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the Detection of PCOS Using Machine learning Algorithms and Feature Selection by K-Means clustering

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Abstract:

polycystic ovarian syndrome, which affects between 5 and 10 percent of adolescent women, is one of the most prevalent endocrine system conditions (PCOS). Infertility and failure to ovulate are symptoms, as are cardiovascular conditions, type 2 diabetes, etc. PCOS can be found by biochemical, clinical, and ultrasonographic techniques. It is well recognized that early detection and intervention can lower the risk of developing PCOS. Therefore, it is essential to understand which classification model and features contribute significantly to the prediction of disease, which is the goal of this study. Despite the employment of several tools, Naïve Bayes exhibits accuracy performances of 89.51% with 6 chosen features.

Keywords: PCOS, Genetic algorithm, Machine learning, KNN, SVM.

Introduction:

Polycystic ovary syndrome or PCOS, is a widely known endocrine disorder in women distinguished by hyperandrogenism during reproductive age [1]. This syndrome was initially reported by Stein and Leventhal in 1935 and it was known for a long time as the Stein-Leventhal syndrome [2]. Women with PCOS often could suffer from menstrual problems and infertility issues [3]. Moreover, it may contribute to long-term health problems like diabetes and heart disease, cancer of the uterus and mood disorders [4]. The exact causes of PCOS are very complex and still not clear. There are many suggested etiologies for PCOS, but it is not fully supported, in general, this hormonal imbalance consists of a combination of excess androgen and insulin resistance[5]. In addition to environmental and genetic factors that contribute to this hormonal imbalance, all of these causes the development of PCOS[6]. Symptoms and signs vary among women with PCOS, it includes a combination of hyperandrogenism (hirsutism, alopecia, acne, high blood testosterone), obesity and severe menstrual irregularity [7]. According to the "Rotterdam

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European Society for Human Reproduction and Embryology and the American Society for Reproductive Medicine (ESHRE/ASRM) Sponsored PCOS Consensus Workshop" (Rotterdam ESHRE/ASRM, 2004), the presence of two of the three diagnosis criteria: "oligomenorrhoea, polycystic ovaries morphology, and hyperandrogenemia" [8].

The various machine learning classification methods have been used for the detection of various diseases such as breast cancer, heart and ovarian, etc. [13]. Some of the classification algorithms used for the prediction of PCOS dataset are reviewed below:

Denny, Amsey, et al. (2019). [9], Identified PCOS using 8 features. Principal Component Analysis was used to minimize the extracted features a random forest classifier was utilized and it was obtained an accuracy of 89.02%.

He proposed **V. Thakre1 and et al. (2020).** [10], proposed a number of statistical parameters with five machine learning classifiers to detct PCOS. In that study, found that the random forest classifier is more reliable than the others, with an accuracy of 90.9%.

Silva, I. S., et al. (2021). [11], they used 72 patients with PCOS and 73 healthy women. Focusing on only 10 features. For data classification, they suggested BorutaShap method, followed by the Random Forest algorithm, which gave an accuracy of 86%.

Munjal, Sanjay K., et al. (2020). [12], proposed a genetic algorithm to select the most representative features to detect PCOS. Several than one classifier including extra trees, random forest, and decision tree were adopted in that It was the best result Extra trees with 88% accuracy.

2. DIFFERENT MACHINE LEARNING ALGORITHMS

There are a number of algorithms used in Machine Learning for the prediction of target values based on various input values. The algorithms taken under use for the prediction of PCOS are as under:

2.1 Support Vector Machine (SVM)

A classification technique, where each data item is plotted in n- dimensional space as a point. It comes under supervised machine learning model. The support vectors lie near the margin of the classifier.

2.2 Naïve Bays

One of the powerful classification methods evolved on Bays' theorem with independence between predictors as an assumption. The Naïve bays classifier assumes that there is no relation between the presences of a particular feature over the other features. The status of a particular feature does not affect the status of another feature.

2.3 Decision tree

Decision tree is a tree structure which looks similar as flow chart. Every node in the tree represents the test of an attribute, every branch represents the output of the test, and every leaf means a class or distribution of classes. The top node is root node, from root node to one leaf consists a classification rule. So, decision tree is easy to be transferred to classification rules. It has many algorithms, but the main idea

is using from top to bottom induction method. And the most important part is choosing which attribute to be the node as well as the evaluation of whether the tree is correct.

2.4 KNN

Nearest-neighbor classifiers are based on learning by analogy, that is, by comparing a given test tuple with training tuples that are similar to it. The training tuples are described by n attributes. Each tuple represents a point in an n-dimensional space. In this way, all the training tuples are stored in an n-dimensional pattern space. When given an unknown tuple, K-NN classifier searches the pattern space for the k training tuples that are closest to the unknown tuple. These k training tuples are K-NN of the unknown tuple.

3. OBJECTIVES

A. To diagnose PCOS based on clinical symptoms associated with the using popular machine learning algorithms on random data samples.

B. To compare performance of different algorithms and determine the best possible algorithm among them.

4. METHODOLOGY

The sound methodology is the key of a successful research. The methodology adopted to carry on this study is shown in the figure 1.



Fig. 1. Block diagram of the proposed method

4.1 Data Sources and Attributes Description

The PCOS dataset for this study was obtained from Kaggle [14]. From previous studies, it is clear that researchers and clinicians are using different disease datasets to study machine learning classification methods, similar to the PCOS dataset [13]. The original PCOS dataset contains 541 instances with 42 attributes, one of which is the patient file number (not taken into consideration for data analysis). Finally, there are 41 attributes in total, with 40 as input attributes and PCOS as a class label [Positive (Yes) and Negative (No)]. The dataset has an uneven distribution of class labels (i.e., 364 instances of class label = 0 and 177 instances of class label =1) and missing values. The data was collected from ten different hospitals in Kerala, India, and is now available on the Kaggle website. Continuous, nominal, and ordinal expressions The description of the attributes is shown in Table 1.

4.2 Preprocessing of data

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The data is preprocessed to identify missing values before being used to diagnose PCOS via various machine learning algorithms.

| No | Attributes | No | Attributes | | | | | |
|------------------------------|------------------------|-------------------------|----------------|---------------------------|--|-------------------------|-------------|-----------|
| 1 | Patient File number | | 22 Thyroi | d-Stimulating H | ormone: T | SH (mll | J/L) 2 | PCOS |
| (clas | ss label) | 23 Anti | -Müllerian Ho | ormone: AMH (r | ng/mL) | 3 | Age (Yrs) | |
| 24 | Prolactin: PRL (ng/mL) |) | | 4 Weight (F | <g)< td=""><td></td><td>25</td><td>5 Vit D3</td></g)<> | | 25 | 5 Vit D3 |
| (ng/ | /mL) | 5 | Height (Cr | n) | 2 | 26 Pro | ogesteron | e: PRG |
| (ng/ | /mL) 6 | BMI: body | mass index | 27 | BP _Syste | olic (mn | nHg) | |
| 7 | Blood Group | | 28 Rando | m Blood Sugar: | RBS (mg/c | II) | 8 Pi | ulse rate |
| (bpm) 29 Weight gain (Y/N) 9 | | | | RR (| breaths/n | ths/min) | | |
| 30 | hair growth (Y/N) | | 10 | Haemoglobin: | Hb(g/dl) | | 31 9 | Skin |
| darl | kening (Y/N) | | 11 Menstrua | I Cycle: Cycle(R | /I) | 32 H | air loss (Y | /N) |
| 12 | Cycle length (days) | | 33 Pimple | es (Y/N) | | | 13 | Marriage |
| Stat | cus (Yrs) 34 | 4 Fast fo | od (Y/N) | | 14 | 1 Pregr | nant (Y/N) | |
| 35 | Reg. Exercise (Y/N) | | 15 | 5 No. of abortio | ons | | 36 | BP |
| _Sys | stolic (mmHg) | | 16 Follicle | e No. (R) | | 37 | BP _Dias | tolic |
| (mmHg) | | 17 LI | 17 LH (mIU/mL) | | 38 | 38 Follicle stimulating | | |
| hormone: FSH (mIU/mL) | | 18 | 18 FSH/LH | | 39 | 39 Follicle No. (L) | | |
| 19 | Hip (inch) | | 40 Avg. F | ^s ize (L) (mm) | | | 20 V | Vaist |
| (inch) | | 41 Avg. F size (R) (mm) | | | 21 Waist: Hip Ratio | | | |
| 42 | Endometrium (mm) 4 | | | | | | | |

Table 1 Attributes description and their units

4.3 k-means based feature selection

We proposed a feature selection algorithm based on k mean clustering. The proposed model used K-mean clustering as a feature selection using city block distance [15]. It was found that the K-mean clustering was capable to select a high the most influential features and eliminating the irrelevant ones. In this experiment, six features were selected and fed into classifiers. The extracted features were individually tested, and all results were recorded. Table 2 shows the detection rate based on features. The follicle feature scored the highest detection compared to other features. However, weight gain was recorded with the lowest detection rate.

| TABLE 2. RANKING OF BEST 6 FEATURES using k-means ARRANGED IN DESCENDING ORDER | | | | | | |
|--|---------|--|--|--|--|--|
| Features | Weights | | | | | |
| Follicle No. (L) | 78.62 | | | | | |
| Skin darkening | 77.50 | | | | | |
| Follicle No. (R) | 77.32 | | | | | |

| hair growth | 77.32 |
|--------------------|-------|
| Cycle length(days) | 74.53 |
| Weight gain | 74.34 |

4.4 Classification

We use MATLAB to implement classification algorithms, diagnose problems, and validate model performance. To diagnose PCOS, seven machine learning algorithms were trained and evaluated on random samples of data with 42 independent variables as symptoms. The dependent variable PCOS has a strong correlation with these independent variables and has two possible values: '1' or '0.' Among the algorithms used are knn1, knn3, knn5, SVM (Support Vector Machine), Decision tree, and Nave Bayes. Table 1 depicts the list of variables used in the diagnosis of PCOS.

4.5 Validation of models

Three widely used parameters, accuracy, sensitivity, and specificity, presented in Eqs. (1) to (5), to evaluate machine learning algorithms, are used in this paper to demonstrate the capability of the proposed method to diagnose PCOS.

| $\operatorname{Recall} = \frac{TP}{TP + FN} .$ | (2) |
|---|-----|
| Precision $= \frac{TP}{TP+FP}$. | (3) |

$$F\text{-measure}=2 \times \frac{\text{Precision} \ast \text{Recall}}{\text{Precision} + \text{Recall}}$$
(4)

$$Specificity = \frac{TN}{TN + FP}$$
(5)

accuracy =
$$\frac{TP + TN}{TP + TN + FP + FN}$$
. (6)

TP shows a person does not have PCOS and identified as a nonPCOS patient, and TN shows a PCOS patient correctly identified as a PCOS patient. FN shows the patient has PCOS but is predicted as a healthy person. Moreover, FP shows the patient is a healthy person but predicted as a PCOS patient.

5. RESULTS

In this section, the effect of k means on PCOS detection was discussed. Table 3 shows the classification results for different classifiers. the Linear SVM algorithm and naive bayes algorithm both gave the same accuracy which is 89.51%. However, the naive bayes algorithm recorded an increase in Sensitivity, F-Measure, Prec., and Specificity, reaching 85.94%, 86.61%, 87.30%, and 91.84%, respectively. While in the Linear SVM algorithm, the rating scales recorded 84.91%, 84.11%, 83.33% and 91.74% for Sensitivity, F-Measure, Prec. and Specificity, respectively.

6. CONCLUSION

PCOS is a disorder caused by hormonal imbalance in young women's bodies, and it is a very common problem that affects a large percentage of women worldwide. An early diagnosis of the disorder can aid in its treatment and management. We applied popular machine learning algorithms, such as Linear SVM, Gaussian SVM, Decision Tree, Nave Bayes, KNN1, KNN3, and KNN5, to clinical data of PCOS patients

Table 3. Typical performance test results for machine learning algorithms using Selected features by K-mean cluster.

| | | | | | | | Conf | usion | | |
|------|---------------|-------------|------------|-------|-------------|----------|---------|-------|---|---------|
| S.N. | Network | Sensitivity | <i>F</i> - | Prec. | Specificity | Accuracy | Matrix | | | |
| | | (%) | Measure | (%) | (%) | (%) | Predict | | | |
| | | | (%) | | | | Class | | | |
| | | | | | | | 0 | 1 | | |
| 1 | Linear SVM | 84.91 | 84.11 | 83.33 | 91.74 | 89.51 | 100 | 9 | 0 | |
| | | | | | | | 8 | 45 | 1 | |
| 2 | Gaussian SVM | 83.33 | 81.82 | 80.36 | 89.81 | 87.65 | 97 | 11 | 0 | |
| | | | | | | | 9 | 45 | 1 | SS |
| 3 | Decision Tree | 78.85 | 80.39 | 82.00 | 91.82 | 87.65 | 101 | 9 | 0 | Cla |
| | | | | | | | 11 | 41 | 1 | ən. |
| 4 | Naïve Bayes | 85.94 | 86.61 | 87.30 | 91.84 | 89.51 | 90 | 8 | 0 | ^{4}L |
| | | | | | | | 9 | 29 | 1 | |
| 5 | KNN1 | 84.62 | 87.57 | 73.33 | 90.24 | 88.89 | 111 | 12 | 0 | |
| | | | | | | | 6 | 33 | 1 | |
| 6 | KNN3 | 76.92 | 80.00 | 83.33 | 92.73 | 87.65 | 102 | 8 | 0 | |
| | | | | | | | 12 | 40 | 1 | |
| 7 | KNN5 | 74.55 | 78.85 | 83.67 | 92.52 | 86.42 | 99 | 8 | 0 | |
| | | | | | | | 14 | 41 | 1 | |

based on symptoms. The performance validation metrics recall, accuracy, precision, and F- statistics showed that the Nave Bayes algorithm performed best in diagnosing PCOS with an accuracy of 89.51 percent, followed by the Linear SVM algorithm with an accuracy of 89.51 percent. Thus, based on the data, it is concluded that the Nave Bayes algorithm is the best suitable algorithm for PCOS diagnosis. The study's future scope may include the use of various or large data sets for disease diagnosis.

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