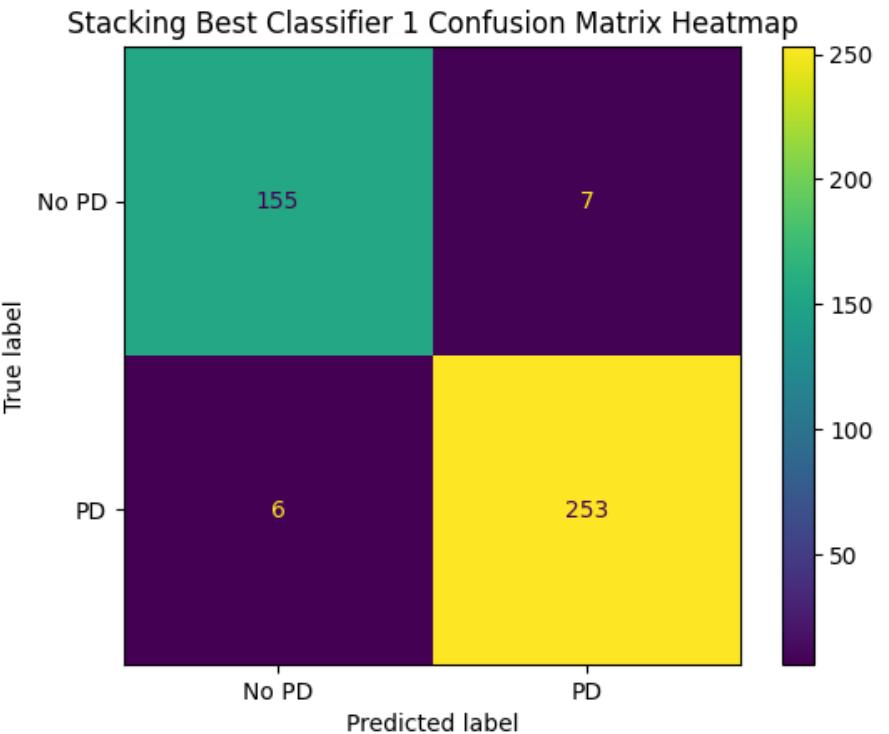
253 of 259 Parkinson's Disease patients were detected

01

MODEL 1: Test Results

01

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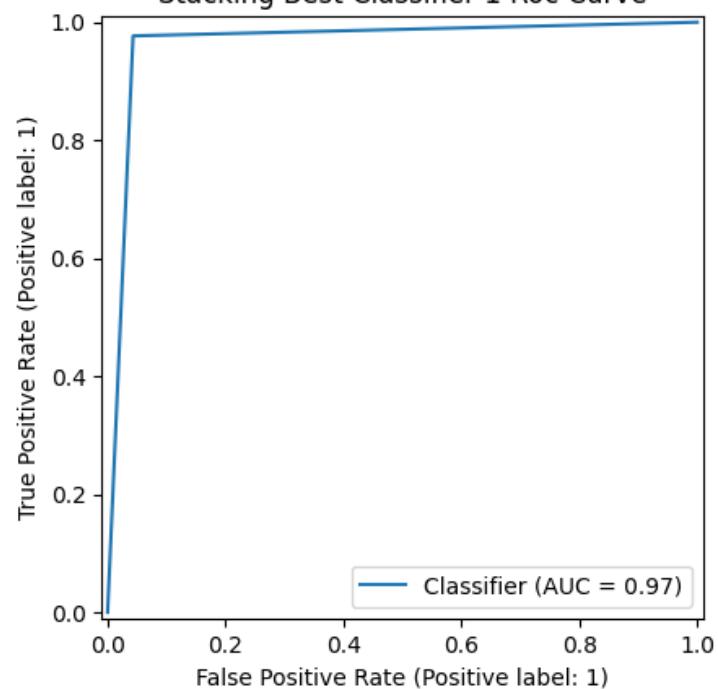
01

MODEL 1: Test Results

Stacking Best Classifier 1 Confusion Matrix Heatmap



Stacking Best Classifier 1 Roc Curve

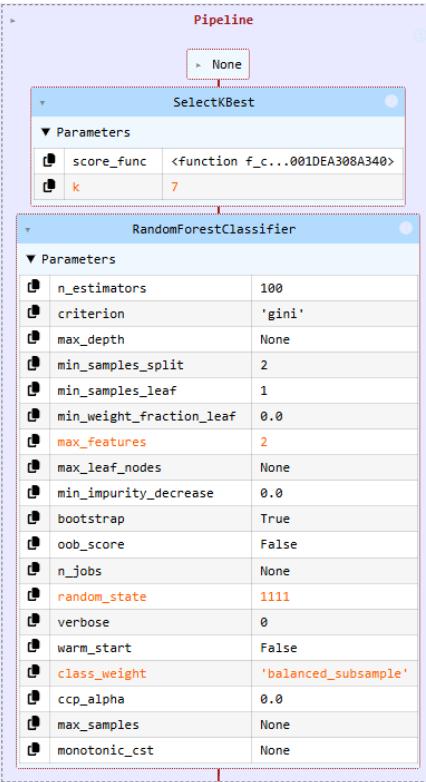


02

MODEL 2: Pipeline and Hyperparameters

02

MODEL 2: Pipeline and Hyperparameters



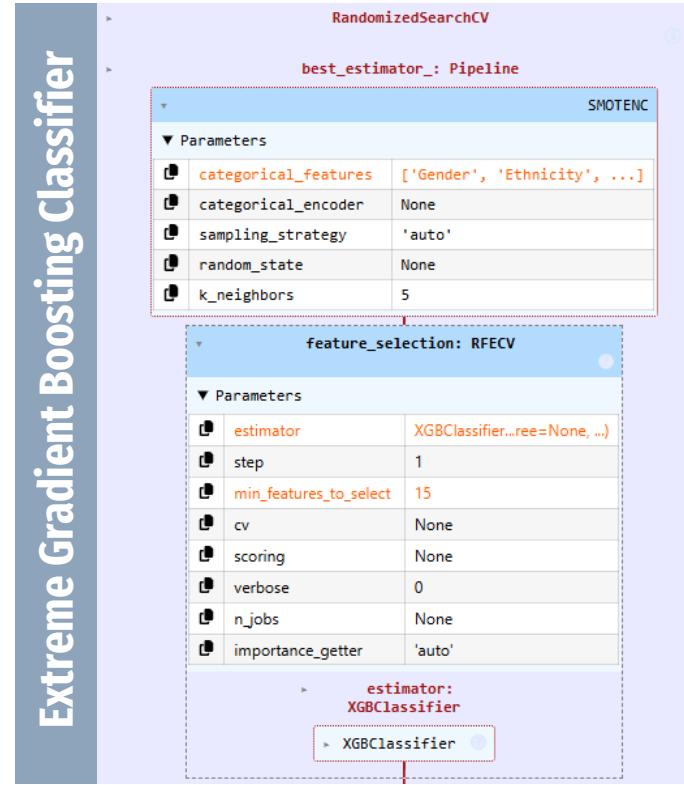
Random Forest Classifier



02

MODEL 2: Pipeline and Hyperparameters

Extreme Gradient Boosting Classifier



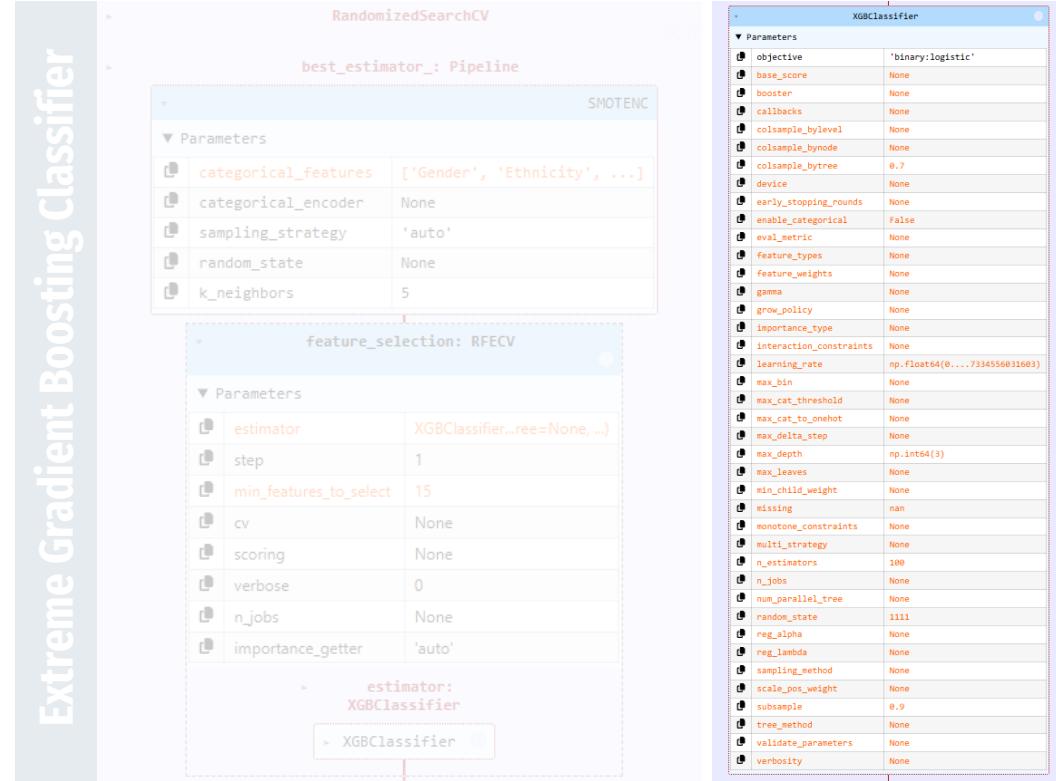
02

MODEL 2: Pipeline and Hyperparameters

Random Forest Classifier

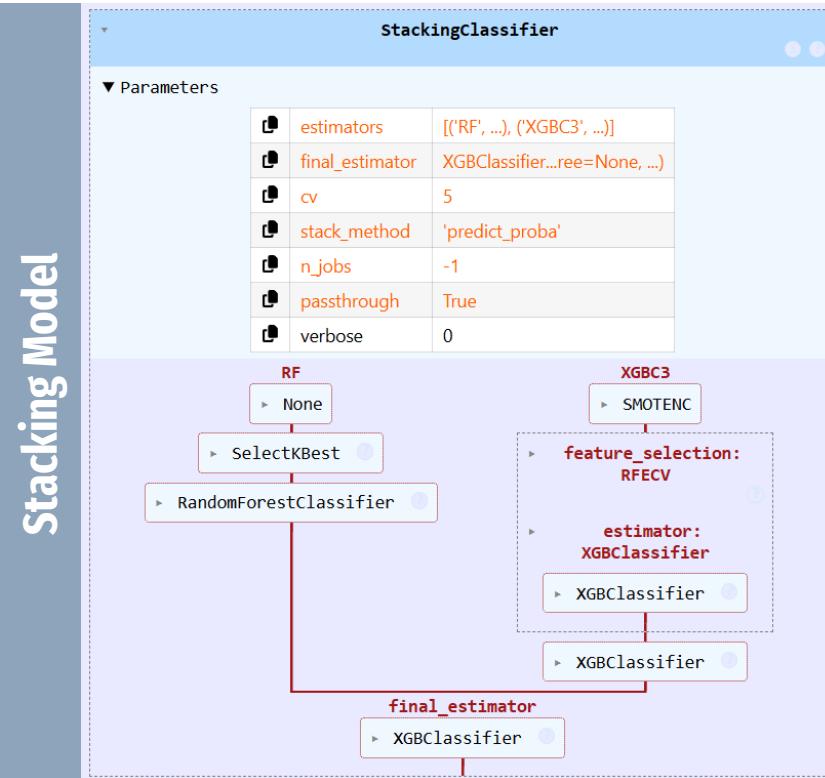


Extreme Gradient Boosting Classifier



02

MODEL 2: Pipeline and Hyperparameters

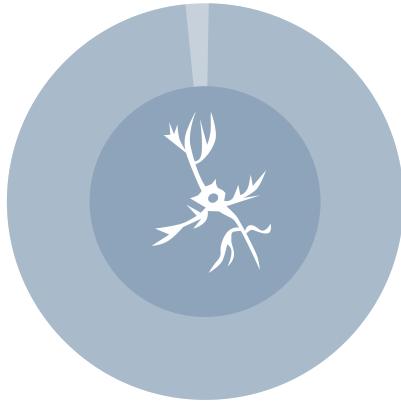


02

MODEL 2: Test Results

02

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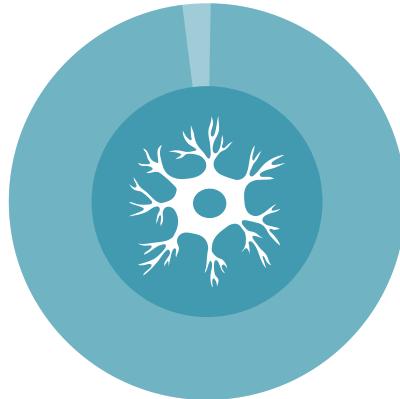
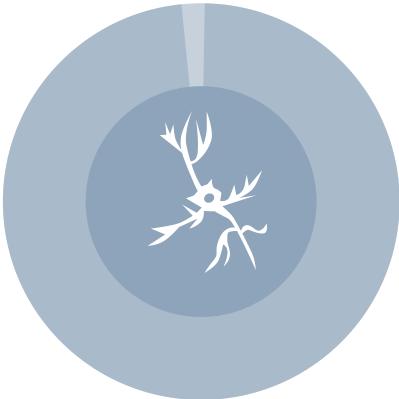


96.4% Accuracy

This model predicted correctly 406 of 421 patients' diagnosis

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MODEL 2: Test Results



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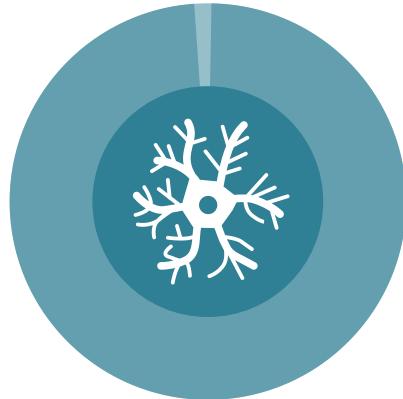
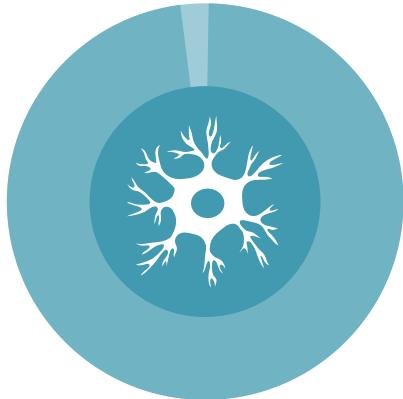
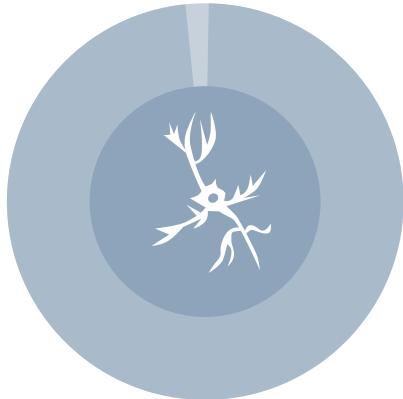
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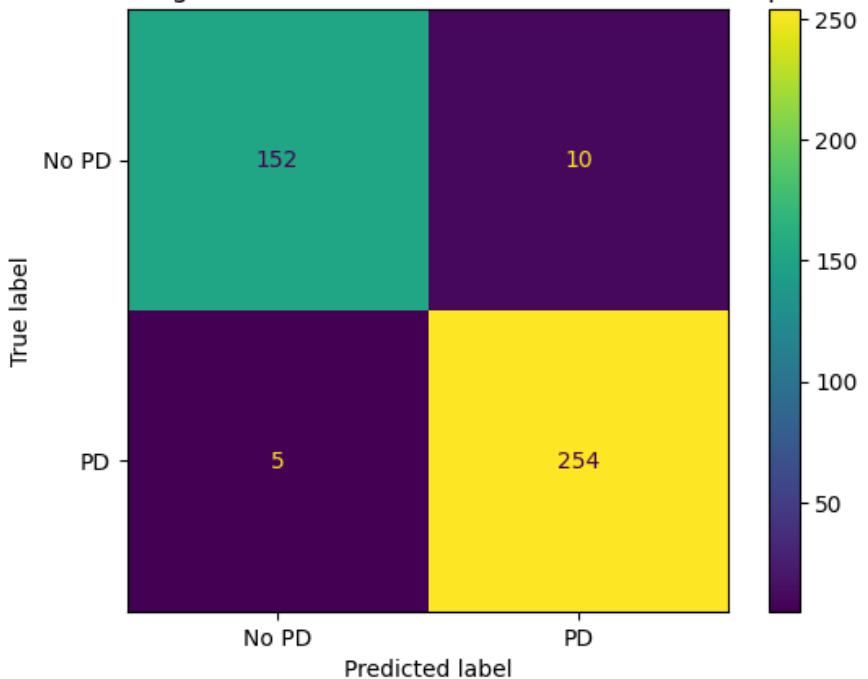
98.1% Sensitivity

254 of 259 Parkinson's Disease patients were detected

02

MODEL 2: Test Results

Stacking Best Classifier 2 Confusion Matrix Heatmap



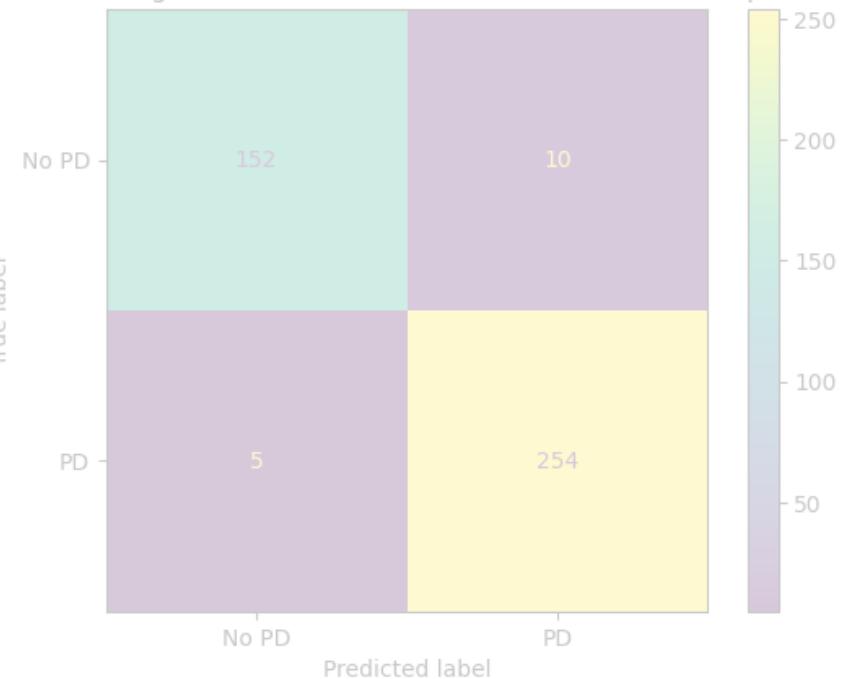
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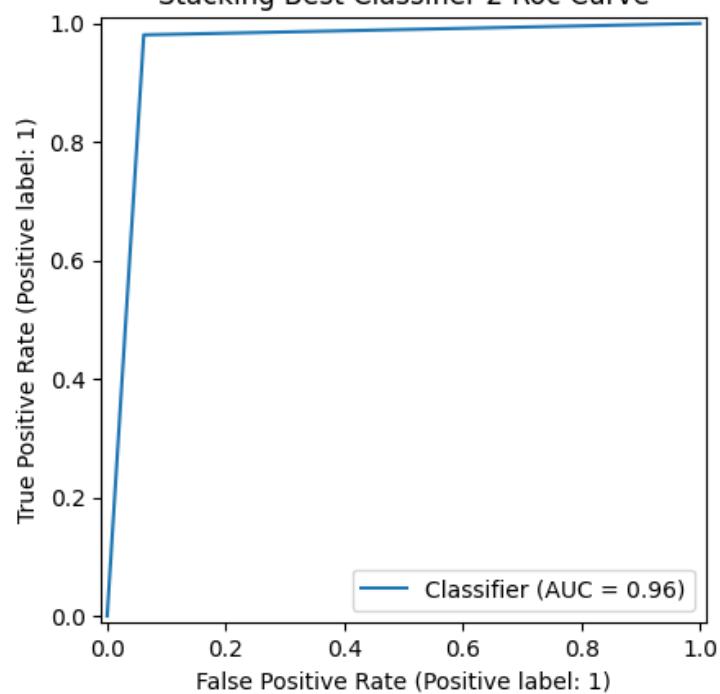
02

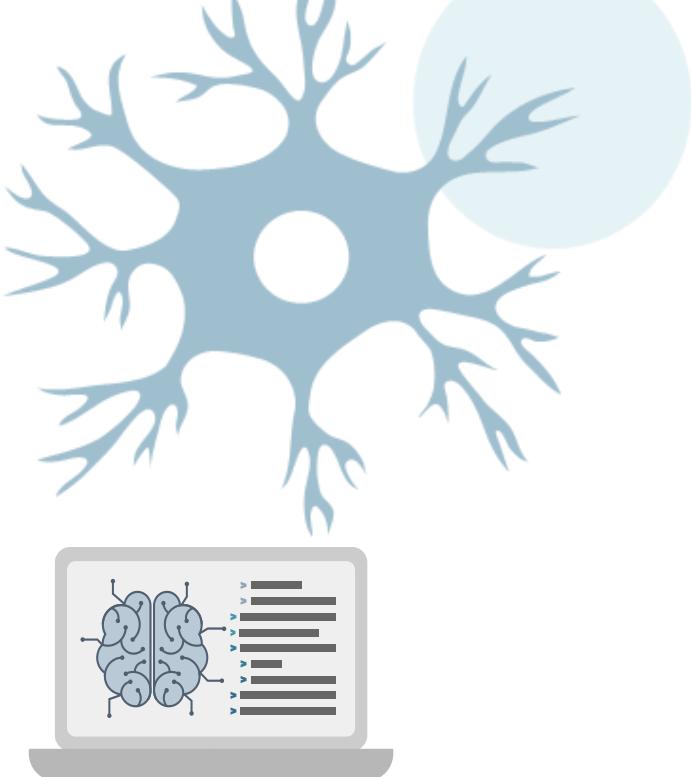
MODEL 2: Test Results

Stacking Best Classifier 2 Confusion Matrix Heatmap

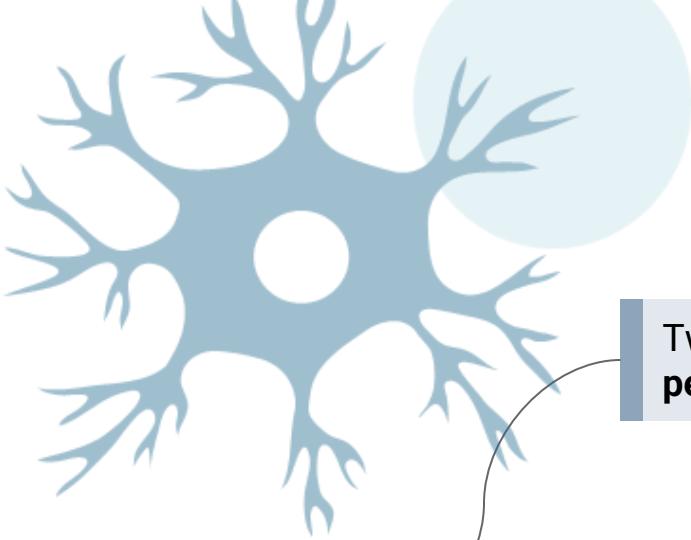


Stacking Best Classifier 2 Roc Curve





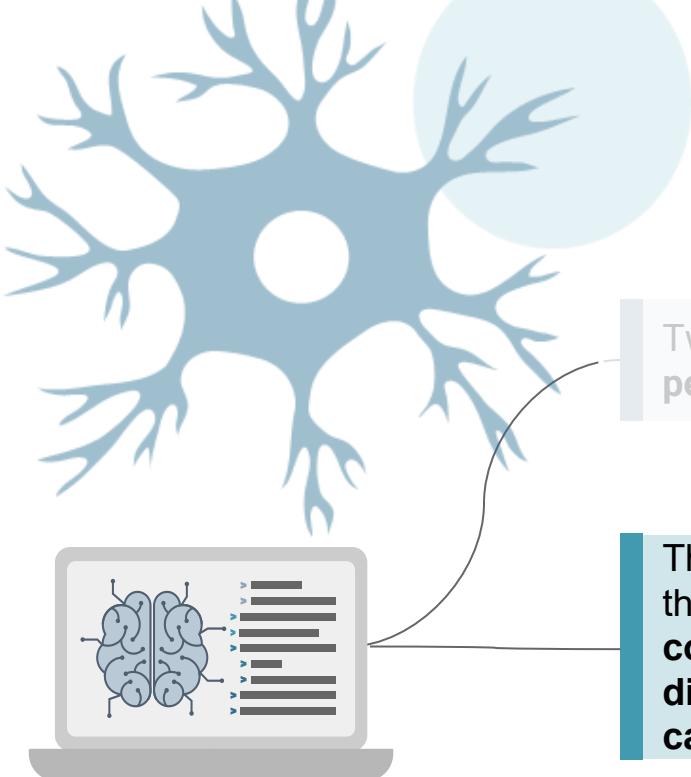
Conclusions



Conclusions

Two machine learning predictive models with impressive performance were developed successfully.

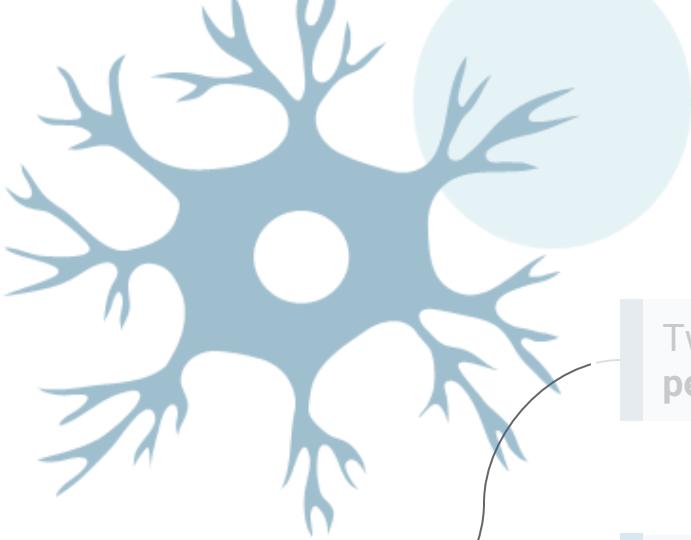




Conclusions

Two machine learning predictive models with impresive performance were developed sucessfully.

These models can support clinicians in **recognizing subtle signs** that might otherwise go unnoticed, **gain deeper insight** into the **complex interactions** that contribute to neurodegenerative disorders and reach **more personalized and proactive patient care**.



Conclusions

Two machine learning predictive models with impresive performance were developed sucessfully.



These models can support clinicians in recognizing subtle signs that might otherwise go unnoticed, gain deeper insight into the complex interactions that contribute to neurodegenerative disorders and reach more personalized and proactive patient care.

Timely diagnosis will improve the **effectiveness of available therapeutic interventions, influencing patient outcomes and long-term quality of life.**

Strengths and Limitations

Strengths

- Predictive models obtained impressive sensitivity scores (~98%).
- Components of the model are scalable and fast-performing, allowing fast prediction of huge datasets in seconds.
- Models have great flexibility. Consequently it can be adjusted and re-trained with new data with training times of less than one minute in train dataset of 2000 registers.

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Limitations

- Although the models obtained impressive sensitivity scores (~98%), specificity was compromised (93-96%), leading to some healthy patients incorrectly diagnosed as Parkinson's Disease patients.
- The model has not been tested in datasets with missing values.



**THANKS FOR YOUR
ATTENTION**



[Github Repository](#)



[Streamlit Webpage](#)