



Heap Based Optimization with Deep Quantum Neural Network Based Decision Making on Smart Healthcare Applications

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Abstract: The concept of smart healthcare has seen a gradual increase with the expansion of information technology. Smart healthcare will use a new generation of information technologies, like artificial intelligence, the Internet of Things (IoT), cloud computing, and big data, to transform the conventional medical system in an all-around way, making healthcare highly effective, more personalized, and more convenient. This work designs a new Heap Based Optimization with Deep Quantum Neural Network (HBO-DQNN) model for decision-making in smart healthcare applications. The presented HBO-DQNN model majorly focuses on identifying and classifying healthcare data. In the presented HBO-DQNN model, three stages of operations were performed. Data normalization is applied to pre-process the input data at the initial stage. Next, the HBO algorithm is used in the second stage to choose an optimal set of features from the healthcare data. At last, the DQNN model is exploited for healthcare data classification. A series of experiments were carried out to portray the promising classifier results of the HBO-DQNN model. The extensive comparative study reported the improvements of the HBO-DQNN method over other existing models with maximum accuracy of 97.05% and 95.72% under the colon cancer and lymphoma dataset.

Keywords: Heap-based optimization; smart healthcare; decision making; intelligent models; artificial intelligence

1 Introduction

Smart Health can be referred to as a field that grows as a subsection of Electronic Health (e-Health) and smart cities [1]. Smart cities are defined as “cities strongly originated on information and communication technology that invest in social and human capital for enhancing the standard of life of people by efficient mobility participatory governance, sustainability and wise management of resources, at the same time they guarantee the security and privacy of the people.” e-Health is defined as “an evolving domain in the incorporation of public health business and medical informatics, denoting to health services and data enhanced or delivered via the Internet and related technology [2].



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Broadly, the term symbolizes not just a technical expansion but also an attitude, a state of mind, a commitment to networked and a way of thinking, and global thinking, for the improvement of health care globally, locally, and regionally by utilizing information and communication technology [3]. As a result, computational medicine seems to be an innovative subject. Depending on computer technology and large biomedical data, computational medicine was a disciplinary subject merging mathematics, medicine, biology, computer science, and many more [4]. It employs the technique of artificial intelligence (AI) to perceptively comprehend the principle and physiological system of human illnesses through analyses in big data. It offers valuable data and direction for disease estimation, medical services, and clinical diagnosis [5–9].

Considering the pharmaceutical sector, conventionally, novel drug research will suffer from a high failure rate, long periods, and substantial investment [10]. Conversely, the research related to computing medicine could complete the preclinical drug research and expansion in approximately 1 to 2 years, with low resource consumption and a high success rate denoting that the domain of medicinal health was progressively entering the time of digitization and intelligence. But biomedical data can be sparse, high-dimension, loud, and messy, making it tough to extract the rich data behind such data efficiently [11]. Then, a suitable technique was required to process great totals of bio-medical data to gain effectual data. In machine learning (ML), DL is an optimistic gem in artificial intelligence (AI). As a division of ML, it was shown that DL was a potential technique and exceeded conventional ML in areas like speech recognition, CV, and NLP [12]. The main step of the ML technique, termed feature engineering, was to utilize domain insincerely and expert knowledge to distil attributes from data and examine the attributes by ML techniques (like SVM, RF, and so on). In the period of big data, the physical abstraction was biased and insufficient such that it could not accomplish a high-performance method for particular errands [13]. Dissimilar to conventional ML methods, DL spares the physical need to derive features, enhancing resource and time efficacy. DL was applied by a neural network (NN) containing neurons. Every layer of NNs has more neurons, and the output of the upper layer was taken as input for the successive layer [14]. The NN will transform the actual input to the output through the link among layers and the non-linear processing approach.

This study designs a new Heap Based Optimization with Deep Quantum Neural Network (HBO-DQNN) model for decision-making in smart healthcare applications. The presented HBO-DQNN model majorly focuses on identifying and classifying healthcare data. In the presented HBO-DQNN model, three stages of operations were performed. Data normalization is applied to pre-process the input data at the initial stage. Next, the HBO algorithm is used in the second stage to choose an optimal set of features from the healthcare data. At last, the DQNN model is employed for healthcare data classification. A series of experiments were carried out to portray the promising classifier results of the HBO-DQNN model.

2 Related Works

Praveen et al. [15] introduce an innovative Oppositional GSO (OGSO) method related to clustering with DNNs termed the OGSO-DNN technique for distributed healthcare mechanisms. The OGSO technique has been implemented in this paper for choosing the Cluster Head (CH) from presented IoT gadgets. The chosen CH sends data to the cloud server, which then accomplishes DNN-related classification procedures for medical diagnosis. In [16], the capsule network was implemented for ECG signal categorization in this study. Depending on the original network structure, 1D convolutional neural network (CNN) and LSTM network were included as a parallel feature extracting layer for extracting the spatial and temporal attributes of ECG signal. Moreover, the improvised routing

technique was modelled, employing the prior probability of sub-capsules as a weighting factor for routing method categorization to weaken the effect of noise capsules.

In [17], dynamic and interoperable communication frameworks (DICFs) to regulate the functions of wearable medical gadgets were presented. The structure was accountable for monitoring, decision-making, and controlling the functionality and functioning period of the wearable sensor (WS) as a portion of smart healthcare trailing applications. In this structure, the nature of the wireless sensing gadgets and their basic attributes were accountable for devising a fully operative and continuous functioning of the sensing gadgets. Hoang et al. [18] devise a new technique utilizing a novel segmentation method and wide-ShuffleNet for classifying skin lesions. Firstly, the author computes the entropy-related weighting and first-order cumulative moment (EW-FCM) of the skin image. To separate the lesions from the background, such values were employed. After that, the author input the segmentation outcome into a novel DL framework wide-ShuffleNet and determined the type of skin lesion.

Mohapatra et al. [19] devise a smart healthcare system to detect numerous gastrointestinal (GI) abnormalities by utilizing time-frequency studies and CNNs. The initial stage of the work includes an image pre-processing stage, then extracting approximate DWT coefficients. Every class of decomposed images can be later given as input to certain concerned CNN techniques to train and test in 2 distinct classifier levels for recognizing its estimated value. In [20], a wearable sensor-related mechanism was modelled for activity estimation utilizing RNN on an edge device (i.e., laptop or personal computer). The input data of the mechanism can be gained from many wearable healthcare sensors like ECG, gyroscope sensors, magnetometers, and accelerometers. In [21], skin data will be accumulated through dedicated hardware. The gathered samples will be uploaded to the cloud for more processing utilizing a new multimodal data fusion structure, which executes skin lesion segmentation after classification. A hybrid structure was modelled, which uses the complementary strengths of 2 CNN structures for generating the segmented images for lesion segmentation.

3 The Proposed Model

This study designed a new HBO-DQNN method for decision-making in smart healthcare applications. The presented HBO-DQNN method primarily concentrates on detecting and classifying healthcare data. In the presented HBO-DQNN model, three stages of operations were performed, as elaborated below. Fig. 1 represents the block diagram of the HBO-DQNN system.

3.1 Data Normalization

Normalization has distinct meanings in statistics, the simple utility of which was to regularize variables or data. It can be a technique that presents data in a similar field when they are not [22]. Otherwise, a data miner can face circumstances where the data properties add values in various domains or ranges. Large-value features will have a higher impact on cost function than low-value features. In building a metamodel from data, earlier model training starts. Information was divided into equivalent values normalized to values among one and zero scales to minimize the effect of the total scale and have nearly every input in the same range. The min-max method was a popular and simple normalizing technique in medicinal imaging. Through the assumption of attribute X , such it has a mapping from the dataset among X_{\min} and X_{\max} , the min-max normalization (X_{norm}) is attained by the following:

$$X_{norm} = \frac{X - X_{\min}}{X_{\max} - X_{\min}}. \quad (1)$$

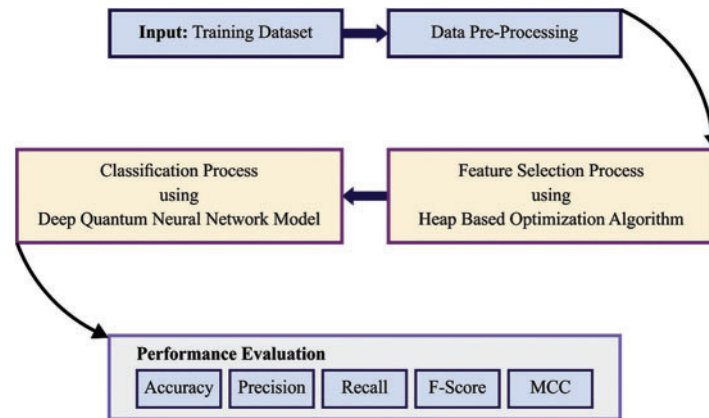


Figure 1: Block diagram of HBO-DQNN system

3.2 Feature Selection Using HBO Algorithm

In this stage, the HBO algorithm selects an optimal feature set from the healthcare data. Askari et al. [23] recommended a new MH called HBO based on the job description titles and employee responsibilities. The corporate rank hierarchy (CRH) was considered the common basis used in the corporation. The HBO depends on the 4 main phases; 1) employee self-contribution, 2) interaction between colleagues, 3) communication with the immediate boss, and 4) CRH. The mathematical modelling of HBO phases is listed as follows.

CRH: It can be modelled through the heap data structure. Here, searching agent fitness is defined using the index of the search agent, and key nodes in the population can be implemented by node value in a heap.

Communication with immediate boss: usually, the upper level in the central organizational structure is in charge of imposing restrictions and strategies; subordinates (children) will follow the immediate boss (parental node). To model these behaviours, the position of all search agents \vec{x}_i is upgraded following the parental node B as follows:

$$X_i^k(t+1) = B^k + \gamma \lambda^k |B^k - X_i^k(t)| \quad (2)$$

In Eq. (2), the existing iteration can be characterized as r , the k -th vector component denoted in the following. The parameter λ^k denotes the vector component. $\vec{\lambda}$ can be estimated as follows:

$$\vec{\lambda} = 2r - 1 \quad (3)$$

From the expression, the parameter γ can be defined as follows:

$$\gamma = \left| 2 - \frac{(t \bmod \frac{T}{c})}{\frac{T}{4c}} \right| \quad (4)$$

Now, C and T correspondingly symbolize variable and max iteration count. Usually, γ is linearly reduced from 2 to 0.

Communication among colleagues: In HBO, colleagues were agents, and the positions of all the agents are given below.

$$\chi_i^k(t+1) = \begin{cases} S_r^k + \gamma \lambda^k |S_r^k - X_i^k(t)|, f(\vec{S}_r) < f(\vec{x}_i(t)) \\ x_i^k + \gamma \lambda^k |S_r^k - x_i^k(t)|, f(\vec{S}_r) \geq f(\vec{x}_i(t)) \end{cases} \quad (5)$$

From the expression, the objective function (f) aims to estimate the fitness of the search agents. If $f(\vec{S}_r) < f(\vec{x}_i(t))$, Eq. (5) aim is to allow the search agent to search the region around (S_r^k) ; if not, around x_i

Employee's self-contribution: now, the self-contribution is executed by storing the preceding employee's location as;

$$x_i^k(t+1) = x_i^k(t) \quad (6)$$

A roulette wheel can be applied to separate the population into p_1 , p_2 , and p_3 proportions to preserve the balance between exploration and exploitation; p_1 enables the search agent to upgrade the location [24]. Moreover, p_1 , p_2 , and p_3 proportions are estimated, whereby t defines the present iteration, and T specifies the maximal iteration count.

$$p_1 = 1 - \frac{t}{T} \quad (7)$$

$$p_2 = p_1 + \frac{1 - p_1}{2} \quad (8)$$

$$p_3 = p_2 + \frac{1 - p_1}{2} = 1 \quad (9)$$

A common methodology to upgrade the searching agent location is as follows;

$$X_i^k(t+1) = \begin{cases} X_i^k(t), p \leq p_1 \\ B^k + \gamma \lambda^k |B^k - x_i^k(t)|, p > p_1 \text{ and } p \leq p_2 \\ S_r^k + \gamma \lambda^k |S_r^k - x_i^k(t)|, \\ p > p_2 \text{ and } p \leq p_3 \text{ and } f(\vec{S}_r) < f(\vec{x}_i(t)) \\ X_i^k + \gamma \lambda^k |S_r^k - X_i^k(t)|, \\ p > p_2 \text{ and } p \leq p_3 \text{ and } f(\vec{S}) \geq f(\vec{x}_i(t)) \end{cases} \quad (10)$$

Now, p in $[0, 1]$ designates an arbitrary integer. Note that Eq. (2) boosts exploitation and convergence, Eq. (6) raises exploration, and Eq. (5) endorses exploitation and exploration. The HBO system's fitness function (FF) will consider the classifier accuracy and the number of selected features. It maximizes classifier accuracy and diminishes the set size of the selected features. Therefore, the following FF was employed for assessing individual solutions, as established in Eq. (11).

$$Fitness = \alpha * ErrorRate + (1 - \alpha) * \frac{\#SF}{\#All_F} \quad (11)$$

Whereas error rate refers to the classification error rate employing particular features. The error rate has been calculated as the percentage of incorrect classifiers (by DQNN technique) to the count of classifiers made, written as a value betwixt zero and one. (ErrorRate was the complement of classifier

accuracy), $\#SF$ stands for the count of selective features, and $\#All_F$ stands for the elements in the novel database. α is employed for controlling the significance of classifier qualities and subset lengths. During this experiment, α is set to 0.9.

3.3 Healthcare Data Classification

Finally, the DQNN model is exploited for healthcare data classification. The quantum network comprises many quantum perceptrons [25]. Every perceptron parallels the arbitrary unitary operator having n output rubies and m input qubits, resulting in $(2^{m+n})^2 - 1$ complex parameter. This operator is significant for the neural network weight matrix. Density matrices denote the state. In these cases, the output and input matrices are pure states that imply:

$$\rho^{in} = |\psi^{in}\rangle\langle\psi^{in}|, \rho^{out} = |\psi^{out}\rangle\langle\psi^{out}| \quad (12)$$

The quantum network comprises the input layer (input density matrix), L hidden layer (the unitary operator), and an output layer ($L + 1 \equiv out$). The deep quantum network can be described by

$$\rho^{out} = tr_{l \neq out} [U (\rho^{in} \otimes |0_{out}, 0_L, \dots, 0_1\rangle\langle 0_{in}, 0_1, \dots, 0_L, 0_{out}|) U^T] \quad (13)$$

Whereas the trace is taken over each layer except the output layer, $|0, \dots, 0\rangle$ denotes the ground state, and U represents the resultant quantum circuit or unitary transformation.

$$U = U^{out} U^{(L)} U^{(L-1)} \dots U^{(1)} \quad (14)$$

$\{U^{(l)}\}_{l=1}^L$ shows unitary operators ($U^{(l)} U^{(l)\dagger} = U^{(l)\dagger} U^{(l)} = I, \forall l$). Fig. 2 represents the framework of the QNN technique [26].

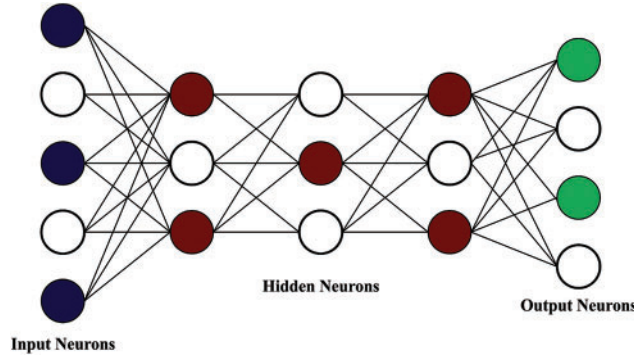


Figure 2: Architecture of QNN

The symbol \dagger characterizes Hermitian transpose (transpose of complex conjugate). The product of the non-commuting unitary operator factorizes every unitary operator:

$$U^{(l)} = \prod_j^{m_l} U_j^{(l)} \quad (15)$$

Consequently, the net positive is formulated by the composition of 0 & a series of completely positive transition maps.

$$\rho^{out} = \mathcal{E}^{out} (\mathcal{E}^{(L)} (\dots \mathcal{E}^{(2)} (\mathcal{E}^{(1)} (\rho^{in})) \dots)) \quad (16)$$

Whereas the transition map over layer l can be equivalent to

$$\mathcal{E}^{(l)}(X^{(l-1)}) = \underset{\wedge}{tr}_{l-1} \left[\prod_{j=m_l}^1 U_j^{(l)} (X^{(l-1)} \otimes |0_l, \dots, 0_1, 0_{in}\rangle \langle 0_{in}, 0_1, \dots, 0_L, 0_{out}|) \prod_j^{m_l} U_j^{(l)\dagger} \right] \quad (17)$$

And the trace being taken over layer $l - 1$. The architecture of Eq. (16) was reminiscent of the dense deep NN, where all the transition maps are integrated into the activation function. The training cost function depends on fidelity which was a fundamentally sole measure for pure states, estimating the nearness among the true output $|\psi_t^{out}\rangle$ and network output ρ_t^{out} ,

$$\mathcal{L} = \frac{1}{t_f - t_i + 1} \sum_{t=t_i}^{t_f} \langle \psi_t^{out} | \rho_t^{out} | \psi_t^{out} \rangle, \quad L \in [0, 1] \quad (18)$$

Here, t denotes a specific sample of the trained set (time later at time t). The cost function differs between zero and one, with the latter happening while the quantum network outputs are equivalent to the true outputs. In the training stage, the unitary operator is upgraded as follows:

$$U_j^{(l)} \rightarrow \exp [i\varepsilon K_j^{(l)}] U_j^{(l)}, \quad i \equiv \sqrt{-1} \quad (19)$$

Afterwards the training iteration, the cost function differs as

$$\Delta C = \frac{\varepsilon}{t_f - t_i + 1} \sum_{t=t_i}^{t_f} \sum_{l=1}^{L+1} tr [\sigma_t^{(l)} \Delta \mathcal{E}^{(l)} (\rho_t^{(l-1)})] \quad (20)$$

Whereas

$$\rho_t^{(l)} = \mathcal{E}^{(l)} (\dots \mathcal{E}^{(2)} (\mathcal{E}^{(1)} (\rho_t^{in})) \dots) \quad (21)$$

is the density matrix for layer l ,

$$\sigma_t^l \triangleq F^{(l+1)} (\dots F^{(L)} (F^{out} (|\psi_t^{out}\rangle \langle \psi_t^{out}| \dots))) \quad (22)$$

And (P) indicates the adjoint channel of the wholly positive map (\mathcal{E}):

$$F^{(l)}(X^{(l-1)}) = tr_{l-1} \left[\prod_{j=m_l}^1 U_j^{(l)\dagger} (|0_{out}, 0_L, \dots, 0_1, 0_{in}\rangle \langle 0_{in}, 0_1, \dots, 0_l| \otimes X^{(l-1)}) \prod_j^{m_l} U_j^{(l)} \right] \quad (23)$$

The training model is defined. Firstly, a first unitary operator is randomly selected. Then, the density matrices are recursively evaluated for each layer and training pair. Then, the variable matrixes $K_j^{(l)}$ is evaluated, with the trace taken on each qubit unaffected using $U_j^{(l)}$ and η , being the learning rate. Lastly, the unitary operator was upgraded. Steps 2 and 3 were repetitive until the cost function obtained a maximal (preferable one). Consequently, the variable matrixes might be estimated. The two layers are necessary, significantly decreasing the memory needed. Accordingly, the dimension of the matrix scale has several layers, thereby ensuring that the deep quantum neural network is computationally tractable.

$$U_j^{(l)}, \quad j \sim p(j), \quad \sim p(l), \quad (24)$$

$$\forall (|\psi_t^{in}\rangle, |\psi_t^{out}\rangle) \in \{(|\psi_t^{in}\rangle, |\psi_t^{out}\rangle)\}_{t=t_i}^{t_f} \wedge \forall [l] \Rightarrow \rho_t^{(l)} = tr_{l-1} \mathcal{E}^l (\rho_t^{(l-1)}) \\ = tr_{l-1} \left[\prod_{j=m_l}^1 U_j^{(l)} (\rho_t^{(l-1)} \otimes |0_{out}, 0_L, \dots, 0_1\rangle \langle 0_{in}, 0_1, \dots, 0_L, 0_{out}|) \prod_j^{m_l} U_j^{(l)\dagger} \right], \quad (25)$$

$$U_j^{(l)} \rightarrow \exp [i\varepsilon K_j^{(l)}] U_j^{(l)}, \quad (26)$$

$$K_j^{(l)} = \eta \frac{2^{m_{l-1}}}{t_f - t_i + 1} \sum_{t=t_i}^{t_f} tr_{C U_j^{(l)}} M_j^{(l)}, \tag{27}$$

where

$$M_j^{(l)} = \left[\prod_{\alpha=j}^1 U_{\alpha}^{(l)} (\rho_t^{(l-1:l)}) \prod_{\alpha=1}^j U_{\alpha}^{(l)\dagger} \times \prod_{\alpha=j+1}^{m_l} U_{\alpha}^{(l)\dagger} (I_{l-1} \otimes \sigma_t^{(l)}) \prod_{\alpha=m_l}^{j+1} U_{\alpha}^{(l)} \right] \tag{28}$$

$$\rho_t^{(l-1:l)} \hat{=} \rho_t^{(l-1)} \otimes |0_{out}, 0_L, \dots, 0_1\rangle \langle 0_{in}, 0_1, \dots, 0_L, 0_{out}|, \tag{29}$$

$$\sigma_t = F^{(l+1)} (\dots F^{out} (|\psi_t^{out}(l)\rangle \langle \psi_t^{out}(l)|) \dots) \tag{30}$$

4 Results and Discussion

This section examines the performance of the HBO-DQNN method on two datasets, namely colon cancer and Lymphoma datasets. Table 1 represents the details of two datasets.

Table 1: Dataset details

Class	No. of samples	Total no. of samples
Colon cancer (Genes = 2000)		62
Tumor	40	
Normal	22	
Lymphoma (Genes = 4026)		45
Germinal centre B-Like (GCL)	23	
Activated B-Like (ACL)	22	

Table 2 and Fig. 3 illustrate the overall classification outcomes of the HBO-DQNN technique on the Colon Cancer Dataset. The outcomes inferred that the HBO-DQNN method had reached effectual outcomes under all aspects. For example, on 80% of the TR database, the HBO-DQNN model has obtained an average $accu_y, prec_n, reca_l, F_{-score}$, and MCC of 96.61%, 95.76%, 95.92%, 96.31%, and 95.84% respectively. Meanwhile, on 20% of the TR database, the HBO-DQNN approach has reached an average $accu_y, prec_n, reca_l, F_{-score}$, and MCC of 97.05%, 95.43%, 96.05%, 96.43%, and 95.27% correspondingly. Concurrently, on 70% of TR databases, the HBO-DQNN approach has achieved an average $accu_y, prec_n, reca_l, F_{-score}$, and MCC of 96.51%, 95.73%, 97.22%, 96.68%, and 97.44% correspondingly. At last, on 30% of the TR database, the HBO-DQNN system has gained an average $accu_y, prec_n, reca_l, F_{-score}$, and MCC of 96.64%, 95.98%, 96.86%, 96.91%, and 96.63% correspondingly.

Table 2: Result analysis of HBO-DQNN system with distinct measures under colon cancer dataset

Class	Accuracy	Precision	Recall	F-score	MCC
Training phase (80%)					
Tumor	95.48	95.89	95.64	95.99	95.56
Normal	97.74	95.62	96.19	96.62	96.11
Average	96.61	95.76	95.92	96.31	95.84
Testing phase (20%)					

(Continued)

Table 2: Continued

Class	Accuracy	Precision	Recall	F-score	MCC
Tumor	96.86	95.04	95.45	96.36	95.31
Normal	97.24	95.81	96.65	96.50	95.23
Average	97.05	95.43	96.05	96.43	95.27
Training phase (70%)					
Tumor	96.05	95.13	97.72	96.21	97.25
Normal	96.97	96.32	96.71	97.15	97.62
Average	96.51	95.73	97.22	96.68	97.44
Testing phase (30%)					
Tumor	95.45	96.71	95.98	97.61	97.84
Normal	97.82	95.25	97.74	96.21	95.42
Average	96.64	95.98	96.86	96.91	96.63

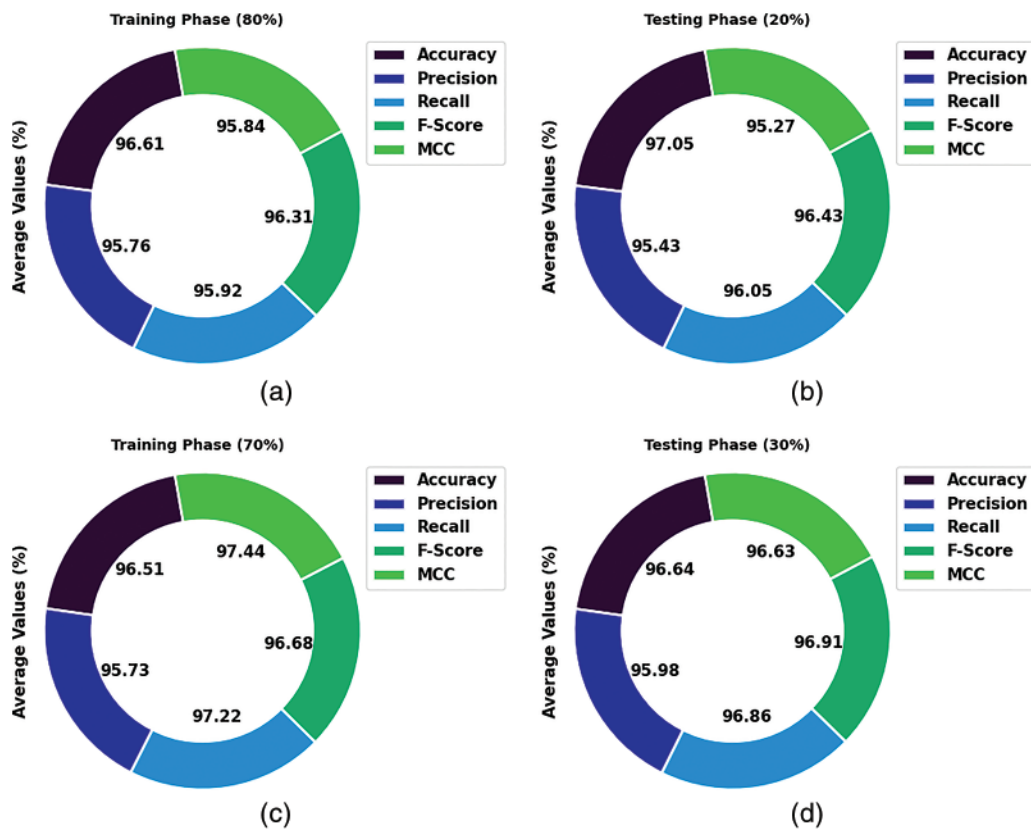


Figure 3: Average analysis of HBO-DQNN system under colon cancer dataset (a and b) TR and TS database of 80:20 and (c and d) TR and TS database of 70:30

The training accuracy (TR_{acc}) and validation accuracy (VL_{acc}) obtained by the HBO-DQNN system under the Colon cancer database is exposed in Fig. 4. The simulation result stated that the

HBO-DQNN approach had attained increased values of TR_{acc} and VL_{acc} . In certain, the VL_{acc} looked that better than TR_{acc} .

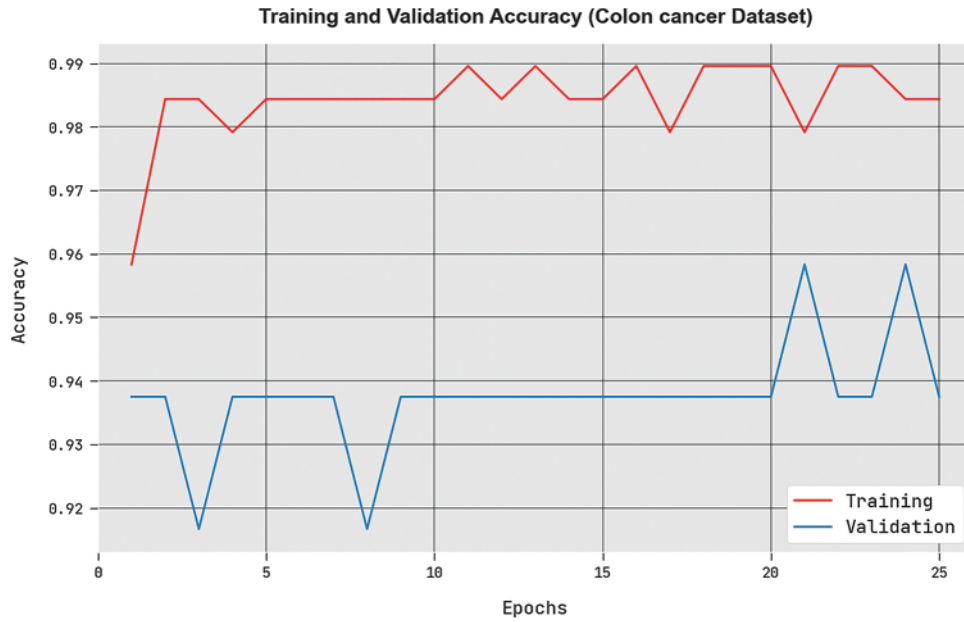


Figure 4: TR_{acc} and VL_{acc} analysis of HBO-DQNN system under colon cancer dataset

The training loss (TR_{loss}) and validation loss (VL_{loss}) acquired by the HBO-DQNN approach under the Colon cancer database are shown in Fig. 5. The simulation values represented by the HBO-DQNN method have exhibited minimal values of TR_{loss} and VL_{loss} . Particularly, the VL_{loss} is lesser than TR_{loss} .

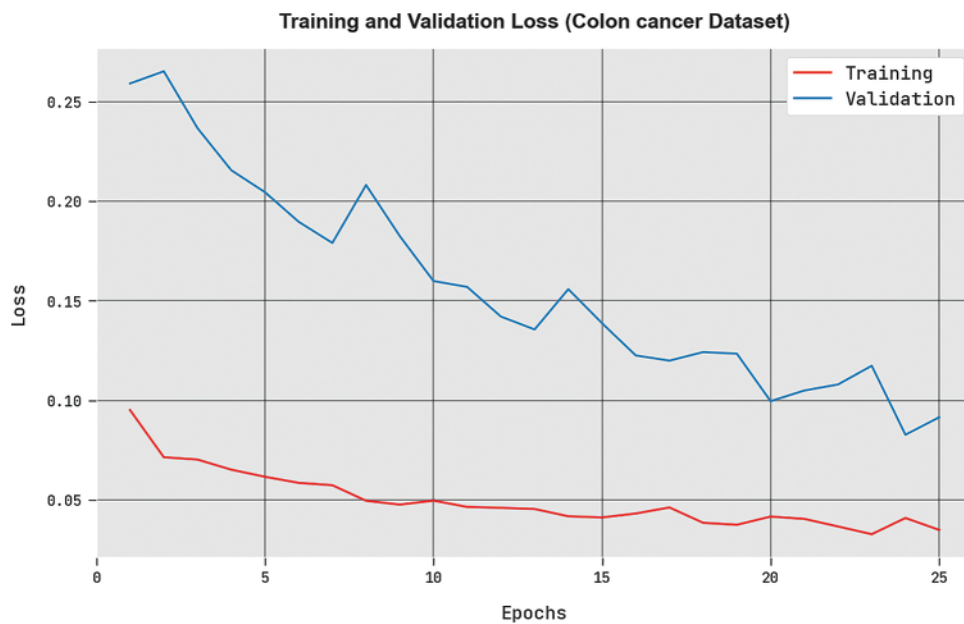


Figure 5: TR_{loss} and VL_{loss} analysis of HBO-DQNN system under colon cancer dataset

A detailed ROC analysis of the HBO-DQNN technique under the Colon cancer database is displayed in Fig. 6. The outcomes displayed the HBO-DQNN method’s capability in classifying different classes in the test database.

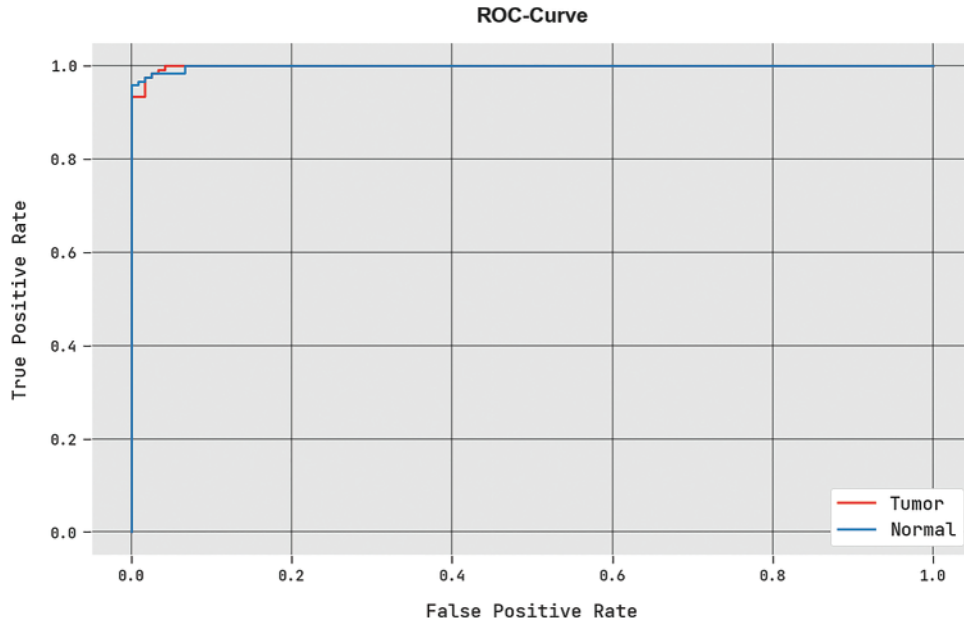


Figure 6: ROC analysis of HBO-DQNN system under colon cancer dataset

Table 3 and Fig. 7 demonstrate the overall classification outcomes of the HBO-DQNN method on the Lymphoma Dataset. The results show the HBO-DQNN technique has reached effectual outcomes in all aspects. For instance, on 80% of the TR database, the HBO-DQNN method has gained an average $accu_y$, $prec_n$, $reca_t$, F_{score} , and MCC of 94.79%, 94.59%, 94.98%, 94.78%, and 95.09% correspondingly.

Table 3: Result analysis of HBO-DQNN system with distinct measures under the lymphoma dataset

Class	Accuracy	Precision	Recall	F-score	MCC
Training phase (80%)					
Tumor	95.12	94.66	94.11	95.30	95.34
Normal	94.46	94.51	95.85	94.25	94.84
Average	94.79	94.59	94.98	94.78	95.09
Testing phase (20%)					
Tumor	94.79	94.61	95.72	94.08	94.83
Normal	94.39	94.34	94.96	94.79	94.24
Average	94.59	94.48	95.34	94.44	94.54
Training phase (70%)					
Tumor	95.35	94.81	95.21	95.94	95.47
Normal	95.22	95.32	95.46	95.32	95.31
Average	95.29	95.07	95.34	95.63	95.39

(Continued)

Table 3: Continued

Class	Accuracy	Precision	Recall	F-score	MCC
Testing phase (30%)					
Tumor	95.69	95.85	94.45	94.51	96.00
Normal	95.74	94.98	95.48	94.88	94.79
Average	95.72	95.42	94.97	94.70	95.40

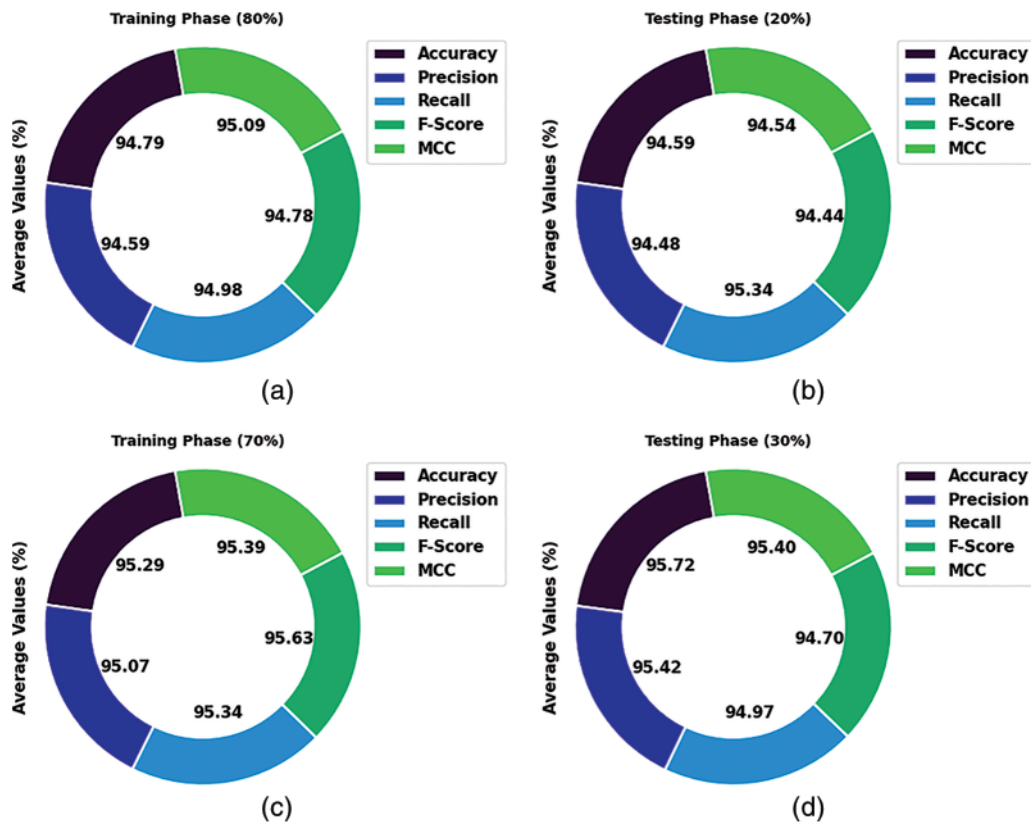


Figure 7: Average analysis of HBO-DQNN system under lymphoma dataset (a and b) TR and TS database of 80:20 and (c and d) TR and TS database of 70:30

In the meantime, on 20% of the TR database, the HBO-DQNN technique has achieved an average $accu_y$, $prec_n$, $reca_t$, F_{-score} , and MCC of 94.59%, 94.48%, 95.34%, 94.44%, and 94.54% correspondingly. Simultaneously, on 70% of the TR database, the HBO-DQNN method has attained an average $accu_y$, $prec_n$, $reca_t$, F_{-score} , and MCC of 95.29%, 95.07%, 95.34%, 95.63%, and 95.39% correspondingly. At last, on 30% of the TR database, the HBO-DQNN approach has attained an average $accu_y$, $prec_n$, $reca_t$, F_{-score} , and MCC of 95.72%, 95.42%, 94.97%, 94.70%, and 95.40% individually.

The TL_{acc} and VL_{acc} gained by the HBO-DQNN approach under the Lymphoma database are displayed in Fig. 8. The simulation values exhibited by the HBO-DQNN method have reached the maximal values of TR_{acc} and VL_{acc} . Mostly the VL_{acc} is greater than TR_{acc} .

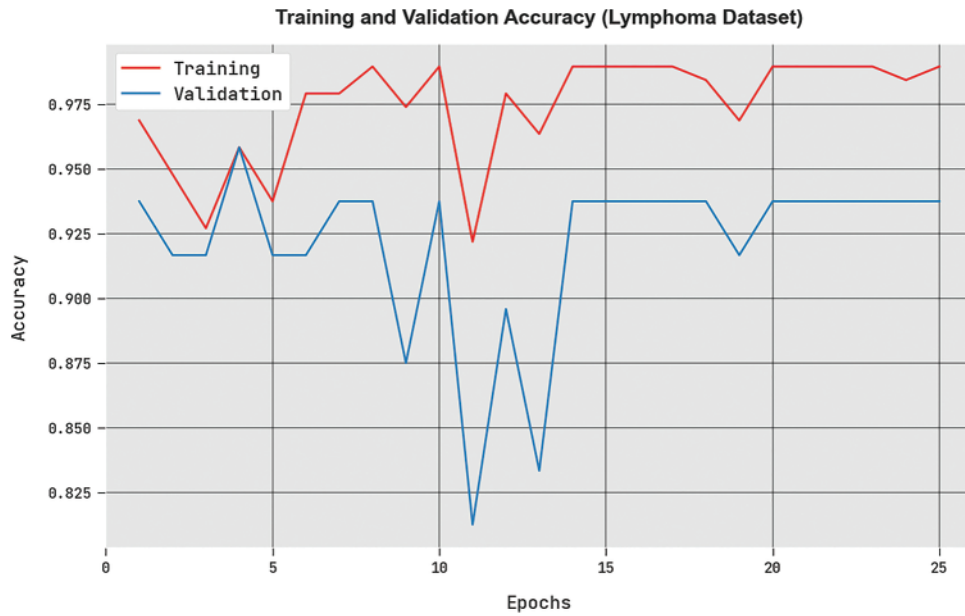


Figure 8: TR_{acc} and VL_{acc} analysis of HBO-DQNN system under the lymphoma dataset

The TR_{loss} and VL_{loss} attained by the HBO-DQNN method under the Lymphoma database are displayed in Fig. 9. The simulation values exhibited by the HBO-DQNN method have manifested the least values of TR_{loss} and VL_{loss} . Particularly, the VL_{loss} is lesser than TR_{loss} .

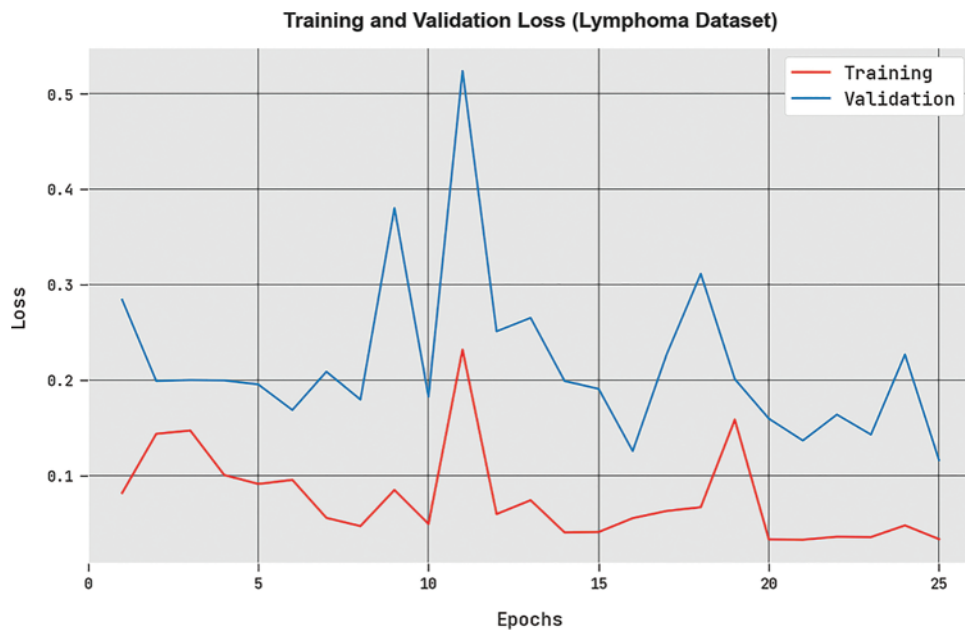


Figure 9: TR_{loss} and VL_{loss} analysis of HBO-DQNN system under the lymphoma dataset

A brief ROC study of the HBO-DQNN approach under the Lymphoma database is portrayed in Fig. 10. The outcomes denoted the HBO-DQNN technique has displayed its capability in classifying different classes in the test database.

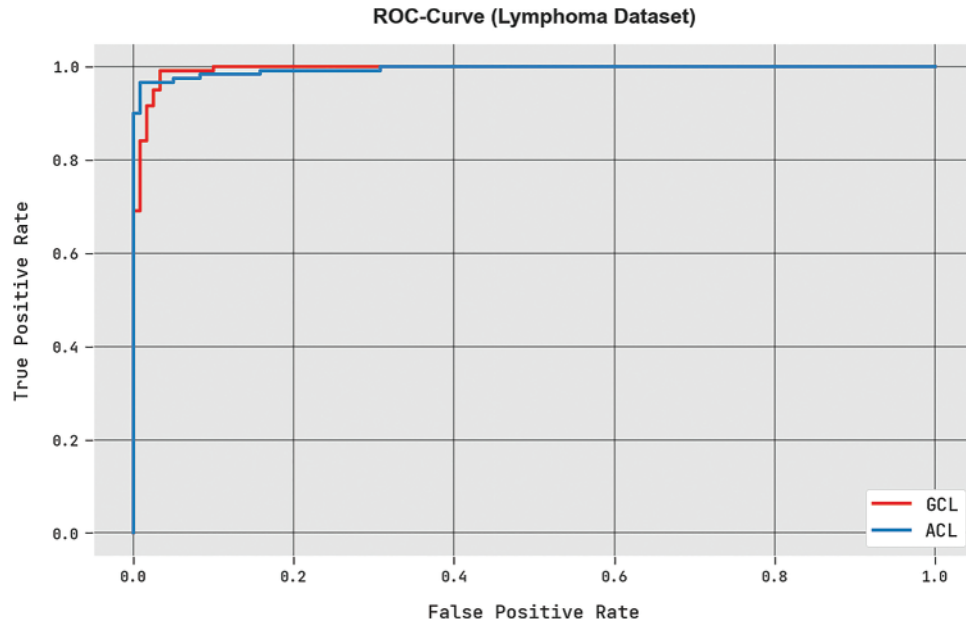


Figure 10: ROC analysis of HBO-DQNN system under the lymphoma dataset

A comprehensive comparison study of the HBO-DQNN method with contemporary techniques is given in Table 4 [27]. Fig. 11 examines a close $accu_y$ inspection of the HBO-DQNN method with other ML methods on the colon cancer dataset. The simulation values implied that the KNN, SVM-RBF, SVM-Quadratic, and SVM-Polynomial models had reduced $accu_y$ of 95.87%, 95.46%, 95%, and 95.19%, respectively. Although the ANN and SVM-linear models have resulted in closer $accu_y$ values of 96.46% and 96.25%, the presented HBO-DQNN model has shown a maximum $accu_y$ of 97.05%.

Table 4: Accuracy analysis of HBO-DQNN system with existing approaches under two database

Methods	Accuracy (%)	
	Colon cancer	Lymphoma
HBO-DQNN	97.05	95.72
KNN model	95.87	93.96
ANN model	96.46	93.59
SVM-linear	96.25	94.52
SVM-RBF	95.46	93.53
SVM-quadratic	95.00	94.18
SVM-polynomial	95.19	93.91

Finally, Fig. 12 examines a comparative $accu_y$ analysis of the HBO-DQNN technique with other ML models on the lymphoma dataset. The simulation values exhibited the KNN, SVM-RBF, ANN,

and SVM-Polynomial models have shown reduced $accu_y$ of 93.96%, 93.53%, 93.59%, and 93.91%, respectively. Though the SVM-linear and SVM-Quadratic methods have resulted in closer $accu_y$ values of 94.52% and 94.18%, the presented HBO-DQNN approach has shown a maximum $accu_y$ of 95.72%.

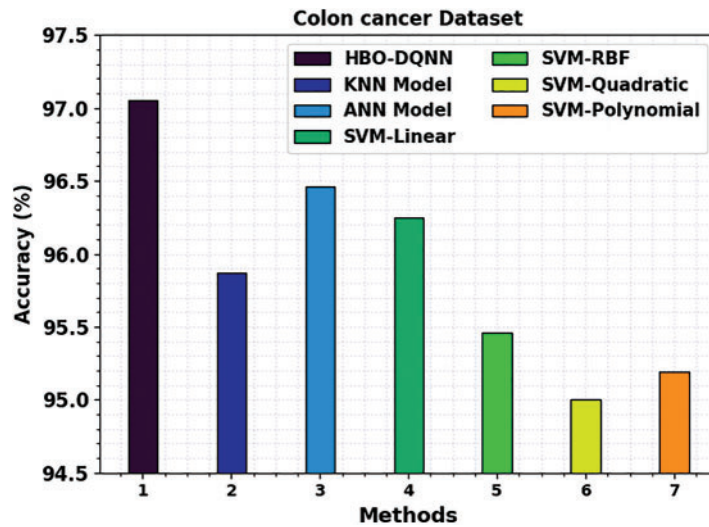


Figure 11: $Accu_y$ analysis of HBO-DQNN system under colon cancer dataset

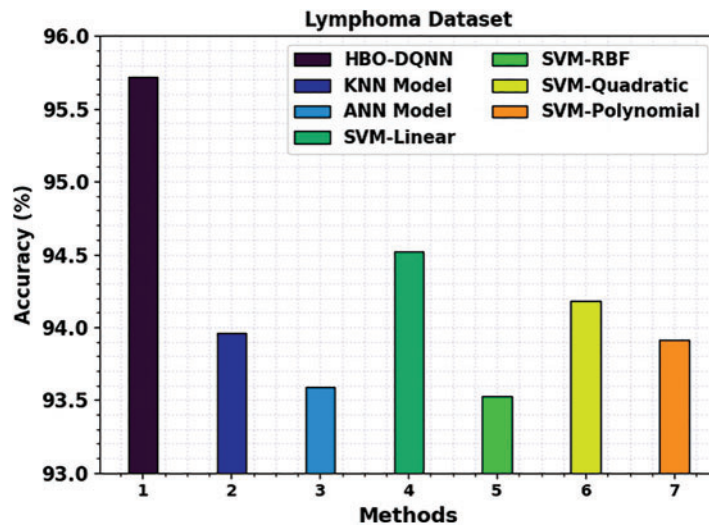


Figure 12: $Accu_y$ analysis of HBO-DQNN system under the lymphoma dataset

5 Conclusion

In this study, a new HBO-DQNN method was designed for decision-making in smart healthcare applications. The presented HBO-DQNN method primarily concentrates on detecting and classifying healthcare data. In the presented HBO-DQNN model, three stages of operations were performed. Data normalization is applied to pre-process the input data at the initial stage. Next, the HBO algorithm is used in the second stage to choose an optimal set of features from the healthcare data. At

last, the DQNN model was used for healthcare data classification. A series of experiments were carried out to depict the promising classifier results of the HBO-DQNN model. The extensive comparative analysis reported the improvements of the HBO-DQNN method over other existing models with maximum accuracy of 97.05% and 95.72% under the colon cancer and lymphoma dataset. In future, hybrid DL classifiers can be involved to boost overall performance.

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