ORIGINAL ARTICLE

Artifcial Intelligence Model for Parkinson Disease Detection Using Machine Learning Algorithms

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Received: 12 January 2023 / Accepted: 21 February 2023 / Published online: 14 March 2023 © The Author(s), under exclusive licence to Springer Science+Business Media, LLC 2023

Abstract

In order for Parkinson's disease (PD) treatment and examination to be logical, a key requirement is that estimates of disease stage and severity are quantitative, reliable, and repeatable. The PD research in the past 50 years has been overwhelmed by the subjective emotional evaluation of human's understanding of disease characteristics during clinical visits. The Parkinson's disease data set contains 23 features and 197 instances, of which 8 patients are sound and 23 patients, are analyzed as PD patients. Relying on chi² test, extra trees classifier, and correlation matrix as feature extraction strategies and relying on Decision Trees, K-Nearest Neighbors, Random Forests, Bagging, AdaBoosting, and Gradient Boosting as supervised AI calculations for permutation calculations. The calculation is based to obtain higher classifer accuracy, as well as ROC curves accuracy. Three conspicuous component selection strategies allow each of the 23 features to select 10 best performing features. The DT classifer has a higher accuracy of 94.87% in a dataset with 23 attributions, just like a dataset with 11 features. These results are also checked by ROC curve $(AUC=98.7\%)$. This calculation significantly separates PD patients from patients at the individual level, thus ensuring the use of computer-based fndings in clinical practice.

Keywords Parkinson's disease · Machine learning · Classifers · ROC curve · Decision Tree

Introduction

Parkinson's disease (PD) is a dangerous disease that occurs on the earth after Alzheimer's disease. Countless people around the world have experienced this disease. PD is a reformist and long-term focus sensory system degenerative disorder that seriously affects the elderly [[1](#page-11-0)]. The significant side efects of PD are developmental weakness, such as delayed back development, muscle extension, hindered standing and balance, loss of procedural development, changes in speech, and changes in composition. The undergoing PD has no steady progress in dopamine under the body framework. The sound problem is a potential side efect for PD patients [[2\]](#page-11-1). Such patients have problems with speaking, such as volume level and irregular pronunciation. The sound problems of these issues can be evaluated for early PD analysis. Diagnosing and monitoring PD through speech signals is more accurate and equally powerful. The

 \boxtimes Saurabh Pal drsaurabhpal@yahoo.co.in result information is often used by neurologists to analyze PD through voice recording systems to help patients and get clear opinions. The new symptom model of Parkinson's infection has been released, and the main model rules for Parkinson's disease of the Movement Disorder Society have been established. Their goal is to help standardize clinical examinations [[3\]](#page-11-2). The confrmation of Parkinson's infection is usually guided by certain methods, such as observational evaluation and evaluation of patient clinical records. These strategies, like the abrupt strategies that distinguish PD, are not reliable in terms of accuracy and feasibility. The Medical Foundation announced that the current determination framework has not yet accurately distinguished Parkinson's disease. To overcome these limitations, we need a reliable technique that can be used to identify and help prevent PD. In this association, part of the AI strategy is critical to the identifcation, avoidance, and treatment of PD [\[4](#page-11-3)].

In order to overcome the aforementioned problems, this article proposes a new coordination strategy based on chi^2 , extra trees classifer and correlation matrix to select the outline of the appropriate features [\[5](#page-11-4)]. These calculations have been used to process a large number of features and rank them as needed. Compared with a separate demonstration,

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the mixture of the three calculations provides excellent execution [\[6](#page-11-5)]. Then, at that time, we used the selected features to train and test six classifers to predict PD patients. This article describes as follows:

- First, three calculation methods are proposed for selecting suitable features, that is, the chi^2 , extra trees classifer, and Correlation matrix, which allocates an appropriate load to each component in the feature set, locates the feature according to the weight, and fnally resolves the correlation.
- Secondly, the presentation of decision tree, K-nearest neighbors, random forest, Bagging, AdaBoosting, and Gradient Boosting has been evaluated using selected features. The results show that compared with the frst feature list, DT has created key results based on the features of the chi² , extra trees classifer, and correlation matrix. In addition, ROC curve has been drawn for each selected object, including all the features approved by the result obtained by the classifer.
- We have conducted extensive investigations on realworld data sets, and the results show that compared with partners, the proposed analysis techniques (including selective classifer-AI classifer) have achieved key results in terms of high accuracy and low computational cost.

The rest of the paper is divided into seven parts. Section ["Literature Review](#page-1-0)" describes the writing survey. Section "[Materials and Methods"](#page-2-0) introduces the tools and techniques for each model/device/program/calculation used in the discussion, and their importance in advancing the proposed method. The Sect. "[Experimental Setup"](#page-4-0) experiment is arranged to simulation environment and the required boundary and data set depiction. Section "[Results](#page-4-1) [and Discussion](#page-4-1)" discusses the legitimacy and adequacy of the model through decomposition of the results, and fnally Sect. ["Conclusion](#page-9-0)" describes the conclusion of work and the scope of future inspections.

Literature Review

In the writing, the suggested PD analysis strategies, obstacles and benefts are summarized in Table [1](#page-1-1) for better arrangement, just as the importance of the strategies we proposed. Nonetheless, these technologies are limited in selecting the outline of the appropriate features, and therefore sufer from a lack of PD recognition and profciency issues.

Das and Tsanas et al. [\[17,](#page-11-6) [18\]](#page-11-7), proposed unique artificial intelligence-based techniques that have been created to analyze PD patients. Little et al. [\[7\]](#page-11-8) proposed a technique for distinguishing Parkinson's disease using speech signal information. They distinguished between 23 PD patients and 8 able-bodied subjects. SVM is used to characterize Parkinson's disease and healthy individuals. The accuracy of the proposed strategy was recorded at 91.4%. In another survey [\[18](#page-11-7)], 132 features were selected based on the signs of dysphonic discourse. Calculations using specular selection (FS) like LASSO, Relief, MRMR and LLBFS [\[18](#page-11-7)]. In addition, the model uses the feature selection calculation to select 10 features from 132 features, which are used for the sequence of Parkinson's disease and the entity. In contrast, Sakar et al. [[8\]](#page-11-9) reported that a large number of voice recordings from 40 subjects were collected, of which 20 were Parkinson's disease subjects and 20 were non-Parkinson's subjects. 26 speech signals containing daily pronunciation, words, numbers and vowels were recorded. They used the Praat acoustic inspection program to record the speech [\[19](#page-11-10)]. In addition, a theme (LOSO) and S-LOO approval strategy were used to check the presentation of K-NN and SVM classifers [[20](#page-11-11)]. Exploratory work [[7\]](#page-11-8) proposed a strategy that relies on ML calculations, using speech signals to diagnose Parkinson's disease, conveys the calculations of feature

selection, such as help, LLBS, LASSO, and mRMR, and the proposed method achieves excellent results in terms of accuracy. Sakar et al. [\[8](#page-11-9)] established an analysis framework using SVM and achieved an accuracy of 92.75%. In addition, Der et al. [[9\]](#page-11-12) by using a fufy-based indirect change strategy combined with SVM, proposed a model for analyzing PD, and an accuracy of 93.47% was achieved. Andre et al. [[10\]](#page-11-13) proposed a symptom framework for PD recognition using the backwoods classifer based on the strategy of change and the ideal way. The framework achieved an accuracy of 84.01%. Chai et al. [[14](#page-11-17)] planned another academic project to identify PD. The SVM and mitigation calculations are coordinated with the simplifed calculations for bacterial removal, and critical accuracy is achieved. Emarie et al. [[19\]](#page-11-10) cultivated a program that uses fufy theory, K-NN, and PCA to diagnose PD, and achieved an accuracy of 96.07%. Tsanas [\[18](#page-11-7)] planned the use of PSO and improved FKNN to find the determination strategy of PD, and obtained an accuracy of 97.47%. To this end, Gok [[11](#page-11-14)] used the results of the Rotation Forest Ensemble (RFE) KNN classifer to propose a PD analysis framework with an accuracy of 98.46%. Along this path, Das [\[15\]](#page-11-18) studied the scheduling and execution of ANN, strategy recurrence (LR) and (decision tree) DT. Compared with other LRs and DTs, the grouping execution of ANN is very good in terms of accuracy, and an accuracy rate of 92.9% is obtained. A PD discovery framework was proposed in [[12](#page-11-15)], using mRMR to feature certain calculations and complex and respected ANN classifers. The proposed framework has achieved 98.12% accuracy. Taking into account the writing of the survey, we infer that to efectively derive PD, a very smart judgment framework is required. In the planning of the PD analysis framework, current investigations [\[21,](#page-11-20) [22](#page-11-21)] have used unique grouping calculations, for

example, strategy recurrence [[23\]](#page-12-0), support vector machines [[13](#page-11-16)], k-NN [\[23\]](#page-12-0), DT, NB [[24](#page-12-1)] and ANN discovered PD. Among these classifers, compared with diferent classifers, the help vector machine performs very well. In view of the occasional extra features that will afect the performance of the characterization, just like the computational complexity of the model, the grouping execution of the classifer can be improved by selecting the appropriate element determination technology. Notable element selection and boundary improvement calculations include: help, mRMR, LASSO, LLBFS, Genetic-Algorithm (GA), Particle-Swarm Optimization, Whale- Optimization-Algorithm (WO), Natural Product Fly Enhancement (FFO), Diferential Flowering Fertilization and Bacterial Elimination and Refinement (BFO) have been used in the feature selection of the existing trial selection outline features.

Materials and Methods

The specifc details and basic ideas used in the proposed model are described below in Fig. [1](#page-2-1). This section proposes six machine learning classifers and three feature selection methods.

Machine Learning Classifers

Decision Tree

There is a classifer whose graphical explicit feature determination is an essential part of the learning cycle: the selection table. The entire question of the study selection table includes selecting the correct credits to be combined.

Fig. 1 The fowchart describes each step/phase of the method

Usually, this is done by estimating the cross-approval execution of the various feature subsets of the form and selecting the best performing subset. Fortunately, the leave-one-out cross-approval is very gentle for this classifer. Obtaining cross-approval errors from the selection list obtained from the preparation information is just a matter of controlling the class check associated with each table entry, because the design of the table will not be changed or erased as occasions increase [[25,](#page-12-2) [26](#page-12-3)]. To a large extent, the feature space is searched through the pursuit of best priority, because this method is less likely to fall into the largest neighborhood than other methods, such as forward selection.

K‑Nearest Neighbor

In order to handle the diferent marks, a directional calculation is used, which is adjusted and a bunch of name signals are obtained. To assign another point, it fnds the nearest point and makes a decision on that point, so it assigns the nearest mark [[27,](#page-12-4) [28](#page-12-5)]. The following distance work is used to evaluate KNN.

$$
\sqrt{\sum_{i=1}^{k} (x_i - y_i)^2}
$$
 Euclidean function

$$
\sum_{i=1}^{k} |x_i - y_i|
$$
 Manhattan function

$$
\left(\sum_{i=1}^{k} (|x_i - y_i|)^q\right)^{1/q}
$$
 Minkowski function

Random Forest

Random forest classifer is a comprehensive learning system for collecting, backing off, and various efforts that can be performed with the help of decision trees. These decision trees can work during planning, and the benefts of this category can be portrayal or retrogressive. With the help of such unpredictable remote areas, people can resolve their affinity for over-adaptation to arrangement sets [[29\]](#page-12-6).

At the random forest level, it is completely expected on all trees. The importance of the entire part of each tree is evaluated and isolated by the complete number of trees:

$$
RFfii = \frac{\sum_{j \in all trees} norm}{T}fiji.
$$

where, RFf*ii* is the signifcance of highlight I determined from all trees in the Random Forest model. Norm f*iij* is the standardized element signifcance for I in tree *j*. T is the absolute number of trees.

Bagging

The idea of bagging (deciding grouping, average recurrence type problems, and uninterrupted ward income factors) is suitable for prescient information mining space, adding expected orders (predictions) from many models, or models from various learning information of similar types. It is also used to solve the inherent instability of the results, while applying complex models to the index of usually little information. Assuming that the task of information mining is to build a model with a foresighted arrangement, there are usually very few data sets to prepare the model. We can generate sub-examples (with substitutions) from the data set multiple times and apply, for example, tree classifers (such as CART and CHAID) to progressive examples. In fact, it is common to develop completely diferent trees for various examples, outlining the instability of the model that is usually obvious with a small number of data sets [[30](#page-12-7)]. One strategy for determining individual predictions (for novel perceptions) is to use all the trees found in various examples and apply some basic democracy: the last feature is a feature that various trees often predict.

AdaBoosting

AdaBoosting or Adaptive Boosting is an AI used for metacomputation. Diferent learning indicators are usually used to further improve execution efficiency. The benefits of other learning evaluations will be combined into a weighted whole, which is stable with the last beneft of the supported classifers. AdaBoosting is versatile and can guarantee substitute students who are powerless due to the misclassifcation of past classifers [[31](#page-12-8)]. AdaBoosting perceives large amounts of data and one condition. On some issues, overftting is not as defensive as other learning measures. Each substitute may be weak, but as long as everyone performs better than any theory, the last model may eventually be severely afected by a strong substitute.

$$
E_t = \sum_i E[F_{t-1}(x_i) + \alpha_t h(x_i)]
$$

Among them, $F_{t-1}(x)$ is the Boosted classifier, E (F) is the error function, $F_t(x) = a_t h(x) = \text{trail learner}, h(x_i)$ is the test in the learning set, t is the number of iteration, α_t is the distribution coefficient, E_t is the boost result of the classifier.

Gradient Boosting

Gradient boosting is an AI method for recurrence and characterization problems. It gives an expectation model as a bunch of general forecasting models and selection trees. Like other upgrade methods, it builds models in an ingenious way and summarizes them by allowing self-affirmation to be recognizable by appalling work [[32](#page-12-9)].

Extensive use of "gradient improvement" follows strategy 1 to limit target work. In each cycle, we adjust the basic students to the negative point of the negative tendency and continue to increase the normal value, and add it to the previously emphasized motivation.

$$
F_m(x) = F_{m-1}(x) - \gamma_m \sum_{i=1}^n \nabla F_{m-1} L(y_i, F_{m-1}(x_i)),
$$

$$
\gamma_m = \frac{\text{argmin}}{\gamma} \sum_{i=1}^n L(y_i, F_{m-1}(x_i)) - \gamma \nabla F_{m-1} L(y_i, F_{m-1}(x_i))
$$

where $L(y, F(x))$ is a differentiable loss function.

Feature Selection Method

Suppose we consider the list of capabilities to be processed as x with n features. The feature selection is picking m, out discrete advancement problem n contains the set, that is, $m \le n$ (24). Display and execute a classifier that is basically unafected by features. Therefore, it is fundamentally important to deal with unimportant features from the feature set [\[33,](#page-12-10) [34](#page-12-11)].

Chi2 Test

The χ 2 (chi2) test involves determining the calculation of χ 2 between each component and the target and selecting the ideal number of features with the best χ 2 score using the following equation $[35]$ $[35]$ $[35]$:

$$
\chi 2 = \sum_{i=1}^{n} \frac{(O_i - E_i)^2}{E_i}
$$

where O_i is the Observation in class *i*. E_i is the observations in class *i* if there was no relationship between the feature and target.

Extra Trees Classifer

For extracting salient features between data set elements by applying the element importance of the model, the model scores each information component and the higher the score, the more components in the income variable [[36](#page-12-13)]. We apply the ET classifier to evaluate the five main features of the data set.

Correlation Matrix

Correlation is an attribute to check whether the characteristics of the data set are associated with the target variable. The relationship may be positive or negative. To a certain extent, if the single meaning of a feature is expanded, it will increase the value of fairness, while if the single meaning of the relevance is expanded, it will reduce the objective value [[37](#page-12-14)]. Through the heat map, it can undoubtedly discover which features are most suitable for the target variable.

Experimental Setup

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The data set used in this examination can be accessed online on the UCI machine learning repository [[38\]](#page-12-15), which contains the acoustic features of 31 patients. 23 of these patients are experiencing PD. The data set has 197 instances, of which 23 are acoustic features that are separate from the patient. The exploratory meeting aims to discover the features that improve the expected PD performance (see Table [2\)](#page-5-0). The analysis was done on Jupyter Notebook (Anaconda3), Python adapted to 3.8 and 32-bit Windows 7 framework, 4 GB RAM, and Intel® Core™ i3-4600U CPU @ 2.10 GHz 2.70 GHz. The size of the preparation set and test set is 80% and 20%, respectively. In order to evaluate the performance of each classifer, the results have been taken into account for accuracy. Finally, we analyzed the results obtained from the experiment. Table [2](#page-5-0) describes the details of the PD patient data set.

Results and Discussion

Pre-preparation methods, for example, deleted missing feature, standard scalar, and min–max scalar have been applied to the data set to successfully prepare and test with the classifer. These factual strategies are the basis for a basic understanding of the data set. The data set has 197 instances and 22 real value features and an output object class. Figure [2](#page-6-0) is a correlation matrix, which is a two-dimensional depiction of information, where colors indicate values. The correlation matrix provides a quick visual summary of the data. More complex matrix allows

Attribute's name	Description	Status of sound recording count	
Name	Subject name and recording number	PD: 1-147 (23-people) Healthy: 0-48 (8-people)	
MDVP: Fo(Hz)	Average vocal fundamental frequency		Count of Status
MDVP:Fhi(Hz)	Maximum vocal fundamental frequency		
MDVP: Flo(Hz)	Minimum vocal fundamental frequency	140	
$MDVP: Jitter(\%),$ MDVP: Jitter(Abs), MDVP:RAP, MDVP:PPO, Jitter:DDP	Measures of variation in fundamental frequency	120	
MDVP:Shimmer, MDVP:Shimmer(dB), Shimmer:APQ3,	Measures of variation in amplitude.	100 80	
Shimmer:APQ5, MDVP:APO, Shimmer:DDA		Count 60	
NHR, HNR	Ratio of noise to tonal components in the voice		
Status	Health status 1-Parkinson's Disease (PD) 0-Healthy	40 20	
RPDE, D ₂	Two non-linear dynamical complexity measures	0	
DFA	Signal fractal scaling exponent	0	
Spread1, Spread2, PPE	Nonlinear measures of fundamental frequency- variation		Status

Table 2 Description of PD patient's dataset

observers to understand complex data sets. In addition, the links between factors indicate that when the value of one variable changes, the other variable usually moves in a certain direction. Understanding this relationship is helpful because we can use the value of one variable to predict the value of another variable.

Result Based on Feature Selection

In this section, the test results of feature selection calculation chi2, extra tree classifers, and correlation matrix have been explained and discussed in detail. No element is selected in any feature selection algorithm: MDVP:Jitter (%), MDVP:RAP, MDVP:PPQ, and Jitter:DDP and RPDE. Subsequently, these characteristics have little effect on the confrmation of PD.

The features selected by χ chi² calculation are shown in Table [3](#page-7-0).

The features selected by the extra tree classifer are shown in Fig. [3.](#page-7-1)

² Springer

In addition, the characteristics of the correlation matrix selection are represented in Table [4](#page-8-0) and Fig. [4,](#page-8-1) respectively.

After applying the three feature selection techniques, 10 best features and one output class are selected from each method mentioned in Tables [3](#page-7-0) and [4](#page-8-0) and Figs. [3](#page-7-1) and [4](#page-8-1).

Root Mean Square Error (RMSE) play an important role in the performance of classifers. It is defned as the values predicted by a classifer and the values actually observed. The values of RSME for training and testing datasets are similar if we have developed the good classifer; in other case if the RMSE values are much higher in testing of data than training data the classifer developed is not good. The RMSE values is calculated using the formula

$$
RMSE = \sqrt{\frac{1}{n} \sum_{i=1}^{n} (y_i - \hat{y}_i)^2}
$$

Chi² , extra tree classifers, and correlation matrix have been used for efective training and testing of the classifer DT, KNN, RF, Bagging, AdaBoosting, and Gradient

Table 3 Feature selected by chi² algorithm

Attributes	Score	
MDVP:Flo(Hz)	456.626628	
MDVP: Fo(Hz)	316.985398	
MDVP:Fhi(Hz)	227.402656	
HNR	22.691579	
MDVP:Shimmer(dB)	3.210348	
PPE.	2.151107	
D ₂	1.381600	
Spread ₂	1.232614	
Shimmer:DDA	0.462793	
NHR	0.457699	

Boosting. Thus the experimental results of feature selected by chi2, extra tree classifers, and correlation matrix with classifers are reported in Table [5.](#page-9-1) The experimental results show that the classifer classifcation performances is same for classifer DT which is 94.87% on reduced feature sets as well as the full feature set. While on the other hand the classifiers (KNN % accuracy $= 82.05$ for without feature Selection & chi² feature selection & Extra trees classifier, RF accuracy = 92.30% for without feature Selection & chi² feature selection, Bagging accuracy= 92.30% for without feature Selection, AdaBoosting accuracy=89.74% for

Correlation matrix, Gradient Boosting accuracy=94.87% for chi² feature selection & Correlation matrix) have the higher performance accuracy.

Based on these statistical results, we conclude that DT is signifcantly better than other peers in accuracy, so the proposed method is suitable for PD identifcation. Therefore, using chi2, extra tree classifiers, and correlation matrix FS algorithms and classifers (DT, KNN, RF, Bagging, AdaBoosting, and Gradient Boosting) to select more appropriate features helps the model efectively diagnose PD. The features selected by the proposed FS algorithm include MDVP:Fo (Hz), MDVP:Fhi (Hz), MDVP:Flo (Hz), MDVP:Jitter (Abs), MDVP:Shimmer, MDVP:Shimmer (dB), Shimmer:APQ3, Shimmer:APQ5, MDVP:APQ, Shimmer:DDA, NHR, HNR, DFA, spread2, D2, and PPE. In short, the proposed method can be used to detect PD, especially in the early detection of PD. Figure [5](#page-9-2) represents the performance of classifiers with and without feature selection.

Result Based on ROC Curve

ROC curve were assessed to each set and subset of PD patients to recognize affectability (true positive rate) against the investigation group. Region under the ROC curve (AUC) was assessed to gauge how well the

Fig. 3 Feature selected by Extra Trees Classifer

Table 4 Features selected by correlation matrix

Feature name	Score
PPE	0.53
spread2	0.45
MDVP:Shimmer	0.37
MDVP:APQ	0.36
Shimmer:APO5	0.35
Shimmer:APO3	0.35
MDVP:Shimmer(dB)	0.35
Shimmer:DDA	0.35
MDVP: Fo(Hz)	-0.38
MDVP:Flo(Hz)	-0.38

classifers can recognize a dataset with full features and with diminished features between the investigation groups [[38](#page-12-15)[–41\]](#page-12-16). Figure [6](#page-10-0) shows the area under the classifier performance measurement curve.

By analyzing the ROC curve, the gradient boosting classifer has a higher performance accuracy of 98.7%, with 11 features reduced by χ chi² method.

The results obtained in this paper shows highest performance as compared to other papers. The comparison has been made in following Table [6.](#page-11-22)

Fig. 4 Highly correlated features by correlation matrix

No. of features	$%$ Accuracy				RMSE $(\%)$
	Without feature selection (23-Attrib- utes)	$Chi2$ feature selection Technique (11- Attrib- utes)	Extra trees classifier feature selection Technique (11- Attributes)	Correlation matrix feature selection technique (11- Attributes)	
DT	94.87	94.87	94.87	94.87	6.85
KNN	82.05	82.05	82.05	79.48	7.98
RF	92.30	92.30	89.74	87.17	6.96
Bagging	92.30	89.74	89.74	87.17	6.72
AdaBoosting	87.17	84.61	87.17	89.74	5.52
Gradient boosting 92.30		94.87	92.30	94.87	6.12

Table 5 Performance measurement of the classifers with full features vs. reduced features

■ % Accuracy Without feature Selection (23-Attributes)

Conclusion

This research aims to solve the problem of speech performance execution in Parkinson's disease using feature elimination and the execution of multiple classifers. Parkinson's disease is a dangerous human disease, and diferent people all over the world have experienced this disease. In this way, a reliable method is needed to fully confrm PD. In this article, we propose a reliable technique that uses appropriate AI to confrm the approach of Parkinson's disease. In particular, DT, KNN, RF, Bagging, AdaBoosting, and Gradient Boosting have been applied to grouping of

Parkinson's disease and sound subjects. Chi², extra tree classifers, and merging techniques based on correlation matrix have been accepted for the selection of relevant features. In addition, the K-fold cross-validation strategy has been used to determine the ideal value of the super boundary of the best model. In addition, evaluation measurements have been used to evaluate the presentation of the proposed model. The test results show that the DT group efectively evaluated PD and physical subjects. The high presentation of our strategy is due to the feature selection that determines the high enough features of the calculation. In terms of accuracy, the proposed strategy achieved amazing results and

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achieved 94.87% accuracy and AUC (%98.7). In addition, the suggested strategy can be easily used in medical service associations. In future work, since deep neural tissue will naturally select appropriate features for characterization and

Fig. 6 Area Under the Classifer (AUC) performance measurement curve. **a** Represents ROC curve with 23 features, **b** Represents chi2 feature selection (11 features), **c** Represents extra trees classifer fea-

> artifcial intelligence calculations need to include selection calculations, in future work, deep neural work methods will be used to sort Parkinson's disease and entities. The proposed strategy will be applied to other data sets to identify

Table 6 Comparison of proposed method with other literature

Author	Machine learning clas- sifier	Accuracy $(\%)$
Li and Li (2022) [42]	LR	80.00
	SVM	84.00
	DT	59.00
	KNN	62.00
Trabassi et al. (2022) [43]	SVM	88.00
	RF	87.00
	DT	85.00
	KNN	82.00
	MLP-ANN	84.00
Mamun et al. [44]	Random Forest	92.31
	XGBoost	90.00
	AdaBoost	90.00
	Bagging	87.18
Thakur et al. $[45]$	LR	92.00
	SVM	90.00
Proposed model	DT	94.87
	KNN	82.05
	RF	92.30
	Bagging	92.30
	AdaBoosting	89.74
	Gradient Boosting	94.87

comparative types of diseases. Treatment and recuperation after a given illness are of the utmost importance. In this way, we will gradually reduce infection and apply it for recuperation in future.

Funding The authors received no fnancial support for the research, authorship, and/or publication of this article.

Data Availability The data set used in this exploration work can be found on UCI machine learning repository.

Declarations

Conflict of Interest The authors have no affiliation with any organization with a direct or indirect fnancial interest in the subject matter discussed in the manuscript.

References

- 1. S.Y. Lim, S.H. Fox, A.E. Lang, Overview of the extranigral aspects of Parkinson disease. Arch. Neurol. **66**(2), 167–172 (2009)
- 2. S. Perez-Lloret, M.V. Rey, A. Pavy-Le Traon, O. Rascol, Emerging drugs for autonomic dysfunction in Parkinson's disease. Expert Opin. Emerg. Drugs **18**(1), 39–53 (2013)
- 3. K. Seppi, D. Weintraub, M. Coelho, S. Perez-Lloret, S.H. Fox, R. Katzenschlager et al., The Movement Disorder Society evidencebased medicine review update: treatments for the non-motor

symptoms of Parkinson's disease. Movement Dis. **26**(S3), S42– S80 (2011)

- 4. K.H. Yu, A.L. Beam, I.S. Kohane, Artifcial intelligence in healthcare. Nat. Biomed. Eng. **2**(10), 719–731 (2018)
- 5. L. Ma, T. Fu, T. Blaschke, M. Li, D. Tiede, Z. Zhou et al., Evaluation of feature selection methods for object-based land cover mapping of unmanned aerial vehicle imagery using random forest and support vector machine classifers. ISPRS Int. J. Geo-Inform. **6**(2), 51 (2017)
- 6. A.D. Macleod, I. Dalen, O.B. Tysnes, J.P. Larsen, C.E. Counsell, Development and validation of prognostic survival models in newly diagnosed Parkinson's disease. Mov. Disord. **33**(1), 108–116 (2018)
- 7. M. Little, P. McSharry, E. Hunter, J. Spielman, L. Ramig, Suitability of dysphonia measurements for telemonitoring of Parkinson's disease. Nature Precedings (2008). [https://doi.org/10.1038/npre.](https://doi.org/10.1038/npre.2008.2298.1) [2008.2298.1](https://doi.org/10.1038/npre.2008.2298.1)
- 8. C.O. Sakar, O. Kursun, Telediagnosis of Parkinson's disease using measurements of dysphonia. J. Med. Syst. **34**(4), 591–599 (2010)
- 9. D.C. Li, C.W. Liu, S.C. Hu, A fuzzy-based data transformation for feature extraction to increase classifcation performance with small medical data sets. Artif. Intell. Med. **52**(1), 45–52 (2011)
- 10. Spadoto, A. A., Guido, R. C., Carnevali, F. L., Pagnin, A. F., Falcão, A. X., & Papa, J. P. (2011). Improving Parkinson's disease identifcation through evolutionary-based feature selection. In *2011 Annual International Conference of the IEEE Engineering in Medicine and Biology Society* (pp. 7857–7860). Ieee.
- 11. M. Gök, An ensemble of k-nearest neighbours algorithm for detection of Parkinson's disease. Int. J. Syst. Sci. **46**(6), 1108–1112 (2015)
- 12. M. Peker, B. Sen, D. Delen, Computer-aided diagnosis of Parkinson's disease using complex-valued neural networks and mRMR feature selection algorithm. J. Healthcare Eng. **6**(3), 281–302 (2015)
- 13. L. Naranjo, C.J. Perez, J. Martin, Y. Campos-Roca, A two-stage variable selection and classifcation approach for Parkinson's disease detection by using voice recording replications. Comput. Methods Programs Biomed. **142**, 147–156 (2017)
- 14. Z. Cai, J. Gu, H.L. Chen, A new hybrid intelligent framework for predicting Parkinson's disease. IEEE Access **5**, 17188–17200 (2017)
- 15. A.U. Haq, J.P. Li, M.H. Memon, A. Malik, T. Ahmad, A. Ali et al., Feature selection based on L1-norm support vector machine and efective recognition system for Parkinson's disease using voice recordings. IEEE Access **7**, 37718–37734 (2019)
- 16. S. Yadav, M.K. Singh, Hybrid machine learning classifer and ensemble techniques to detect Parkinson's disease patients. SN Computer Sci. **2**(3), 1–10 (2021)
- 17. R. Das, A comparison of multiple classifcation methods for diagnosis of Parkinson disease. Expert Syst. Appl. **37**(2), 1568–1572 (2010)
- 18. A. Tsanas, M.A. Little, P.E. McSharry, L.O. Ramig, Nonlinear speech analysis algorithms mapped to a standard metric achieve clinically useful quantifcation of average Parkinson's disease symptom severity. J. R. Soc. Interface **8**(59), 842–855 (2011)
- 19. J. Howell, When technology is too hot, too cold or just right. Emerg. Learn. Design J. **5**(1), 2 (2017)
- 20. C.W. Hsu, C.J. Lin, A comparison of methods for multiclass support vector machines. IEEE Trans. Neural Netw. **13**(2), 415–425 (2002)
- 21. H.L. Chen, G. Wang, C. Ma, Z.N. Cai, W.B. Liu, S.J. Wang, An efficient hybrid kernel extreme learning machine approach for early diagnosis of Parkinson's disease. Neurocomputing **184**, 131–144 (2016)
- 22. N. Singh, V. Pillay, Y.E. Choonara, Advances in the treatment of Parkinson's disease. Prog. Neurobiol. **81**(1), 29–44 (2007)
- 23 X. Wu, V. Kumar, J.R. Quinlan, J. Ghosh, Q. Yang, H. Motoda et al., Top 10 algorithms in data mining. Knowl. Inform. Syst. **14**(1), 1–37 (2008)
- 24. F. Pernkopf, Bayesian network classifers versus selective k-NN classifer. Pattern Recogn. **38**(1), 1–10 (2005)
- 25. V. Chaurasia, S. Pal, Applications of machine learning techniques to predict diagnostic breast cancer. SN Comput. Sci. **1**(5), 1–11 (2020)
- 26. M.K. Pandey, M.K. Singh, S. Pal, B.B. Tiwari, Prediction of phishing websites using stacked ensemble method and hybrid features selection method. SN Comput. Sci. **3**(6), 488 (2022)
- 27. Z. Soumaya, B.D. Taoufq, N. Benayad, B. Achraf, A. Ammoumou, A hybrid method for the diagnosis and classifying parkinson's patients based on time–frequency domain properties and K-nearest neighbor. J. Med. Sig. Sensors **10**(1), 60 (2020)
- 28. R. Aggrawal, S. Pal, Sequential feature selection and machine learning algorithm-based patient's death events prediction and diagnosis in heart disease. SN Comput. Sci. **1**(6), 344 (2020)
- 29. H. Byeon, Best early-onset Parkinson dementia predictor using ensemble learning among Parkinson's symptoms, rapid eye movement sleep disorder, and neuropsychological profle. World J. Psychiatr. **10**(11), 245 (2020)
- 30. A.K. Tiwari, Machine learning based approaches for prediction of Parkinson's disease. Mach. Learn Appl. **3**(2), 33–39 (2016)
- 31. L. Ali, C. Zhu, N.A. Golilarz, A. Javeed, M. Zhou, Y. Liu, Reliable Parkinson's disease detection by analyzing handwritten drawings: construction of an unbiased cascaded learning system based on feature selection and adaptive boosting model. IEEE Access **7**, 116480–116489 (2019)
- 32. I. Karabayir, S.M. Goldman, S. Pappu, O. Akbilgic, Gradient boosting for Parkinson's disease diagnosis from voice recordings. BMC Med. Inform. Decis. Mak. **20**(1), 1–7 (2020)
- 33. V. Chaurasia, S. Pal, Stacking-based ensemble framework and feature selection technique for the detection of breast cancer. SN Computer Sci. **2**(2), 1–13 (2021)
- 34 V. Chaurasia, A. Chaurasia, Novel method of characterization of heart disease prediction using sequential feature selection-based ensemble technique. Biomed. Mater. Dev. (2023). [https://doi.org/](https://doi.org/10.1007/s44174-022-00060-x) [10.1007/s44174-022-00060-x](https://doi.org/10.1007/s44174-022-00060-x)
- 35. V. Chaurasia, S. Pal, Data mining techniques: to predict and resolve breast cancer survivability. Int. J. Comput. Sci. Mobile Computing IJCSMC **3**(1), 10–22 (2014)
- 36. Chaibub Neto, E. L. I. A. S., Bot, B. M., Perumal, T., Omberg, L., Guinney, J., Kellen, M., et al. (2016). Personalized hypothesis

tests for detecting medication response in Parkinson disease patients using iPhone Sensor data. In *Biocomputing 2016: Proceedings of the Pacifc Symposium* (pp. 273–284).

- 37. A. Zhan, S. Mohan, C. Tarolli, R.B. Schneider, J.L. Adams, S. Sharma et al., Using smartphones and machine learning to quantify Parkinson disease severity: the mobile Parkinson disease score. JAMA Neurol. **75**(7), 876–880 (2018)
- 38. <https://archive.ics.uci.edu/ml/datasets/parkinsons>Accessed 4 July 2021
- 39. Yadav, D. C., & Pal, S. (2022). Measure the superior functionality of machine intelligence in brain tumor disease prediction. In *Artifcial Intelligence-Based Brain-Computer Interface* (pp. 353–368). Academic Press, London.
- 40. D.C. Yadav, S. Pal, An ensemble approach for classifcation and prediction of diabetes mellitus disease, in *Emerging trends in data driven computing and communications*. ed. by R. Mathur, C.P. Gupta, V. Katewa, D. SinghJat, N. Yadav (Springer, Singapore, 2021), pp. 225–235
- 41. S. Pal, Chronic kidney disease prediction using machine learning techniques. Biomed. Mater. Dev. (2022). [https://doi.org/10.1007/](https://doi.org/10.1007/s44174-022-00027-y) [s44174-022-00027-y](https://doi.org/10.1007/s44174-022-00027-y)
- 42. A. Li, C. Li, Detecting parkinson's disease through gait measures using machine learning. Diagnostics **12**(10), 2404 (2022)
- 43. D. Trabassi, M. Serrao, T. Varrecchia, A. Ranavolo, G. Coppola, R. De Icco et al., Machine learning approach to support the detection of Parkinson's disease in IMU-based Gait analysis. Sensors **22**(10), 3700 (2022)
- 44. Mamun, M., Mahmud, M. I., Hossain, M. I., Islam, A. M., Ahammed, M. S., & Uddin, M. M. (2022, October). Vocal Feature Guided Detection of Parkinson's Disease Using Machine Learning Algorithms. In *2022 IEEE 13th Annual Ubiquitous Computing, Electronics & Mobile Communication Conference (UEMCON)* (pp. 0566–0572). IEEE.
- 45. M. Thakur, S. Dhanalakshmi, H. Kuresan et al., Automated restricted Boltzmann machine classifer for early diagnosis of Parkinson's disease using digitized spiral drawings. J. Ambient Intell. Human Comput. **14**, 175–189 (2023). [https://doi.org/10.](https://doi.org/10.1007/s12652-022-04361-3) [1007/s12652-022-04361-3](https://doi.org/10.1007/s12652-022-04361-3)

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