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Diagnosis of Prostate Cancer using Soft Computing Paradigms

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Diagnosis of Prostate Cancer using Soft Computing Paradigms

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Abstract- The process of diagnosing of prostate cancer using traditional methods is cumbersome because of the similarity of symptoms that are present in other diseases. Soft Computing (SC) paradigms which mimic human imprecise data manipulation and learning capabilities have been reviewed and harnessed for diagnosis and classification of prostate cancer. SC technique based on Adaptive Neuro-Fuzzy Inference System (ANFIS) facilitated symptoms analysis, diagnosis and prostate cancer classification. Age of Patient (AP), Pains in Urination (PU), Frequent Urination (FU), Blood in Semen (BS) and Pains in Pelvic (PP) served as input attributes while Prostate Risk (PR) served as output. Matrix laboratory provided the programming tools for system implementation. The practical function of the system was assessed using prostate cancer data collected from the University of Uyo Teaching Hospital. A 95% harmony observed between the computed and the expected output in the ANFIS model, showed the superiority of the ANFIS model over the fuzzy model. The system is poised to assist medical professionals in the domain of diagnosis and classification of prostate cancer for the promotion of management and treatment decisions.

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I. Introduction

Prostate cancer is a common disease in elderly men (Leonard, 2008; Ajape & Babatunde, 2009; Thomas, 2011). The rapid spread of prostate cancer disease stems from unawareness of its early symptoms. Early diagnosis and treatment of prostate cancer reduce the rate of fatality (Ifere & Ananaba, 2012; Ganesh *et al*., 2013; Mfon, 2017). Some symptoms of prostate cancer observed in other diseases make it difficult to obtain precise diagnosis using traditional and hard computing methods. Soft Computing (SC) methodology offers a plausible solution to this problem. SC emulates human processing capabilities. It harnesses imprecision, uncertainty, partial truth as well as learn from previous experience to provide solution in a seemingly impossible scenario. The principal techniques of SC are – fuzzy logic, neural networks, support vector machines, evolutionary computation and probabilistic reasoning (Kurhe *et al*., 2011). The implementation technique of SC is complementary rather than competitive. SC has been successfully applied in medical diagnosis, prediction, pattern recognition, decision support, automotive control and infrastructure monitoring (Obot and Udoh, 2013; Agu *et al*., 2015; Udoh, 2016;Mfon 2017; Udoh *et al*., 2017; Arlan *et al*., 2018). The remainder of the paper is organized in Sections. Section 2 presents related works in soft computing techniques. Section 3 addresses the design of adaptive neuro-fuzzy inference system for prostate cancer diagnosis. Implementation and discussion on the results are carried out in Section 4 while Section 5 presents the conclusion of the work and recommendation for further research.

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II. Related Works

a) Fuzzy Logic

Zadeh (1965) introduced fuzzy logic (FL) as a mathematical tool for dealing with uncertainty. The FL theory provides a mechanism for representing linguistic constructs such as "many," "low," "medium," "often," "few." It is a problem-solving methodology which provides a simple way to draw definite conclusions from vague, ambiguous or imprecise information. FL technique follows the process of fuzzification, inferencing, composition, and defuzzification (Gupta, 1995; Atınç & Kürşat, 2011; Agu *et al*., 2015; Udoh, 2016). Ismail *et al*. (2003) presented a fuzzy logic expert system for classification of prostate cancer risk based on identified symptoms. Input parameters used were Prostate Specific Antigen (PSA), Age and Prostate Volume (PV) while Prostate Cancer Risk (PCR) served as output. The few input parameters employed hindered adequate prostate cancer risk classification. Ganesh *et al*. (2013) presented a prostate cancer hospital-based survival study to aid early diagnosis and remedy of prostate cancer. Lack of detailed clinical analysis and inclusion of intelligent tools for precise diagnosis were weaknesses of the work. Mfon (2017) investigated the intensity of prostate cancer using Mamdani reasoning mechanism of fuzzy logic. The system could diagnose and classify prostate cancer patients but lacked the cognitive ability to learn from previous data of prostate cancer patients.

b) Neural Network

Neural Network (NN) is an information processing paradigm that is inspired by the way biological nervous systems, process information and learns from previous patterns (Akinyokun, 2007; Udoh, 2016). Javed *et al*. (2001) used NN to classify cancers

into four distinct diagnostic categories based on their gene expression signatures. The study demonstrated the potential application of NN for tumor diagnosis and identification of a candidate for therapy. Misop *et al*. (2001) studied a large series of patients with clinically localized prostate carcinoma. The clinical and pathologic data obtained at the time of prostate biopsy were used to develop a NN for prostate cancer risk classification. Bob *et al*. (2002) compared the predictive capabilities of NN and conventional statistical model. Prostate Specific Antigen (PSA) levels were employed in NN together with multivariate analysis for the detection of prostate cancer. The predictive accuracy of NN was superior to that of the statistical model. Joseph & David (2006) surveyed machine learning techniques for prostate cancer prediction. NN models performed better than decision tree models. Maysam & Feddie (2007) reported on the application of artificial intelligence technology in prostate cancer management. While most researchers focused on NN to improve the diagnosis and prognostic prediction, others explored expert systems and fuzzy modeling approaches. The lack of transparency of NN processing technique hinders global scientific community acceptance. However, this could be handled by neuro-fuzzy paradigm.

c) Neuro-Fuzzy Paradigm

Neuro-fuzzy model combines the capabilities of NN and FL (Akinyokun, 2007; Udoh, 2016). Benecchi (2006) proposed a neuro-fuzzy system for predicting the presence of prostate cancer. The system made use of a co-active neuro-fuzzy inference model. The predictive ability of neuro-fuzzy system performed better than that obtained by a total prostate specific antigen. Kuo *et al*. (2015) proposed a fuzzy neural network (FNN) system for prognosis of prostate cancer. The use of cluster analysis helped in the determination of the initial membership function parameters. An integration of artificial immune network and a particle swarm optimization assisted the investigation of input-output relationships. FNN algorithm gave a satisfactory prediction in prostate cancer prognosis. Cosma *et al.* (2016) proposed a neuro-fuzzy model for prediction of pathological state in patients with prostate cancer. The receiver operating characteristic (ROC) points obtained from neuro-fuzzy approach performed better than those obtained from fuzzy c-means, support vector machine (SVM) and Naïve Bayes classifiers. Mustain & Nazrul (2016) presented an ANFIS for predicting lung cancer risk. Cancer historical data were collected from hospital and preprocessed. The use of linear discriminant analysis facilitated attributes dimension reduction for cancer classification. Arlan *et al*. (2018) presented the training of the ANFIS with genetic algorithm for diagnosis of prostate cancer. The results of cancer gene profiles classification were satisfactory and superior to results obtained from neural networks.

III. Methodology

The method followed for prostate cancer diagnosis in this work is depicted in Figure 1. It comprises four major stages namely: 1. Data collection and preprocessing; 2. ANFIS design and training 3; ANFIS parameters checking and 4. Prostate Cancer Diagnosis.

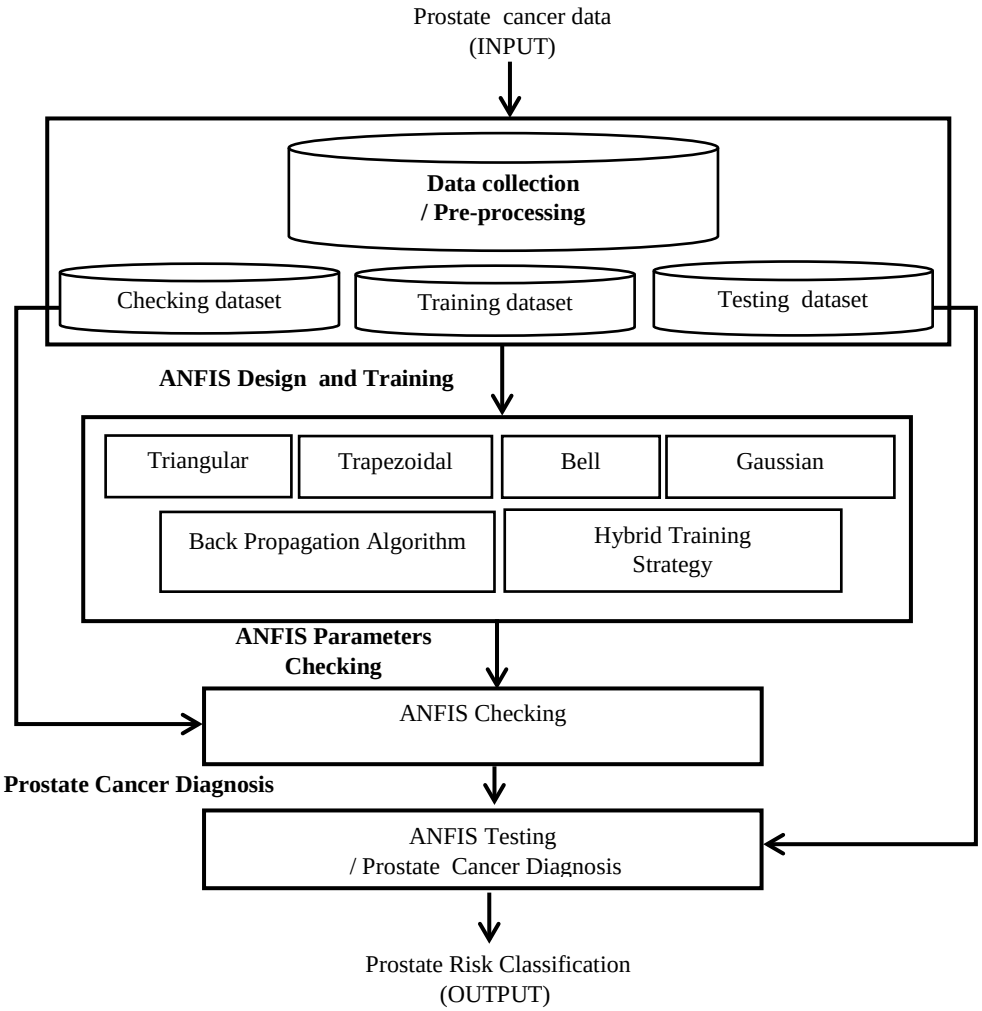

Figure 1: Block Structure of Stages in Prostate Cancer Diagnosis

a) Data Collection and Processing

A collection of 510 prostate cancer dataset within nine months (July 2017 to March 2018) from the University of Uyo Teaching Hospital, Uyo, (UUTH), Nigeria, assisted the assessment of the practical function of the system. The attributes: Age of Patient (AP), Pain in Urination (PU), Frequent Urination (FU), Blood in Semen (BS) and Pains in Pelvic (PV) served as input while Prostate Risk (PR) served as output. The splitting of the dataset in the ratio of 8:1:1, translated into 408, 51 and 51 datasets for system training, checking and testing respectively.

b) ANFIS Design and Training

ANFIS design consists of five layers. The first and the fourth layers consist of adaptive nodes which have parameters to be learned while the second, third and fifth layers are fixed nodes and contain no learning parameters. The system employed Sugeno inference mechanism whose reasoning methodology shows the output of each rule as a sequential combination of each rule input variable plus the constant term as shown in Equation 1.

IF a is X_1 AND b is Y_1 AND… AND c is Z_1 THEN $f_1 = p_1a + q_1b + \ldots + r_1c + s_1$ (1)

where a, b, c are the inputs or antecedent parameters, X, Y, Z are the fuzzy sets of inputs parameters, f is the fuzzy set of output parameters and p, q, r, and s are consequent parameters.

Layer 1 is the input layer. It has AP, PU, FU, BS, and PP as inputs. Every node i in layer 1 has a node function

$$O_i^1 = \mu X_i(a) \quad (2)$$

where a is the input to node i, and X_i is the linguistic label (Low, Moderate and High) associated with this node function. O_i^1 is the membership function of X_i and it specifies the degree to which the given input satisfies the quantifier X_i. Different Types of Membership functions such as Triangular, Bell,

Gaussian and Trapezoidal are employed in ANFIS. The general form of a triangular membership function is shown in Equation 3.

$$\mu_X(a) = \begin{cases} 1 & \text{if } a = y \\ \dfrac{a-x}{y-x} & \text{if } x \le a < y \\ \dfrac{z-a}{z-y} & \text{if } y \le a < z \\ 0 & \text{if } z \le a \end{cases} \quad (3)$$

where x, y, z are the parameters of the membership function (MF) governing triangular shape, X_i is the linguistic variable, *a* is prostate cancer input, *x* and *y* are the parameters of the membership function such that $x \le a < y$. Layer 2 is the rule node. Every node in layer 2 computes the firing strength of each rule as given in Equation 4. Layer 3 is the normalization layer. Every node in layer 3 calculates the ratio of the ith rule's firing strength to the sum of all rules's firing strengths as shown in Equation 4. Layer 4 is the defuzzification layer which consists of consequent nodes for computing the contribution of each rule to the overall output as shown in Equation 6. Layer 5 is the output layer (a single node that computes the overall output, Prostate Risk (PR). The output as shown in Equation 7 is computed as summation of prostate cancer signals.

$$O_i^2 = w_i = \mu X_i(a) * \mu Y_i(b) * \mu Z_i(c) \quad (4)$$

$$O_i^3 = \overline{w}_i = \frac{w_i}{\sum_{1=1}^{n} w_i} \quad (5)$$

$$O_i^4 \equiv \overline{w}_i f_i = \overline{w}_i (p_i a + q_i b + \ldots + r_i c + s_i) \quad (6)$$

$$O_i^5 = \sum_i \overline{w}_i f_i = \frac{\sum_i w_i f_i}{\sum_i w_i} \quad (7)$$

The training and parameters adjustments in ANFIS are facilitated either by hybrid learning algorithm or the back propagation algorithm. The hybrid learning algorithm converges faster than the traditional back propagation method. It comprises the combination of least square method in the forward pass and back propagation gradient descent procedure in the backward pass. In the forward pass, the node output goes forward until layer 4 and the consequent parameters are updated by least square method. In the backward pass, the error signal propagates backwards and the premise parameters are updated by gradient method. (Udoh *et al.,* 2017).

IV. Results and Discussion

The system as shown in Figure 2 was implemented in an environment characterized by MatLab 2015a programming tools. Prostrate cancer data samples of sizes 408, 51 and 51 records facilitated system training, checking and testing respectively. Figures 3 and 4 depict the loading of training and checking data as well as training and checking error interface respectively. The results of training and checking errors carried out in 20 iterations using hybrid learning process with Triangular, Trapezoidal, Bells or Gaussian membership functions are presented in Table 1.

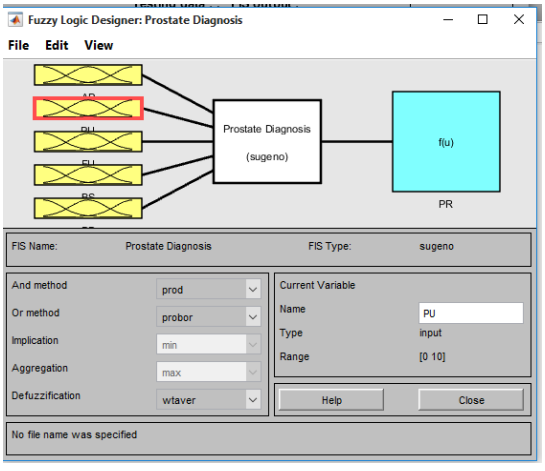

Figure 2: Prostate Cancer Diagnosis Implementation

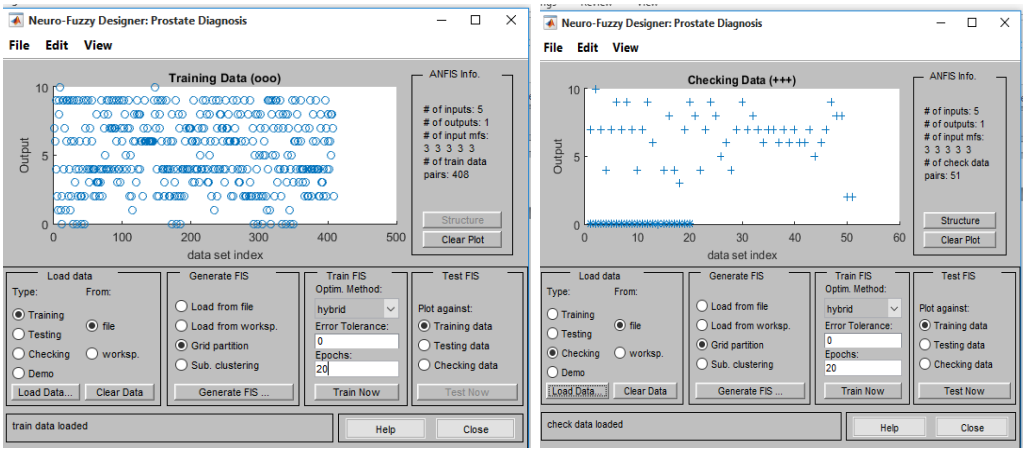

Figure 3: Loading of Training and Checking Data

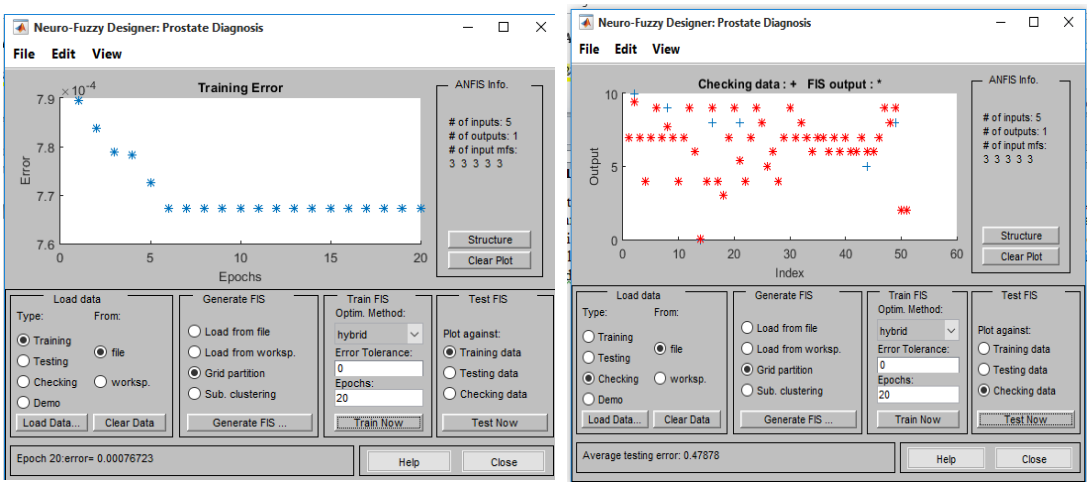

Figure 4: Training and Checking Error Interface

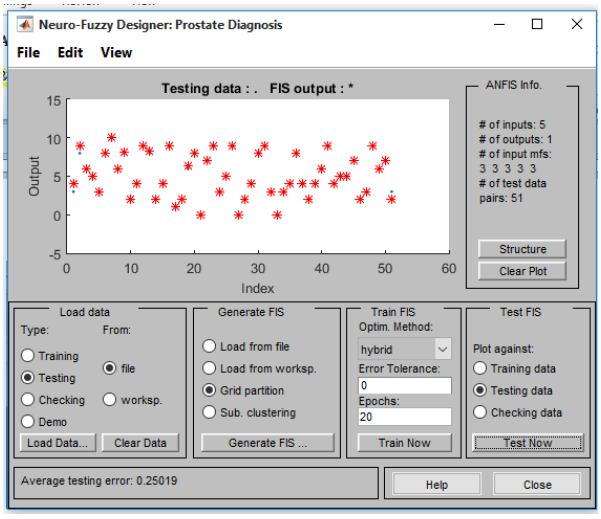

Figure 5: Testing Error Interface

As shown in Figure 5. The 51 testing data samples were loaded to ascertain the functionality of the trained and checked ANFIS. An average testing error of 0.25019 was observed between the computed and the expected output. The testing and checking errors derived from the experiment using different membership functions are depicted in Table 1.

Table 1: Training and Checking Errors Based on Different Membership Functions

Iteration No.	Triangular MF		Trapezoidal MF		Bells MF		Gaussian MF	
	Training Error	Checking Error	Training Error	Checking Error	Training Error	Checking Error	Training Error	Checking Error
1	0.000148	0.531080	0.197052	1.020400	0.002829	0.619600	0.001827	0.606912
2	0.000145	0.530330	0.197007	1.014800	0.002764	0.679068	0.001760	0.612429
3	0.000141	0.529580	0.196963	1.009600	0.002699	0.745846	0.001696	0.617943
4	0.000138	0.528840	0.196919	1.004500	0.002635	0.819051	0.001633	0.623423
5	0.000135	0.528110	0.196815	0.999700	0.002571	0.897450	0.001572	0.628843
6	0.000132	0.527370	0.196831	0.995100	0.002508	0.979547	0.001514	0.634181
7	0.000129	0.526650	0.196786	0.990700	0.002445	1.063660	0.001458	0.639416
8	0.000127	0.525920	0.196742	0.986400	0.002382	1.148020	0.001403	0.644531
9	0.000124	0.525200	0.196697	0.982300	0.002320	1.230880	0.001352	0.649510
10	0.000121	0.524490	0.196653	0.978400	0.002260	1.310590	0.001302	0.654341
11	0.000119	0.523770	0.196608	0.974600	0.002200	1.385740	0.001255	0.659013
12	0.000116	0.523070	0.196564	0.970900	0.002143	1.455150	0.001209	0.663518
13	0.000114	0.522360	0.196519	0.967400	0.002087	1.517930	0.001166	0.667850
14	0.000112	0.521660	0.196475	0.964000	0.002034	1.573510	0.001125	0.672004
15	0.000110	0.520960	0.196430	0.960685	0.001983	1.621590	0.001086	0.675976
16	0.000108	0.520270	0.196385	0.957500	0.001935	1.662100	0.001048	0.679766
17	0.000105	0.519580	0.196341	0.954400	0.001889	1.695200	0.001012	0.683374
18	0.000104	0.518895	0.196296	0.951390	0.001846	1.721220	0.000978	0.686801
19	0.000102	0.518220	0.196251	0.948500	0.001805	1.740570	0.000945	0.690049
20	0.000098	0.517540	0.196207	0.945600	0.001767	1.753790	0.000914	0.693122

Triangular MF gave the best results in terms of training and checking errors, followed by Gaussian MF. The worst checking errors were observed in Bells MF. The results of prostate cancer diagnosis using the ANFIS and the fuzzy paradigms are depicted in Figure 6. The data points in the ANFIS diagnosis matched the expected output more precisely than those in the fuzzy diagnosis. Out of the 20 data points used in the experiment, 19 data points matched with the expected output in the ANFIS model, whereas the fuzzy model had 14 similar data points. In the first instance of the diagnosis, using the ANFIS model the patient with serial number 1 had a high degree of prostate cancer. This corresponds to the expected output from domain experts.

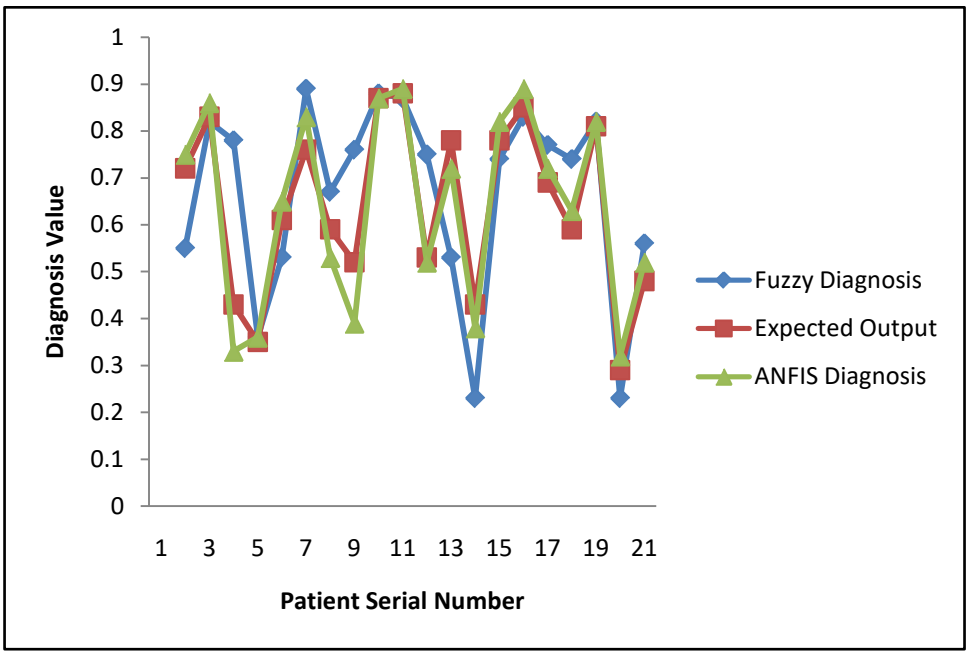

Figure 6: Graph of Prostate Cancer Diagnosis

However, using the same sets of input variables on the fuzzy model presented in (Mfon, 2017), the same patient had a moderate prostate cancer which disagrees with the expected output from domain experts. The summary of the prostate cancer diagnosis results is shown in Table 2.

Table 2: Summary of Prostate Cancer Diagnosis Results

Patient Serial Number	Patient ID Number	Fuzzy Diagnosis Value (Mfon, 2017)	Fuzzy Diagnosis Label	Expected Diagnosis Output	Expected Diagnosis Label	ANFIS Diagnosis Value (Current Study)	ANFIS Diagnosis Label	Class Grade
1	012	0.57	Moderate	0.72	High	0.75	High	A
2	121	0.80	High	0.83	High	0.82	High	A
3	300	0.78	High	0.43	Low	0.33	Low	C
4	705	0.23	Low	0.35	Low	0.36	Low	C
5	131	0.53	Moderate	0.61	Moderate	0.65	Moderate	B
6	125	0.89	High	0.76	High	0.83	High	A
7	171	0.67	Moderate	0.59	Moderate	0.53	Moderate	B
8	165	0.76	High	0.52	Moderate	0.39	Low	C
9	234	0.88	High	0.87	High	0.87	High	A
10	126	0.87	High	0.88	High	0.89	High	A
11	191	0.75	High	0.53	Moderate	0.52	Moderate	A
12	192	0.53	Moderate	0.78	High	0.72	High	B
13	158	0.23	Low	0.39	Low	0.38	Low	C
14	144	0.74	High	0.78	High	0.82	High	A
15	124	0.83	High	0.85	High	0.89	High	A
16	171	0.77	High	0.70	High	0.72	High	B
17	193	0.74	High	0.60	Moderate	0.63	Moderate	A
18	987	0.82	High	0.82	High	0.82	High	A
19	865	0.23	Low	0.29	Low	0.32	Low	C
20	166	0.56	Moderate	0.50	Moderate	0.51	Moderate	B

Both ANFIS and fuzzy models gave high diagnosis in the second instance of the diagnosis. This is in agreement with expected output from domain experts. Nevertheless, the diagnosis value of the ANFIS model was observed to be closer to that of domain experts than the one from the fuzzy model. Investigation showed that 14 out of 20 instances (70%) gave accurate prediction in the fuzzy model while 19 out of 20 instances (95%) gave accurate predictions in the ANFIS model. The results of the experiment shown in Table 2, demonstrated the precision of ANFIS model over fuzzy model in the task of prostate cancer diagnosis.

V. Conclusion and Recommendation

This paper presented a review of prostate cancer diagnosis using soft computing models. Practical function of the ANFIS paradigm was assessed in an environment characterized by matrix laboratory programming tools. The data of prostate cancer patients collected from the University of Uyo teaching hospital, Uyo, Nigeria, was used for system training and testing. A comparison of the results, showed the accuracy of the ANFIS model over the fuzzy model in the task of prostate cancer diagnosis. Future works would employ evolutionary computations and support vector machine for further investigations.

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