

Optimizing the learning process of multi-layer perceptrons using a hybrid algorithm based on MVO and SA

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ABSTRACT

Artificial neural networks (ANNs) are one of the artificial intelligence techniques used in real-world problems and applications encountered in almost all industries such as education, health, chemistry, food, informatics, logistics, transportation. ANN is widely used in many techniques such as optimization, modelling, classification and forecasting, and many empirical studies have been carried out in areas such as planning, inventory management, maintenance, quality control, econometrics, supply chain management and logistics related to ANN. The most important and just as hard stage of ANNs is the learning process. This process is about finding optimal values in the search space for different datasets. In this process, the values generated by training algorithms are used as network parameters and are directly effective in the success of the neural network (NN). In classical training techniques, problems such as local optimum and slow convergence are encountered. Meta-heuristic algorithms for the training of ANNs in the face of this negative situation have been used in many studies as an alternative. In this study, a new hybrid algorithm namely MVOSANN is suggested for the training of ANNs, using Simulated annealing (SA) and Multi-verse optimizer (MVO) algorithms. The suggested MVOSANN algorithm has been experimented on 12 prevalently classification datasets. The productivity of MVOSANN has been compared with 12 well-recognized and current meta-heuristic algorithms. Experimental results show that MVOSANN produces very successful and competitive results.

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1. Introduction

In recent years, meta-heuristic algorithms have been frequently used in computational methods to increase efficiency and quality, reduce costs, and solve various complex problems encountered during the development and management of business processes by using production resources at an optimum level. Although the best solution cannot always be found with meta-heuristic algorithms, they are preferred by researchers because they have features such as ease of application, their effectiveness in complex problems, adaptability to different problems, and so on (Kaya & Fıglalı 2018; Talbi, 2009). The common goal in such algorithms is to get the best solution in the existing circumstances for difficult and complex problems. To obtain the best result, the exploration and exploitation features of the algorithms should be strong, and the balance should be struck between these two processes.

One way to equilibrate between exploration and exploitation may be to combine algorithms (Mirjalili & Hashim, 2010; Mafarja & Mirjalili, 2017). A hybrid algorithm is to combine one algorithm with another algorithm or algorithms using the preferred features of the algorithms. In many studies, it is seen that methods such as improvement, modification and hybridization have been tried to increase the success of algorithms. Hybrid algorithms can be essentially classified into two

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groups. The first group is integrative hybrids, which are constituted by integrating a subsidiary algorithm into a primary algorithm. The second group is collaborative hybrids, a combination of two or more algorithms that run sequentially or in parallel (Ting et al., 2015). Many studies have been carried out on functions such as function optimization and real-world problems with hybrid algorithms.

The process of training for artificial neural networks (ANNs) is an important implementation of meta-heuristic algorithms and hybrid versions of these algorithms. ANNs, generally defined as a mathematical model of nerve cells in the human brain and the connection between these cells, is one of the most researched artificial intelligence techniques today (McCulloch & Pitts, 1943; Anderson, 1995). ANNs are widely used in many fields such as classification, prediction, and optimization. One of the different neural networks (NNs) structures suggested in the literature is the feed-forward multi-layer perceptron (MLP). MLP has multi-layer neuron architecture. At this structure, the first layer is called the input layer, the last layer is the output layer, and the other layers between the first layer and the last layer are named as hidden layers. Neurons in all layers from the first to the last layer are connected to neurons in the following layer. The information flow is forward and there is no feedback. One of the reasons why this technique is preferred by researchers is its suitability for solving nonlinear and complex problems (Faris et al., 2016).

One of the situations that significantly affect performance in MLP and other ANNs structures is the training process. In this process, the output value generated at the end of each iteration is compared with the target value. Depending on the error, the biases and weights are updated. These operations to minimize the error are known as training the network. Supervised, unsupervised and reinforced learning methods are used during the education process. In MLP training, gradient-based and stochastic methods are used for the supervised learning method. Gradient-based conventional methods for instance the back propagation algorithm (Rumelhart et al., 1986) have disadvantages such as local minima, slow convergence, and dependence on initial values (Gori & Tesi, 1992; Gupta & Sexton, 1999). Stochastic methods, such as meta-heuristic algorithms, are other methods recommended as an alternative in the training process to overcome the disadvantages of conventional methods. GA (Seiffert, 2001), ABC (Karaboga et al., 2007), MOA (Mirjalili & Sadiq, 2011), AFS (Hasan et al., 2011), PSO/GSA (Mirjalili et al., 2012), FA (Brajevic & Tuba, 2013; Alweshah, 2014), GWO (Mirjalili et al., 2014a), SSO (Mirjalili et al., 2015), BBO (Mirjalili et al., 2014b), MFO (Yamany et al., 2015), CSSO (Abedinia & Amjady, 2015), MBA (Tuba et al., 2015), MVO (Faris et al., 2016), MPSO (Kolay et al., 2016), SOS (Wu et al., 2016), GSO (Alboaneen et al., 2017), WOA (Aljarah et al., 2018), IMBO (Faris et al., 2018), SSA (Abusnaina et al., 2018; Bairathi & Gopalani, 2019) algorithms can be given as examples of meta-heuristic and hybrid algorithms used in the training of ANNs.

In this study, MVO which is a population-based algorithm and SA which is a single-solution based algorithm used for the training of the MLP, in an integrative hybrid structure. Our aim is to increase the success of MLP training with this hybrid model. Lately, a new hybrid model has been suggested for function optimization, in which MVO and SA algorithms are used together (Yilmaz et al., 2022). However, regarding to the no free lunch (NFL) theorem, a single optimization technique is not sufficient to solve all optimization problems (Wolpert & Macready, 1997). MLP training also means a different optimization problem for each different data set. For this reason, the proposed algorithm for function optimization has been modified and adapted for MLP training. In the literature, a hybrid model that uses MVO and SA algorithms for the training of ANNs not found and this model will be implemented for the first time.

The remaining of this article is established as follows: in the second section, a general description of MLP - NN is given. In the third section elaborate information regarding the suggested hybrid algorithm (MVOSANN) and its components is explained. In the fourth section, it is shown how to use the MVOSANN algorithm to train MLP. Comparing experiment results with other algorithms are given in the fifth section. In this section, also, the performance of the suggested algorithm is interpreted. Finally, in the sixth section, the results of the implementation were evaluated, and recommendations were made for future studies.

2. Feed-forward multilayer perceptron

ANNs are mathematical models whose names and structures are taken inspiration by the human brain. They mimic the nerve cells in the human brain and the connections between these cells. ANNs can be classified into different types to be used for various targets. Feed-forward MLP is one of the commonly used types of ANNs. The input layer, the hidden layers, and the output layer are the parts that constitute the MLP. Neurons are interconnected in one direction and forward. A simple example of a feed-forward MLP architecture with a single hidden layer is shown in Fig.1. Each neuron in one layer connects to all neurons in other layers. Each neuron has a different weight and bias values. If the output of any neuron is above the specified threshold, that node is activated, and data is sent to the next layer. Otherwise, no data is sent to the next layer. The calculation of the weighted sum of the input values is shown below:

$$v_j = \sum_{i=1}^n w_{ij} I_i + \beta_j \quad (1)$$

where, v_j is the net input value of neuron j in the hidden layer, w_{ij} is the connection weight connecting I_i to neuron j , I_i indicates the neuron i in the input layer, β_j is a bias weight of neuron j , and n is the total number of neuron inputs.

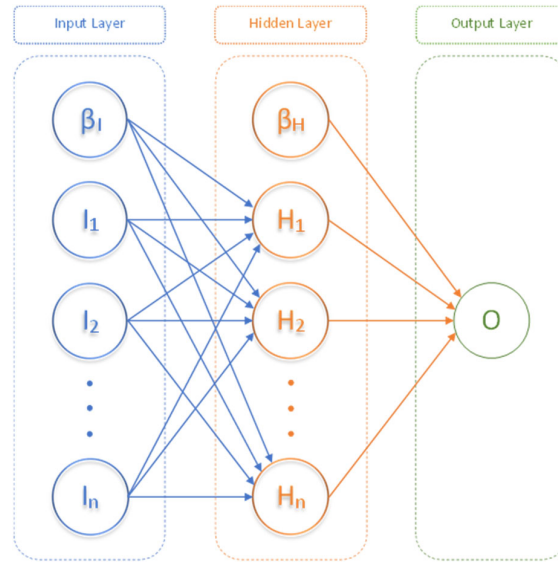


Fig. 1. Multilayer perceptron network with single hidden layer

The outputs of neurons are activated with an activation function which is dependent on the value of the summation function. In general, this function is a nonlinear function like hyperbolic tangent sigmoid. The hyperbolic tangent sigmoid function is shown as follows:

$$f(n) = \frac{2}{1 + e^{-2n}} - 1 \tag{2}$$

In the hidden layer, the output of each neuron is calculated as follows:

$$y_j = f_j \left(\sum_{i=1}^n w_{ij} I_i + \beta_j \right) \tag{3}$$

After the NN is created, the input data presented to it is expected to give the desired output. For the NN to achieve this, it needs to update the network parameters, that is, the weights. This stage is called the training process, which is a tough stage that shows the power of the network. The goal here is to minimize the error among the actual values and the calculated output values. This process is performed using a training algorithm. The system generates an output value by using the values generated at the end of the training and the inputs presented to it.

3. MVOSANN algorithm

3.1. Multi-verse optimizer (MVO)

In 2016, population-based multiverse optimization, taking inspiration from the concepts of wormholes, black holes, and white holes in the big bang theory and in the multiverse theory, was suggested by Seyedali Mirjalili et al. (Mirjalili et al., 2016). In this algorithm, where wormholes, black holes, and white holes are mathematically modelled for exploration, exploitation, and local search, the fitness function for each universe is referred by an inflation rate. Each universe represents a candidate solution, and each object represents a variable in the candidate solution.

In the algorithm process, sometimes, there is an exchange of objects between universes. Universes with high inflation rate prone to send their objects to universes with low inflation rate, while universes with low inflation rate prone to receive objects from universes with high inflation rate. Optimization is started with the above steps. In each new iteration, the universes are ordered accordingly their inflation rates at the end of the previous iteration. The conceptional model of the suggested algorithm is shown in Fig. 2. The following rules apply in optimization:

- The odds that universes have white holes increase if inflation rates are higher.
- The odds that universes have black holes increase if inflation rates are lower.
- Universes with high inflation rate prone to dispatch objects via white holes.
- Universes with low inflation rate prone to get more objects from black holes.
- Wormholes can cause random object movement from the best universe to other universes.

The mathematical model of MVO is as follows:

$$UNV = \begin{bmatrix} o_1^1 & o_1^2 & \dots & o_1^k \\ o_2^1 & o_2^2 & \dots & o_2^k \\ \vdots & \vdots & \ddots & \vdots \\ o_n^1 & o_n^2 & \dots & o_n^k \end{bmatrix} \tag{4}$$

where, n is the number of universes (candidate solutions), k is the number of parameters (variables).

$$o_i^j = \begin{cases} o_h^j & rnl < NRM(UNV_i) \\ o_i^j & rnl \geq NRM(UNV_i) \end{cases} \tag{5}$$

where, o_i^j defines the j^{th} parameter of i^{th} universe, o_h^j defines the j^{th} parameter of h^{th} universe selected by a roulette wheel selection, rnl is a number generated randomly in the range $[0, 1]$, $NRM(UNV_i)$ is normalized inflation rate of the i^{th} universe, UNV_i defines the i^{th} universe.

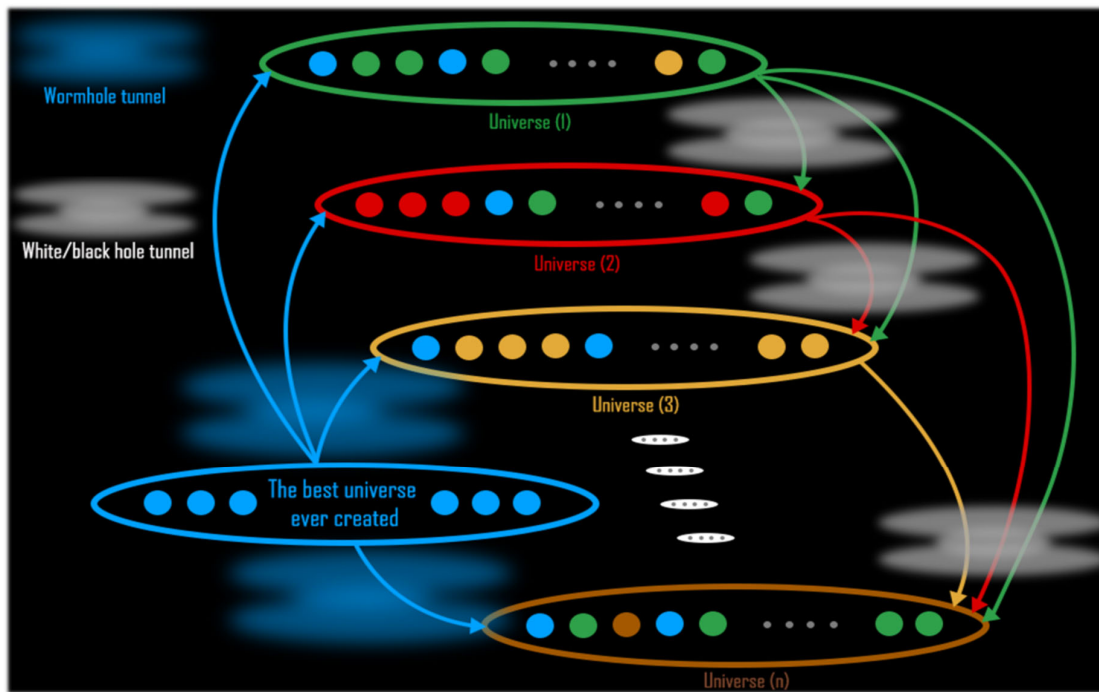


Fig. 2. Conceptual Model of MVO Algorithm.

Wormholes, sometimes randomly replace objects of universes with objects of the best universe to preserve and exploit the diversity of universes. The mathematical model of this mechanism is as follows.

$$o_i^j = \begin{cases} O_j + Travelling_DR \times ((uppb_j - lowb_j) \times rn4 + lowb_j) & rn3 < 0.5 \\ O_j - Travelling_DR \times ((uppb_j - lowb_j) \times rn4 + lowb_j) & rn3 \geq 0.5 \end{cases} \quad \begin{matrix} rn2 < Wormhole_EP \\ rn2 \geq Wormhole_EP \end{matrix} \quad (6)$$

where, O_j defines the j^{th} parameter of best universe generated up to now, o_i^j defines the j^{th} parameter of i^{th} universe, $uppb_j$ is the upper bound of j^{th} variable, $lowb_j$ is the lower bound of j^{th} variable, $Travelling_DR$ and $Wormhole_EP$ are the coefficients, and $rn2, rn3, rn4$ are numbers acquired randomly in the range $[0, 1]$.

MVO has two coefficients: Travelling distance rate ($Travelling_DR$) and Wormhole existence probability ($Wormhole_EP$) (Eq. 6). The mathematical model for both coefficients is as follows:

$$Wormhole_EP = min + l \times \left(\frac{max - min}{L} \right) \quad (7)$$

where, l is the current iteration, L is the maximum iteration, min is the minimum value equal to 0.2, and max is the maximum value equal to 1.

$$Travelling_DR = 1 - \frac{l^{1/p}}{L^{1/p}} \quad (8)$$

where, p defines the accuracy of exploitation on iterations. Earlier and more accurate local search - exploitation is possible with a higher p value. Algorithm 1 shows the pseudocode of MVO.

Algorithm 1. MVO

```

1: Create random universes (UNV)
2: Initialize BestUNV, Wormhole_EP, Travelling_DR
3: SUNV = Sorted universes
4: NRM = Normalize inflation rates (fitness) of the universes
5: while Time < Max_time
6:   Update Travelling_DR, Wormhole_EP
7:   Evaluate the fitness of all universes
8:   for each universe indexed by i
9:     Black_Hole_Index = i;
10:    for each object indexed by j
11:      rn1 = random([0, 1]);
12:      if rn1 < NRM(UNVi)
13:        White_Hole_Index = RouletteWheelSelection(-NRM);
14:        UNV(Black_Hole_Index, j) = SUNV(White_Hole_Index, j);
15:      end if
16:      rn2 = random([0, 1]);
17:      if rn2 < Wormhole_EP
18:        rn3 = random([0, 1]);
19:        rn4 = random([0, 1]);
20:        if rn3 < 0.5
21:          UNV(i, j) = BestUNV(j) + Travelling_DR × ((uppb(j) - lowb(j)) × rn4 + lowb(j));
22:        else
23:          UNV(i, j) = BestUNV(j) - Travelling_DR × ((uppb(j) - lowb(j)) × rn4 + lowb(j));
24:        end if
25:      end if
26:    end for
27:  end for

```

28: end while**3.2. Simulated annealing (SA)**

Simulated annealing (SA) (Kirkpatrick et al., 1983), proposed by Kirkpatrick et al. in 1983, is one of the local search algorithms based on hill climbing method used in solving optimization problems. In the algorithm, where the creation of new solutions is based on predetermined rules or performed randomly, the newly created solution is compared with the current solution in each iteration. To find the global best and not get stuck with the local best, the algorithm, can accept new solutions that only enhance the current solution, as well as worse results that provide predetermined criteria. These criteria are decided by the Boltzmann probability. The following equation (Eq. 9) shows this mechanism:

$$Prb = e^{\left(\frac{-(F(X)-F(X_0))}{T_k}\right)} \quad (9)$$

where, Prb is defined as the acceptance probability. $F(X_0)$ defines the objective function for current solution, $F(X)$ defines the objective function for candidate solution. SA uses the acceptance probability mechanism to decide whether the candidate solution should be considered as the current solution when $F(X)$ is worse than $F(X_0)$. T_k is the temperature value at time k , and in each iteration, value of T_k is calculated as follows:

$$T_{k+1} = T_k \times c \quad (10)$$

where, c refers to the temperature coefficient, T_k is the initial temperature value and T_{k+1} is the temperature at time k . Algorithm 2 presents the pseudocode of SA algorithm.

Algorithm 2. SA

```

1:   $T$  = Temperature value
2:   $T_0$  = Final temperature value
3:   $c$  = Temperature coefficient
4:   $A$  = First solution
5:   $f(A)$  = Fitness value of the first solution
6:  while  $T > T_0$ 
7:       $A'$  = a new solution in the  $A$  neighbourhood
8:       $f(A')$  = Calculate the fitness value of  $A'$ 
9:      if  $f(A') < f(A)$ 
10:          $NSol = A'$ ;
11:          $f(NSol) = f(A')$ ;
12:      else
13:          $\Delta f = f(A') - f(A)$ 
14:          $r = \text{random}[0, 1]$ ;
15:         if  $r > \exp(-\Delta f / T)$ 
16:             $NSol = A'$ ;
17:             $f(NSol) = f(A')$ ;
18:         else
19:             $NSol = A$ ;
20:             $f(NSol) = f(A)$ ;
21:         end if
22:      end if
23:       $A = NSol$ ;
24:       $T = T \times c$ 
25:  end while

```

3.3. MVOSANN algorithm

MVO is a population-based meta-heuristic algorithm that produces very accomplished results in the face of many problems. Despite

the many advantages of the MVO algorithm, it also has disadvantages such as low accuracy, local minimum, and slow convergence (Jia et al., 2019; Song et al., 2020). In this part of the study, to eliminate these disadvantages improvements in the structure of the original MVO algorithm are suggested.

According to the uncertainty principle in physics, the momentum and position of a particle cannot be determined precisely at the same time, there is always an error, uncertainty and blurring possible (Heisenberg, 1985). According to the many-worlds interpretation (MWI), which is based on the uncertainty principle, there is a single and universal wave function for the entire universe as reality itself. This universal wave function, as the wave function of everything, includes all possibilities in the known world and an infinite number of parallel worlds in which every possibility exists (Everett et al., 1973).

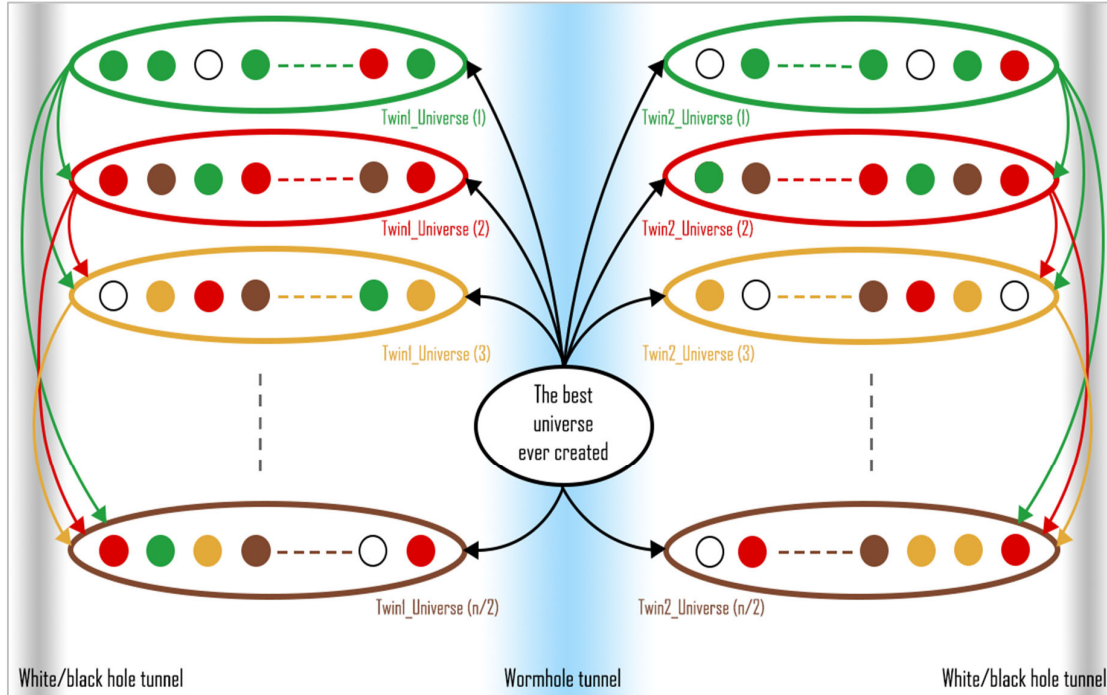


Fig. 3. Conceptual Model of MVOSANN Algorithm

Inspired by the multiverse theory and the big bang theory, the MVO algorithm has been modified based on the uncertainty principle in physics and the MWI. To get the global best without being stuck with local minimums search agents and their twins perform the search by acting according to the general rules of the MVO and the acceptance mechanism of the SA. In this method, we used the SA algorithm in the integrative hybrid model structure, where exploration and exploitation can be increased. Fig. 3 shows the conceptual model of the suggested MVOSANN algorithm.

In the suggested model, unlike the MVO algorithm, two initial populations with the same universes are created (Eq. 11). At each iteration, search agents in this twin population act according to the general rules of the MVO algorithm, to which the SA algorithm's acceptance probability feature is added. Each object represents a variable in the candidate solution and each universe in the populations represents a candidate solution.

$$TP = \left[\begin{array}{cccc} o_1^1 & o_1^2 & \cdots & o_1^k \\ o_2^1 & o_2^2 & \cdots & o_2^k \\ \vdots & \vdots & \ddots & \vdots \\ o_{n/2}^1 & o_{n/2}^2 & \cdots & o_{n/2}^k \end{array} \right] \left[\begin{array}{cccc} o_1^1 & o_1^2 & \cdots & o_1^k \\ o_2^1 & o_2^2 & \cdots & o_2^k \\ \vdots & \vdots & \ddots & \vdots \\ o_{n/2}^1 & o_{n/2}^2 & \cdots & o_{n/2}^k \end{array} \right] \quad (11)$$

where, k is the number of parameters (variables), n is the number of universes (candidate solutions).

In the MVO algorithm, updating the variables in a candidate solution is accomplished with a easy location update (Jia et al., 2019). The values of the fitness function of the solutions generated using the acceptance probability of the SA algorithm will not be prone to decrease constantly, and in some circumstances, also, the solutions with high fitness function values will be accepted and the searching for the global best will be effected. Equation (Eq. 5), which was used in the exploration stage of the original MVO algorithm, is suggested as follows to increase the exploration and exploitation of the universe around the best solution.

Algorithm 3. MVOSANN

```

1: Create twin populations randomly (TP)
2: Initialize BestUNV, Wormhole_EP, Travelling_DR, T, c
3: SUNV = Sorted universes
4: NRM = Normalize inflation rates (fitness) of the universes
5: while Time < Max_time
6:   Update Travelling_DR, Wormhole_EP
7:   Evaluate the fitness of universes of all populations
8:   Sort all universes of all populations
9:   TP[0] = SUNV[0:(N/2)] (Assign the first half of the sorted universes to the first of the twin populations)
10:  TP[1] = TP[0] (Equalize twin population)
11:  for each population indexed by k
12:    for each universe indexed by i
13:      Black_Hole_Index = i;
14:      for each object indexed by j
15:        rn1 = random ([0, 1]);
16:        if rn1 < NRM(SUNV [i,:])
17:          White_Hole_Index = RouletteWheelSelection(-NRM);
18:          TP[k][Black_Hole_Index][j] = SUNV[White_Hole_Index,j];
19:        else
20:          Calculate the fitness's of SUNV[Black_Hole_Index,:] and
                TP[k][Black_Hole_Index,:] (f())
21:           $\Delta f = f(\text{SUNV}[\text{Black\_Hole\_Index},:]) - f(\text{TP}[\text{k}][\text{Black\_Hole\_Index},:])$ 
22:          rn2 = random ([0, 1]);
23:          if rn2 <  $\exp(-\Delta f / T)$ 
24:            White_Hole_Index = RouletteWheelSelection(-NRM);
25:            TP[k][Black_Hole_Index][j] = SUNV[White_Hole_Index,j];
26:          end if
27:        end if
28:        rn3 = random ([0, 1]);
29:        if rn3 < Wormhole_EP
30:          rn4 = random ([0, 1]);
31:          rn5 = random ([0, 1]);
32:          if rn4 < 0.5
33:             $\text{TP}[\text{k}][\text{i}][\text{j}] = \text{BestUNV}(\text{j}) + \text{Travelling\_DR} \times ((\text{uppb}(\text{j}) - \text{lowb}(\text{j})) \times \text{rn5} + \text{lowb}(\text{j}));$ 
34:          else
35:             $\text{TP}[\text{k}][\text{i}][\text{j}] = \text{BestUNV}(\text{j}) - \text{Travelling\_DR} \times ((\text{uppb}(\text{j}) - \text{lowb}(\text{j})) \times \text{rn5} + \text{lowb}(\text{j}));$ 
36:          end if
37:        end if
38:      end for
39:    end for
40:  end for

```


41: $T = T \times c$

42: end while

$$o_i^j = \begin{cases} o_h^j & rn1 < NRM(SUNV[i,:]) \\ o_h^j & rn2 < Prb \\ o_i^j & rn2 \geq Prb \end{cases} \quad rn1 \geq NRM(SUNV[i,:]) \quad (12)$$

where, o_i^j defines the j^{th} parameter of i^{th} universe, o_h^j defines the j^{th} parameter of h^{th} universe chosen by a roulette wheel selection, $rn1$ and $rn2$ are numbers procured randomly in the range $[0, 1]$, $NRM(SUNV[i,:])$ is normalized inflation rate of the i^{th} sorted universe, $SUNV[i,:]$ describes the i^{th} sorted universe. Prb is defined as the acceptance probability (Eq. 9).

At the end of the iteration, the universes of the twin population clusters, which are no longer alike, are sorted collectively according to their inflation rates at each new iteration. The first universe in this ranking is the best universe ever found. After this stage, the first half of the collectively sorted universes with the best values is selected. This selected universe set creates the twin population set of the new iteration. The pseudo code of the MVOSANN algorithm is given in Algorithm 3.

This set of twin populations, with universes containing identical objects at the beginning, diverges from similarity at the end of each iteration. As a result of different exploration and exploitation possibilities in the universes of this twin population cluster at the same time, it is more likely to reach the best inflation value. Because, as mentioned in the uncertainty principle and the MWI, the state of an object cannot be determined simultaneously and there are always parallel worlds where other different possibilities take place.

4. MVOSANN for training MLP

In this section, it is explained how the MVOSANN algorithm is used in training a single hidden layer MLP network. As mentioned in the introduction, the aim of training the neural network is to find the best weight and bias values. This is the main task of the MVOSANN algorithm in this study. Each universe (search agent) in the algorithm consists of the connection weights among the input layer to the hidden layer (Iw_{nm}), the weights among the hidden layer to the output layer (Hw_m), and the bias weights (β_m).

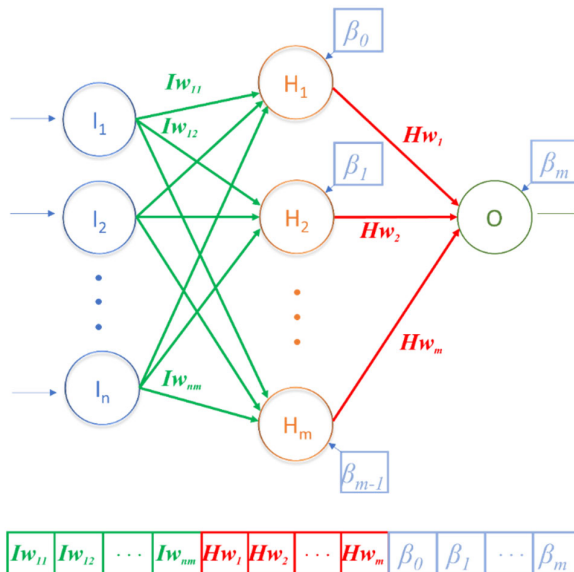


Fig. 4. Representation of the MVOSANN universe for the MLP network

The representation of the universes consisting of real numbers in the range of $[-1, 1]$ is given in Fig. 4 in vector structure.

Equation (Eq. 13) shows how the number of objects in each universe is calculated.

$$IndividualLength = (m \times n) + (2 \times m) + 1 \quad (13)$$

where, m is the number of neurons in the hidden layer, n is the number of input features.

In all data sets, the mean square error (MSE) fitness function, which is based on calculating the difference between the real values and the values predicted by MLP, was used to measure the fitness value of the universes produced by MVOSANN. The aim here is to minimize the MSE value until the last iteration. The mathematical representation of MSE is as follows.

$$MSE = \frac{1}{n} \sum_{i=1}^n (y - \hat{y})^2 \quad (14)$$

where, y is the actual values, and \hat{y} is the predicted values, n is the number of samples in the training dataset.

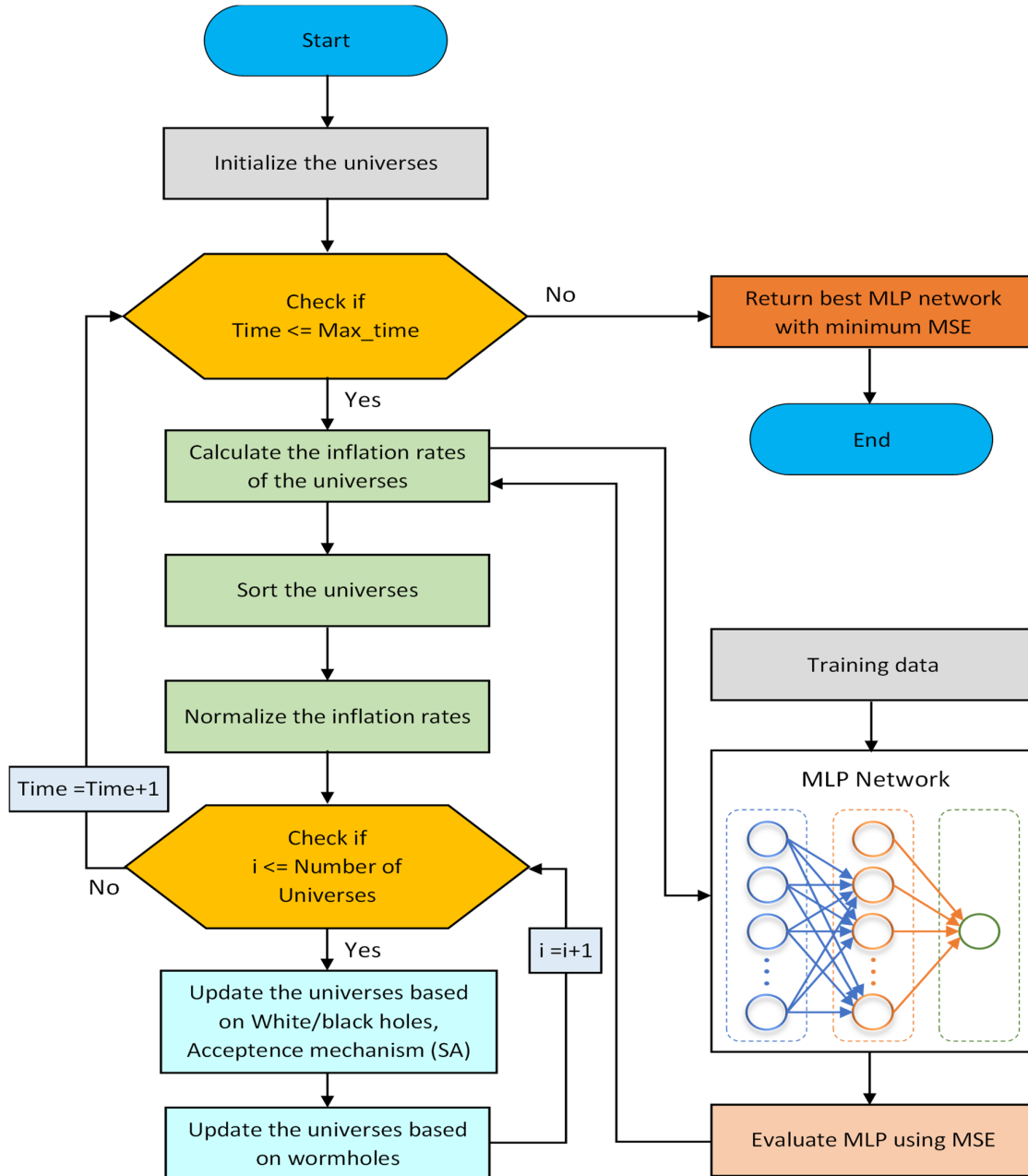


Fig. 5. General steps of the MVOSANN for the MLP network training

The MVOSANN algorithm is designed to train MLP networks as depicted in Fig. 5. We can summarize this approach with the following steps:

1. A predefined number of candidate solution sets are randomly generated, where each universe (candidate solution) represents an MLP network.
2. Universes consisting of all weights, including bias weights, are assigned to MLP networks. All generated MLP networks are evaluated using a fitness function (MSE). The aim is to reach the lowest MSE value according to the examples in the dataset used.
3. In this step, the location update of the universes is performed.
4. Steps 2 and 3 repeat up to the maximum number of iterations.

5. Experimental results

5.1. Datasets and comparison algorithms

In this study, a comprehensive analysis was carried out to examine the effectiveness of the MVOSANN in training MLP neural networks. MVO (Mirjalili et al., 2016) and 12 algorithms prevalently aforesaid in the literature: Grey Wolf Optimizer (GWO) (Mirjalili et al., 2017), Cuckoo Search (CS) (Yang & Deb, 2009), Harris Hawks Optimization (HHO) (Heidari et al., 2019), Genetic Algorithms (GA) (Holland, 1992), Differential Evolution (DE) (Storn & Price, 1997), Particle Swarm Optimization (PSO) (Kennedy & Eberhart, 1995), Firefly Algorithm (FA) (Yang, 2008, 2009), Salp Swarm Algorithm (SSA) (Mirjalili et al., 2017), Sine Cosine Algorithm (SCA) (Mirjalili, 2016), Whale Optimization Algorithm (WOA) (Mirjalili & Lewis, 2016), and JAYA Algorithm (JAYA) (Rao, 2016) were used to evaluate the achievement of the MVOSANN algorithm.

MVOSANN and other algorithms have been tested on 12 data sets used in many studies in the literature. These datasets, selected from the University of California, Irvine Machine Learning Repository (UCI), are: Abalone (Dua & Graff, 2019), Balance Scale (Dua & Graff, 2019), Blood (Yeh et al., 2009), Breast Cancer (Wolberg & Mangasarian, 1990; Bennett & Mangasarian, 1992), Diabetes (Dua & Graff, 2019), Acute Inflammations (Diagnosis I, Diagnosis II) (Czerniak & Zarzycki, 2003), Glass Identification (Dua & Graff, 2019), Iris (Dua & Graff, 2019), Liver disorders (Dua & Graff, 2019), Raisin (Çınar et al., 2020), Vertebral (Dua & Graff, 2019). Information about these datasets, such as the number of features, the number of training samples, and the number of test samples, are given in Table 1.

Table 1

Classification datasets

Dataset	Features	Training Samples	Testing Samples
Abalone	8	2756	1421
Balance Scale	4	412	213
Blood	4	493	255
Breast Cancer	8	461	238
Diabetes	8	506	262
Diagnosis I	6	79	41
Diagnosis II	6	79	41
Glass	10	141	73
Iris	4	99	51
Liver	6	79	41
Raisin	8	594	306
Vertebral	6	204	106

5.2. Experimental setup

Experiments were conducted using the EvoloPy-NN repository created by Faris in 2016 (Faris, 2016; Faris et al., 2016c). Evology-NN is an open-source framework consisting of classic and recent meta-heuristic algorithms coded with Python, with a user-friendly interface. Various studies have been carried out on artificial neural network training using the Evology-NN repository (Faris et al., 2016a; Faris et al., 2016b; Aljarah et al., 2018a; Aljarah et al., 2018b). All algorithms have been experimented on a personal computer with 16 GB (RAM), Intel Core i7-10875H 2.30 GHz (CPU), and 64-Bit Windows 11 (operating system).

Table 2 shows the parameter settings of the algorithms used in this study. All datasets are divided into 34 % for testing, and 66 % for training. The number of search agents for the algorithms is set to 50 and the number of iterations to 100. Each algorithm was run individually 30 times to obtain balanced performance results.

Table 2

Parameters and values of MVOSANN and other algorithms

Algorithm	Parameter	Value
MVOSANN	<i>Maximum wormhole existence probability</i>	1
	<i>Minimum wormhole existence probability</i>	0.2
	p	6
	T	1
	c	0.88
MVO	<i>Maximum wormhole existence probability</i>	1
	<i>Minimum wormhole existence probability</i>	0.2

	<i>p</i>	6
GA	<i>Crossover probability</i>	0.9
	<i>Mutation probability</i>	0.1
	<i>Selection mechanism</i>	Roulette wheel
FA	<i>Alpha</i>	0.5
	<i>Beta</i>	0.2
	<i>Gamma</i>	1
SSA	<i>No custom parameters</i>	
GWO	<i>No custom parameters</i>	
WOA	<i>b</i>	1
	<i>Vmax</i>	6
PSO	<i>c1</i>	2
	<i>c2</i>	2
	<i>wMax</i>	0.9
	<i>wMin</i>	0.2
CS	<i>p_a</i>	0.25
HHO	<i>No custom parameters</i>	
SCA	<i>a</i>	2
JAYA	<i>No custom parameters</i>	
DE	<i>Crossover probability</i>	0.9
	<i>Differential weight</i>	0.5

In this study, the method in which the number of hidden neurons is $2 \times N + 1$ was chosen; N is the number of features in each dataset. For each dataset, all input features values are normalized in the range [0,1] with Min-max normalization technique (Eq. 15).

$$v' = \frac{v_i - \min_A}{\max_A - \min_A} \tag{15}$$

where, v' is the normalized value of v in the range $[\min_A, \max_A]$

5.3. Evaluation metrics

A confusion matrix is a table (Fig. 6), that helps us evaluate the accuracy of a classification model on a dataset with actual values. On this table is a summary of the results predicted by the model in a classification problem (Stehman, 1997). In the model, evaluation was made by calculating the number of correctly predicted samples belong to the class (true positives), the number of incorrectly predicted samples belong to the class (false positives), the number of correctly predicted samples that do not belong to the class (true negatives), and the number of incorrectly predicted samples that do not belong to the class (false negatives). How the metrics used for binary classification and multiclass classification are calculated using the confusion matrix values, are shown in Table 3. (Sokolova & Lapalme, 2009).

		Actual Class	
		Positive	Negative
Predicted Class	Positive	True Positive (TP)	False Positive (FP)
	Negative	False Negative (FN)	True Negative (TN)

Fig. 6. Confusion Matrix

In this study, during the performance evaluation of algorithms, accuracy, recall, precision, error rate and F1 score calculations were made depending on whether each data set was in a binary classification group or a multiclass classification group. In the multi-class classification group, the calculations were made using the micro-average method, considering the class imbalance in the data sets. In addition, the MSE values (Eq. 14) based on the calculation of the difference among the actual values and the predicted values of MLP, and the standard deviations of the accuracy and MSE values were calculated. All algorithms were evaluated for each data set with the results obtained from the calculations. Also, a nonparametric Wilcoxon rank sum test (Wilcoxon, 1992) was applied to the results to analyse the relationship among the algorithms, and the p value was accepted to be less than 0.05 (5E-02) in this test to evaluate the statistically significant difference.

Table 3

Measures for binary classification and multi-class classification

Binary classification		Multiclass classification	
Measure	Formula	Measure	Formula
Accuracy	$\frac{TP + TN}{TP + FN + FP + TN}$	Average_Accuracy	$\frac{\sum_{i=1}^k \frac{TP_i + TN_i}{TP_i + FN_i + FP_i + TN_i}}{k}$
Precision	$\frac{TP}{TP + FP}$	Precision _μ	$\frac{\sum_{i=1}^k TP_i}{\sum_{i=1}^k (TP_i + FP_i)}$
Recall	$\frac{TP}{TP + FN}$	Recall _μ	$\frac{\sum_{i=1}^k TP_i}{\sum_{i=1}^k (TP_i + FN_i)}$
F1_score	$2 \times \frac{Precision \times Recall}{Precision + Recall}$	F1_score _μ	$2 \times \frac{Precision_{\mu} \times Recall_{\mu}}{Precision_{\mu} + Recall_{\mu}}$
Error rate	$\frac{FP + FN}{TP + FN + FP + TN}$	Error_rate _μ	$\frac{\sum_{i=1}^k \frac{FP_i + FN_i}{TP_i + FN_i + FP_i + TN_i}}{k}$

5.3. Results and discussions

To evaluate the suggested MVOSANN algorithm, accuracy, MSE, recall, precision, error rate and F1 score results of all algorithms obtained from 12 data sets were compared. For each dataset and 30 trials, Table 4 shows the accuracy averages,

standard deviations of the mean accuracies of all algorithms, and best accuracies. According to the results, MVOSANN performs much better than all other algorithms by having the best average accuracy values in all data sets. In addition, MVOSANN has low standard deviation values for all data sets, which shows that the proposed algorithm is robust and stable. Considering the best accuracy values, MVOSANN, 5 achieved the highest accuracy values for the data set, and it can be said that it is superior when compared to other algorithms.

Table 4
Results of classification accuracy for all datasets and algorithms

Datasets		Algorithms												
		MVOSA	MVOGA	FA	SSA	PSO	GWO	WOA	CS	HHO	SCA	JAYA	DE	
Abalone	AVE	0.8230	0.821	0.8047	0.8139	0.8100	0.8122	0.8155	0.7877	0.8066	0.8009	0.8013	0.8040	0.8012
	STD	0.0031	0.004	0.0101	0.0080	0.0077	0.0098	0.0058	0.0150	0.0131	0.0100	0.0128	0.0157	0.0199
	BEST	0.8291	0.831	0.8249	0.8270	0.8256	0.8277	0.8242	0.8186	0.8326	0.8136	0.8418	0.8319	0.8291
Balance	AVE	0.8376	0.830	0.7746	0.8144	0.8239	0.8282	0.8194	0.7573	0.7850	0.7624	0.7156	0.7654	0.7019
	STD	0.0490	0.049	0.0787	0.0682	0.0587	0.0664	0.0519	0.0776	0.0919	0.0660	0.1043	0.1273	0.0874
	BEST	0.9296	0.910	0.9296	0.9155	0.9343	0.9343	0.9108	0.9249	0.9014	0.8873	0.8638	0.9437	0.8498
Blood	AVE	0.7906	0.788	0.7837	0.7829	0.7808	0.7843	0.7822	0.7838	0.7829	0.7838	0.7826	0.7835	0.7822
	STD	0.0038	0.005	0.0051	0.0054	0.0038	0.0074	0.0048	0.0067	0.0065	0.0070	0.0074	0.0072	0.0083
	BEST	0.7961	0.796	0.7922	0.7922	0.7882	0.8000	0.7922	0.7922	0.7961	0.7922	0.7922	0.7961	0.7961
Breast Cancer	AVE	0.9777	0.975	0.9678	0.9711	0.9721	0.9655	0.9746	0.9682	0.9664	0.9696	0.9602	0.9578	0.9571
	STD	0.0044	0.005	0.0085	0.0063	0.0063	0.0072	0.0055	0.0089	0.0085	0.0049	0.0107	0.0103	0.0104
	BEST	0.9874	0.983	0.9832	0.9832	0.9832	0.9748	0.9832	0.9790	0.9790	0.9790	0.9832	0.9748	0.9748
Diabetes	AVE	0.7579	0.753	0.7246	0.7490	0.7480	0.7439	0.7497	0.6957	0.7247	0.7118	0.7078	0.7230	0.7289
	STD	0.0057	0.005	0.0236	0.0121	0.0075	0.0136	0.0063	0.0329	0.0229	0.0254	0.0267	0.0231	0.0235
	BEST	0.7710	0.763	0.7748	0.7748	0.7672	0.7748	0.7672	0.7824	0.7672	0.7481	0.7595	0.7672	0.7710
Diagnosis I	AVE	0.9317	0.915	0.8431	0.8455	0.8683	0.8537	0.9049	0.8106	0.9065	0.8350	0.8293	0.8756	0.8236
	STD	0.1057	0.109	0.1170	0.1127	0.1177	0.1133	0.1091	0.1296	0.1150	0.1143	0.1137	0.1321	0.1050
	BEST	1.0000	1.000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000	1.0000
Diagnosis II	AVE	0.8333	0.825	0.7894	0.8138	0.8163	0.8024	0.8146	0.8000	0.8057	0.8073	0.7829	0.7911	0.7911
	STD	0.0129	0.019	0.0606	0.0244	0.0210	0.0407	0.0165	0.0422	0.0359	0.0074	0.0688	0.0596	0.0662
	BEST	0.8537	0.853	0.8293	0.8537	0.8537	0.8537	0.8537	0.8293	0.8293	0.8293	0.8293	0.8293	0.8537
Glass	AVE	0.8810	0.868	0.8625	0.8606	0.8722	0.8750	0.8653	0.8769	0.8810	0.8755	0.8736	0.8514	0.8597
	STD	0.0135	0.025	0.0297	0.0246	0.0191	0.0287	0.0147	0.0455	0.0364	0.0220	0.0395	0.0331	0.0615
	BEST	0.9028	0.930	0.9306	0.9167	0.9167	0.9167	0.9028	0.9306	0.9583	0.9167	0.9444	0.9306	0.9583
Iris	AVE	0.7607	0.741	0.7033	0.7467	0.7273	0.7380	0.7267	0.7020	0.7147	0.7060	0.7067	0.7153	0.7160
	STD	0.0299	0.024	0.0283	0.0325	0.0170	0.0491	0.0184	0.0280	0.0283	0.0253	0.0275	0.0460	0.0373
	BEST	0.8200	0.780	0.7400	0.8400	0.7600	0.9200	0.7600	0.7400	0.7800	0.7400	0.7400	0.8400	0.8200
Liver	AVE	0.7686	0.762	0.6952	0.7395	0.7359	0.7455	0.7517	0.6322	0.6870	0.6630	0.6658	0.6661	0.6661
	STD	0.0132	0.012	0.0361	0.0259	0.0231	0.0234	0.0180	0.0460	0.0307	0.0463	0.0458	0.0375	0.0342
	BEST	0.7966	0.796	0.7712	0.7797	0.7797	0.7797	0.7797	0.7797	0.7119	0.7627	0.7627	0.7542	0.7203
Raisin	AVE	0.8596	0.859	0.8477	0.8527	0.8529	0.8507	0.8558	0.8354	0.8450	0.8402	0.8404	0.8385	0.8369
	STD	0.0041	0.004	0.0074	0.0061	0.0051	0.0131	0.0053	0.0145	0.0123	0.0147	0.0152	0.0148	0.0186
	BEST	0.8693	0.869	0.8595	0.8627	0.8627	0.8660	0.8627	0.8529	0.8627	0.8660	0.8562	0.8627	0.8627
Vertebral	AVE	0.8777	0.873	0.8189	0.8509	0.8604	0.8440	0.8701	0.7833	0.8220	0.7947	0.7906	0.8057	0.8088
	STD	0.0107	0.007	0.0380	0.0368	0.0242	0.0398	0.0178	0.0480	0.0331	0.0362	0.0390	0.0370	0.0327
	BEST	0.8962	0.886	0.8962	0.9151	0.8962	0.8962	0.8962	0.8585	0.8774	0.8679	0.8491	0.8774	0.8774

In 30 trials, for all datasets and all algorithms, the mean of MSE values, the standard deviation of the mean MSE values, and the lowest MSE values are shown in Table 5. As can be seen from the results, MVOSANN has the best average MSE for all datasets. When the standard deviation values are examined, it is also supported that MVOSANN is a robust and stable algorithm. In addition, MVOSANN achieved the lowest MSE value for the 9 dataset and is in the front row compared to other algorithms.

Table 5
Results of MSE for all datasets and algorithms

Datasets	Algorithms													
	MVOSA	MVO	GA	FA	SSA	PSO	GWO	WOA	CS	HHO	SCA	JAYA	DE	
Abalone	AVE	0.126	0.1275	0.1332	0.1299	0.1311	0.1311	0.1290	0.1379	0.1329	0.1340	0.1351	0.1345	0.1352
	STD	0.000	0.0008	0.0017	0.0022	0.0015	0.0025	0.0012	0.0038	0.0010	0.0022	0.0017	0.0022	0.0017
	BES	0.125	0.1259	0.1287	0.1259	0.1279	0.1268	0.1272	0.1299	0.1312	0.1304	0.1316	0.1296	0.1319
Balance	AVE	0.043	0.0461	0.0599	0.0532	0.0531	0.0491	0.0521	0.0616	0.0569	0.0610	0.0632	0.0591	0.0610
	STD	0.002	0.0021	0.0032	0.0045	0.0034	0.0034	0.0029	0.0044	0.0023	0.0033	0.0035	0.0041	0.0032
	BES	0.040	0.0425	0.0519	0.0447	0.0465	0.0424	0.0483	0.0546	0.0516	0.0532	0.0574	0.0494	0.0560
Blood	AVE	0.154	0.1548	0.1581	0.1571	0.1572	0.1558	0.1569	0.1602	0.1579	0.1595	0.1598	0.1588	0.1599
	STD	0.000	0.0009	0.0010	0.0016	0.0011	0.0012	0.0010	0.0038	0.0006	0.0022	0.0012	0.0015	0.0016
	BES	0.152	0.1534	0.1560	0.1544	0.1547	0.1535	0.1547	0.1563	0.1559	0.1563	0.1576	0.1548	0.1554
Breast Cancer	AVE	0.032	0.0339	0.0425	0.0389	0.0386	0.0420	0.0359	0.0478	0.0445	0.0413	0.0510	0.0520	0.0543
	STD	0.001	0.0015	0.0025	0.0028	0.0017	0.0033	0.0016	0.0045	0.0017	0.0013	0.0049	0.0049	0.0047
	BES	0.031	0.0320	0.0362	0.0349	0.0350	0.0342	0.0316	0.0392	0.0405	0.0369	0.0417	0.0430	0.0414
Diabetes	AVE	0.150	0.1517	0.1675	0.1561	0.1557	0.1570	0.1539	0.1842	0.1685	0.1716	0.1755	0.1705	0.1721
	STD	0.001	0.0009	0.0061	0.0027	0.0018	0.0027	0.0010	0.0143	0.0035	0.0069	0.0051	0.0043	0.0045
	BES	0.148	0.1496	0.1571	0.1519	0.1525	0.1517	0.1517	0.1582	0.1627	0.1583	0.1685	0.1623	0.1629
Diagnosis I	AVE	0.090	0.0913	0.0963	0.0931	0.0936	0.0921	0.0928	0.1049	0.0959	0.1001	0.1021	0.0976	0.1003
	STD	0.000	0.0008	0.0019	0.0015	0.0012	0.0029	0.0009	0.0147	0.0015	0.0077	0.0038	0.0028	0.0023
	BES	0.090	0.0898	0.0936	0.0902	0.0912	0.0899	0.0914	0.0941	0.0927	0.0933	0.0951	0.0934	0.0967
Diagnosis II	AVE	0.112	0.1177	0.1459	0.1292	0.1319	0.1169	0.1351	0.1492	0.1409	0.1482	0.1515	0.1444	0.1509
	STD	0.002	0.0046	0.0049	0.0083	0.0071	0.0070	0.0054	0.0089	0.0041	0.0063	0.0066	0.0083	0.0052
	BES	0.107	0.1096	0.1354	0.1123	0.1198	0.1078	0.1247	0.1311	0.1314	0.1347	0.1340	0.1265	0.1394
Glass	AVE	0.028	0.0296	0.0397	0.0361	0.0355	0.0380	0.0325	0.0532	0.0419	0.0408	0.0490	0.0481	0.0534
	STD	0.001	0.0015	0.0037	0.0028	0.0022	0.0037	0.0012	0.0126	0.0017	0.0046	0.0060	0.0043	0.0051
	BES	0.025	0.0266	0.0343	0.0299	0.0288	0.0321	0.0299	0.0368	0.0375	0.0353	0.0395	0.0418	0.0422
Iris	AVE	0.011	0.0126	0.0189	0.0142	0.0167	0.0147	0.0158	0.0231	0.0186	0.0214	0.0215	0.0192	0.0207
	STD	0.001	0.0013	0.0015	0.0030	0.0018	0.0023	0.0017	0.0034	0.0010	0.0020	0.0019	0.0019	0.0020
	BES	0.009	0.0111	0.0161	0.0090	0.0127	0.0110	0.0121	0.0166	0.0161	0.0181	0.0179	0.0158	0.0155
Liver	AVE	0.192	0.1960	0.2157	0.2026	0.2064	0.1972	0.2024	0.2257	0.2147	0.2212	0.2206	0.2158	0.2191
	STD	0.002	0.0029	0.0025	0.0055	0.0038	0.0044	0.0026	0.0064	0.0026	0.0061	0.0039	0.0046	0.0036
	BES	0.189	0.1905	0.2119	0.1927	0.1989	0.1890	0.1969	0.2137	0.2096	0.2089	0.2124	0.2098	0.2126
Raisin	AVE	0.102	0.1036	0.1152	0.1086	0.1108	0.1050	0.1087	0.1249	0.1139	0.1209	0.1186	0.1160	0.1176
	STD	0.001	0.0018	0.0031	0.0038	0.0026	0.0036	0.0017	0.0092	0.0019	0.0058	0.0027	0.0034	0.0029
	BES	0.098	0.0992	0.1100	0.1012	0.1074	0.0990	0.1051	0.1106	0.1092	0.1124	0.1135	0.1100	0.1110
Vertebral	AVE	0.128	0.1314	0.1475	0.1378	0.1398	0.1348	0.1364	0.1631	0.1478	0.1562	0.1559	0.1518	0.1551
	STD	0.001	0.0018	0.0046	0.0057	0.0042	0.0038	0.0040	0.0108	0.0037	0.0069	0.0058	0.0060	0.0061
	BES	0.124	0.1278	0.1385	0.1285	0.1343	0.1265	0.1306	0.1490	0.1345	0.1424	0.1448	0.1412	0.1429

Table 6 contains the results obtained using the Recall, Error rate, F1 score, and Precision evaluation measures of all algorithms for all datasets. According to the results, in the training process, MVOSANN has the best values in 7 data sets (Balance, Blood, Diabetes, Diagnosis I, Glass, Iris, Liver) for precision measurement, 9 data sets (Abalone, Balance, Breast Cancer, Diabetes, Diagnosis I, Glass, Iris, Liver, Raisin) for recall measurement, 11 data sets (Abalone, Balance, Blood, Diabetes, Diagnosis I, Diagnosis II, Glass, Iris, Liver, Raisin, Vertebral) for F1 score measurement, and 11 data sets (Abalone, Balance, Blood, Diabetes, Diagnosis I, Diagnosis II, Glass, Iris, Liver, Raisin, Vertebral) for error rate measurement. In the experimenting stage, MVOSANN has the best values in 7 data sets (Blood, Diabetes, Diagnosis I, Glass, Iris, Liver, Vertebral) for precision measurement, 9 data sets (Abalone, Breast Cancer, Diabetes, Diagnosis I, Diagnosis II, Glass, Iris, Liver, Raisin) for recall measurement, 12 data sets (Abalone, Balance, Blood, Breast Cancer, Diabetes, Diagnosis I, Diagnosis II, Glass, Iris, Liver, Raisin, Vertebral) for F1 score measurement, and 12 data sets (Abalone, Balance, Blood, Breast Cancer, Diabetes, Diagnosis I, Diagnosis II, Glass, Iris, Liver, Raisin, Vertebral) for error rate measurement. The outcomes again show the supremacy of MVOSANN over other algorithms according to these four evaluation measures.

Table 6
Results of precision, recall, F1 score, error rate for all datasets (continued overleaves)

Datasets	Algorithms	Train				Test			
		Precision	Recall	F1 Score	Error	Precision	Recall	F1 Score	Error
Abalone	MVOSANN	0.9027	0.7815	0.8377	0.2055	0.8857	0.8488	0.8668	0.1770
	MVO	0.9018	0.7794	0.8361	0.2074	0.8844	0.8469	0.8652	0.1790
	GA	0.9055	0.7474	0.8184	0.2247	0.8878	0.8159	0.8499	0.1953
	FA	0.9003	0.7719	0.8310	0.2130	0.8804	0.8400	0.8597	0.1861
	SSA	0.9020	0.7650	0.8279	0.2159	0.8804	0.8332	0.8561	0.1900
	PSO	0.8933	0.7745	0.8288	0.2167	0.8771	0.8425	0.8587	0.1878
	GWO	0.9006	0.7749	0.8330	0.2108	0.8795	0.8437	0.8612	0.1845
	WOA	0.8961	0.7220	0.7989	0.2460	0.8794	0.7974	0.8358	0.2123
	CS	0.9037	0.7488	0.8183	0.2251	0.8883	0.8187	0.8514	0.1934
	HHO	0.9019	0.7459	0.8163	0.2276	0.8821	0.8160	0.8475	0.1991
	SCA	0.9017	0.7418	0.8132	0.2306	0.8846	0.8143	0.8473	0.1987
	JAYA	0.8932	0.7593	0.8193	0.2261	0.8776	0.8286	0.8510	0.1960
DE	0.8942	0.7545	0.8161	0.2288	0.8787	0.8230	0.8480	0.1988	
Balance	MVOSANN	1.0000	0.5420	0.7023	0.2468	0.9715	0.7200	0.8246	0.1624
	MVO	1.0000	0.5132	0.6771	0.2623	0.9701	0.7096	0.8149	0.1695
	GA	1.0000	0.3462	0.5098	0.3523	0.9661	0.6107	0.7320	0.2254
	FA	1.0000	0.4089	0.5770	0.3185	0.9786	0.6728	0.7880	0.1856
	SSA	1.0000	0.3980	0.5676	0.3244	0.9807	0.6896	0.8031	0.1761
	PSO	0.9992	0.4596	0.6263	0.2914	0.9490	0.7249	0.8128	0.1718
	GWO	1.0000	0.4191	0.5891	0.3130	0.9777	0.6814	0.7987	0.1806
	WOA	0.9985	0.3110	0.4716	0.3715	0.9825	0.5641	0.7013	0.2427
	CS	0.9989	0.3583	0.5211	0.3460	0.9645	0.6304	0.7435	0.2150
	HHO	1.0000	0.3096	0.4710	0.3720	0.9880	0.5693	0.7122	0.2376
	SCA	0.9956	0.3170	0.4723	0.3689	0.9746	0.4951	0.6226	0.2844
	JAYA	0.9975	0.3785	0.5408	0.3354	0.9505	0.6070	0.7222	0.2346
DE	0.9968	0.3107	0.4651	0.3721	0.9532	0.4835	0.6126	0.2981	
Blood	MVOSANN	0.7821	0.9799	0.8699	0.2236	0.7977	0.9711	0.8759	0.2094
	MVO	0.7795	0.9819	0.8691	0.2256	0.7945	0.9732	0.8748	0.2119
	GA	0.7718	0.9881	0.8667	0.2319	0.7881	0.9789	0.8732	0.2163
	FA	0.7731	0.9855	0.8664	0.2317	0.7886	0.9765	0.8725	0.2171
	SSA	0.7715	0.9864	0.8658	0.2331	0.7877	0.9746	0.8712	0.2192
	PSO	0.7793	0.9816	0.8687	0.2263	0.7915	0.9730	0.8729	0.2157
	GWO	0.7737	0.9841	0.8663	0.2316	0.7891	0.9741	0.8719	0.2178
	WOA	0.7703	0.9899	0.8664	0.2329	0.7852	0.9856	0.8740	0.2162
	CS	0.7729	0.9853	0.8662	0.2321	0.7886	0.9765	0.8725	0.2171
	HHO	0.7697	0.9910	0.8665	0.2330	0.7853	0.9854	0.8740	0.2162
	SCA	0.7722	0.9879	0.8668	0.2316	0.7865	0.9808	0.8728	0.2174
	JAYA	0.7781	0.9823	0.8682	0.2275	0.7904	0.9741	0.8726	0.2165
DE	0.7739	0.9861	0.8671	0.2305	0.7865	0.9801	0.8726	0.2178	
Breast Cancer	MVOSANN	0.9781	0.9598	0.9689	0.0403	0.9832	0.9830	0.9831	0.0223
	MVO	0.9767	0.9591	0.9678	0.0416	0.9820	0.9807	0.9813	0.0246
	GA	0.9824	0.9507	0.9663	0.0433	0.9862	0.9648	0.9753	0.0322
	FA	0.9808	0.9560	0.9683	0.0409	0.9843	0.9718	0.9780	0.0289
	SSA	0.9825	0.9564	0.9693	0.0396	0.9858	0.9718	0.9787	0.0279
	PSO	0.9813	0.9480	0.9643	0.0458	0.9864	0.9611	0.9735	0.0345
	GWO	0.9763	0.9584	0.9673	0.0424	0.9799	0.9817	0.9808	0.0254
	WOA	0.9815	0.9461	0.9634	0.0469	0.9857	0.9658	0.9757	0.0318
	CS	0.9791	0.9480	0.9632	0.0473	0.9853	0.9637	0.9742	0.0336
	HHO	0.9837	0.9555	0.9694	0.0394	0.9845	0.9692	0.9768	0.0304
	SCA	0.9768	0.9406	0.9582	0.0535	0.9818	0.9577	0.9695	0.0398
	JAYA	0.9781	0.9395	0.9583	0.0534	0.9808	0.9550	0.9676	0.0422
DE	0.9726	0.9405	0.9561	0.0565	0.9786	0.9563	0.9672	0.0429	

Table 6. Continued

Datasets	Algorithms	Train				Test			
		Precision	Recall	F1 Score	Error	Precision	Recall	F1 Score	Error
Diabetes	MVOSANN	0.7666	0.5419	0.6348	0.2119	0.7428	0.5191	0.6109	0.2421
	MVO	0.7604	0.5388	0.6306	0.2146	0.7355	0.5122	0.6037	0.2463
	GA	0.7338	0.4674	0.5665	0.2398	0.7074	0.4309	0.5313	0.2754
	FA	0.7464	0.5194	0.6124	0.2235	0.7350	0.4931	0.5899	0.2510
	SSA	0.7549	0.5176	0.6140	0.2211	0.7321	0.4931	0.5890	0.2520
	PSO	0.7589	0.5151	0.6133	0.2206	0.7332	0.4750	0.5758	0.2561
	GWO	0.7591	0.5339	0.6268	0.2161	0.7329	0.4990	0.5936	0.2503
	WOA	0.6768	0.3820	0.4853	0.2704	0.6495	0.3670	0.4669	0.3043
	CS	0.7275	0.4847	0.5805	0.2374	0.6938	0.4479	0.5427	0.2753
	HHO	0.7144	0.4562	0.5550	0.2469	0.6684	0.4243	0.5177	0.2882
	SCA	0.7142	0.4289	0.5308	0.2540	0.6819	0.3962	0.4960	0.2922
	JAYA	0.7377	0.4756	0.5741	0.2373	0.6917	0.4424	0.5361	0.2770
	DE	0.7282	0.4558	0.5565	0.2441	0.7228	0.4312	0.5353	0.2711
Diagnosis I	MVOSANN	1.0000	0.7500	0.8571	0.1266	1.0000	0.8667	0.9142	0.0683
	MVO	1.0000	0.7500	0.8571	0.1266	1.0000	0.8349	0.8943	0.0846
	GA	1.0000	0.7500	0.8571	0.1266	1.0000	0.6937	0.7995	0.1569
	FA	1.0000	0.7500	0.8571	0.1266	1.0000	0.6984	0.8042	0.1545
	SSA	1.0000	0.7500	0.8571	0.1266	1.0000	0.7429	0.8332	0.1317
	PSO	1.0000	0.7500	0.8571	0.1266	1.0000	0.7143	0.8151	0.1463
	GWO	1.0000	0.7500	0.8571	0.1266	1.0000	0.8143	0.8819	0.0951
	WOA	0.9926	0.7417	0.8486	0.1333	0.9856	0.6444	0.7715	0.1894
	CS	1.0000	0.7500	0.8571	0.1266	1.0000	0.8175	0.8821	0.0935
	HHO	1.0000	0.7500	0.8571	0.1266	1.0000	0.6778	0.7889	0.1650
	SCA	1.0000	0.7500	0.8571	0.1266	1.0000	0.6667	0.7813	0.1707
	JAYA	1.0000	0.7500	0.8571	0.1266	1.0000	0.7571	0.8348	0.1244
	DE	1.0000	0.7500	0.8571	0.1266	1.0000	0.6556	0.7758	0.1764
Diagnosis II	MVOSANN	0.8853	0.8812	0.8830	0.1359	0.7993	0.9653	0.8705	0.1667
	MVO	0.8796	0.8848	0.8819	0.1380	0.8242	0.9236	0.8580	0.1748
	GA	0.8917	0.7616	0.8039	0.2059	0.9303	0.7222	0.7999	0.2106
	FA	0.8839	0.8616	0.8709	0.1473	0.9101	0.7931	0.8289	0.1862
	SSA	0.8876	0.8449	0.8646	0.1532	0.9132	0.7903	0.8291	0.1837
	PSO	0.8725	0.8783	0.8750	0.1460	0.8518	0.8542	0.8313	0.1976
	GWO	0.8919	0.8275	0.8562	0.1599	0.9398	0.7514	0.8221	0.1854
	WOA	0.9277	0.6543	0.7558	0.2363	0.9268	0.7431	0.8103	0.2000
	CS	0.8607	0.8362	0.8407	0.1823	0.8910	0.7944	0.8223	0.1943
	HHO	0.9324	0.6783	0.7693	0.2253	0.9744	0.7000	0.8077	0.1927
	SCA	0.8188	0.8072	0.7850	0.2435	0.8715	0.7875	0.8053	0.2171
	JAYA	0.8764	0.7790	0.8071	0.2084	0.9253	0.7347	0.8026	0.2089
	DE	0.8825	0.7420	0.7767	0.2325	0.8928	0.7861	0.8092	0.2089
Glass	MVOSANN	0.9909	0.8892	0.9373	0.1031	1.0000	0.8618	0.9257	0.1190
	MVO	0.9909	0.8851	0.9350	0.1066	1.0000	0.8468	0.9168	0.1319
	GA	0.9872	0.8756	0.9280	0.1176	1.0000	0.8403	0.9129	0.1375
	FA	0.9890	0.8780	0.9302	0.1141	1.0000	0.8382	0.9117	0.1394
	SSA	0.9897	0.8840	0.9338	0.1085	1.0000	0.8516	0.9197	0.1278
	PSO	0.9890	0.8780	0.9301	0.1141	1.0000	0.8548	0.9214	0.1250
	GWO	0.9906	0.8821	0.9332	0.1094	1.0000	0.8435	0.9150	0.1347
	WOA	0.9773	0.8783	0.9247	0.1232	1.0000	0.8570	0.9221	0.1231
	CS	0.9841	0.8821	0.9301	0.1146	1.0000	0.8618	0.9253	0.1190
	HHO	0.9874	0.8892	0.9356	0.1059	1.0000	0.8554	0.9219	0.1245
	SCA	0.9809	0.8694	0.9214	0.1279	0.9988	0.8543	0.9202	0.1264
	JAYA	0.9852	0.8715	0.9246	0.1228	1.0000	0.8274	0.9051	0.1486
	DE	0.9844	0.8585	0.9163	0.1345	0.9993	0.8376	0.9097	0.1403

Table 6. Continued

Datasets	Algorithms	Train				Test			
		Precision	Recall	F1 Score	Error	Precision	Recall	F1 Score	Error
Iris	MVOSANN	1.0000	0.6242	0.7682	0.2480	1.0000	0.6480	0.7856	0.2393
	MVO	1.0000	0.6081	0.7559	0.2587	1.0000	0.6196	0.7645	0.2587
	GA	1.0000	0.5384	0.6997	0.3047	1.0000	0.5637	0.7201	0.2967
	FA	1.0000	0.6010	0.7501	0.2633	1.0000	0.6275	0.7701	0.2533
	SSA	1.0000	0.5672	0.7236	0.2857	1.0000	0.5990	0.7489	0.2727
	PSO	1.0000	0.5934	0.7438	0.2683	1.0000	0.6147	0.7592	0.2620
	GWO	1.0000	0.5601	0.7179	0.2903	1.0000	0.5980	0.7481	0.2733
	WOA	0.9990	0.5202	0.6839	0.3170	1.0000	0.5618	0.7185	0.2980
	CS	1.0000	0.5505	0.7096	0.2967	1.0000	0.5804	0.7336	0.2853
	HHO	1.0000	0.5283	0.6911	0.3113	1.0000	0.5676	0.7235	0.2940
	SCA	1.0000	0.5439	0.7041	0.3010	1.0000	0.5686	0.7242	0.2933
	JAYA	1.0000	0.5601	0.7168	0.2903	1.0000	0.5814	0.7331	0.2847
	DE	1.0000	0.5576	0.7152	0.2920	1.0000	0.5824	0.7346	0.2840
Liver	MVOSANN	0.7262	0.5474	0.6241	0.2759	0.7444	0.6920	0.7170	0.2314
	MVO	0.7113	0.5365	0.6115	0.2852	0.7365	0.6860	0.7101	0.2373
	GA	0.6613	0.4421	0.5243	0.3302	0.6932	0.5273	0.5915	0.3048
	FA	0.6953	0.5161	0.5922	0.2972	0.7144	0.6427	0.6761	0.2605
	SSA	0.6837	0.4937	0.5730	0.3075	0.7119	0.6327	0.6691	0.2641
	PSO	0.7144	0.5260	0.6045	0.2872	0.7250	0.6460	0.6823	0.2545
	GWO	0.6820	0.5140	0.5860	0.3038	0.7251	0.6687	0.6953	0.2483
	WOA	0.6353	0.3414	0.4367	0.3568	0.6169	0.3607	0.4446	0.3678
	CS	0.6743	0.4365	0.5240	0.3266	0.6812	0.5160	0.5789	0.3130
	HHO	0.6747	0.3842	0.4823	0.3361	0.6487	0.4300	0.5114	0.3370
	SCA	0.6509	0.3961	0.4716	0.3501	0.6844	0.4533	0.5166	0.3342
	JAYA	0.6589	0.4354	0.5098	0.3373	0.6627	0.4933	0.5436	0.3339
	DE	0.6849	0.4042	0.4882	0.3363	0.6908	0.4640	0.5261	0.3339
Raisin	MVOSANN	0.8713	0.8897	0.8804	0.1209	0.8291	0.9059	0.8658	0.1404
	MVO	0.8703	0.8872	0.8787	0.1225	0.8304	0.9035	0.8654	0.1405
	GA	0.8764	0.8617	0.8688	0.1300	0.8334	0.8697	0.8508	0.1523
	FA	0.8717	0.8763	0.8740	0.1264	0.8304	0.8867	0.8575	0.1473
	SSA	0.8740	0.8728	0.8734	0.1265	0.8347	0.8802	0.8568	0.1471
	PSO	0.8725	0.8808	0.8762	0.1241	0.8276	0.8867	0.8554	0.1493
	GWO	0.8738	0.8725	0.8731	0.1268	0.8360	0.8852	0.8599	0.1442
	WOA	0.8681	0.8510	0.8591	0.1395	0.8271	0.8490	0.8373	0.1646
	CS	0.8720	0.8618	0.8662	0.1328	0.8288	0.8715	0.8486	0.1550
	HHO	0.8730	0.8504	0.8612	0.1369	0.8341	0.8497	0.8414	0.1598
	SCA	0.8687	0.8637	0.8652	0.1342	0.8257	0.8656	0.8436	0.1596
	JAYA	0.8721	0.8568	0.8639	0.1347	0.8286	0.8551	0.8406	0.1615
	DE	0.8750	0.8455	0.8589	0.1383	0.8348	0.8420	0.8369	0.1631
Vertebral	MVOSANN	0.8738	0.8802	0.8770	0.1634	0.9127	0.9147	0.9136	0.1223
	MVO	0.8732	0.8802	0.8767	0.1639	0.9096	0.9116	0.9105	0.1267
	GA	0.8400	0.8825	0.8600	0.1902	0.8598	0.8902	0.8742	0.1811
	FA	0.8635	0.8810	0.8720	0.1712	0.8901	0.9009	0.8953	0.1491
	SSA	0.8580	0.8933	0.8752	0.1686	0.8886	0.9182	0.9030	0.1396
	PSO	0.8805	0.8627	0.8700	0.1699	0.9027	0.8756	0.8868	0.1560
	GWO	0.8694	0.8832	0.8761	0.1654	0.9040	0.9138	0.9087	0.1299
	WOA	0.7950	0.8719	0.8303	0.2358	0.8262	0.8800	0.8513	0.2167
	CS	0.8402	0.8802	0.8578	0.1931	0.8662	0.8880	0.8753	0.1780
	HHO	0.8169	0.8756	0.8446	0.2134	0.8376	0.8813	0.8585	0.2053
	SCA	0.8167	0.8716	0.8403	0.2193	0.8349	0.8818	0.8557	0.2094
	JAYA	0.8320	0.8810	0.8518	0.2031	0.8581	0.8751	0.8633	0.1943
	DE	0.8394	0.8689	0.8508	0.2008	0.8636	0.8711	0.8643	0.1912

In Fig. 7 and Fig. 8, convergence curves are given to observe the convergence behaviour of the MVOSANN algorithm and to compare its performance with other algorithms. The figure shows that MVOSANN has the fastest convergence rate and the lowest MSE values in all datasets. MVOSANN algorithm has acquired powerful results by surpassing other algorithms in terms of robustness and stability.

The MVOSANN algorithm outclasses all other algorithms used in the comparing. To understand the relationship among the proposed algorithm and other algorithms, non-parametric Wilcoxon rank-sum statistical test was applied. The results of this test are shown in Table 7. The algorithm with the best average value gets the N/A value and is compared with other algorithms. According to table, the MVOSANN algorithm has a p value less than 0.05 for all data sets and for all algorithms, and it is seen to be statistically significant.

MLP training is a problem that has many local solutions and is also difficult to solve. As can be seen from the results obtained, the MVOSANN algorithm shows the best productivity against this difficult problem compared to the other 12 algorithms used for comparison. It has also proven this success in all the different search spaces of 12 different data sets. The reason for this success of MVOSANN is the high exploration - exploitation ability it has and the ability to avoid the local minimum.

Table 7
Wilcoxon rank sum test p -values (N/A = not applicable)

Datasets	Algorithms												
	MVOSANN	MVO	GA	FA	SSA	PSO	GWO	WOA	CS	HHO	SCA	JAYA	DE
Abalone	N/A	4.98E-04	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Balance	N/A	5.10E-05	2.87E-11	2.74E-10	7.03E-11	5.31E-08	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Blood	N/A	4.97E-03	2.87E-11	7.73E-10	1.04E-10	9.18E-07	9.44E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	5.23E-11	3.88E-11
Breast Cancer	N/A	1.54E-04	2.87E-11	3.88E-11	3.51E-11	4.29E-11	1.06E-08	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Diabetes	N/A	2.92E-04	2.87E-11	4.29E-11	2.87E-11	5.77E-11	6.37E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Diagnosis I	N/A	9.77E-04	2.87E-11	3.06E-09	5.23E-11	2.96E-03	1.15E-10	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Diagnosis II	N/A	6.97E-06	2.87E-11	1.86E-10	2.87E-11	0.054609	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Glass	N/A	1.41E-03	2.87E-11	4.29E-11	9.44E-11	2.87E-11	5.23E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Iris	N/A	3.26E-03	2.87E-11	8.93E-05	7.03E-11	8.70E-08	1.04E-10	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Liver	N/A	9.85E-06	2.87E-11	4.01E-10	2.87E-11	1.29E-05	3.88E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Raisin	N/A	4.97E-03	2.87E-11	4.00E-09	2.87E-11	4.22E-05	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11
Vertebral	N/A	5.39E-07	2.87E-11	2.74E-10	2.87E-11	1.63E-08	7.76E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11	2.87E-11

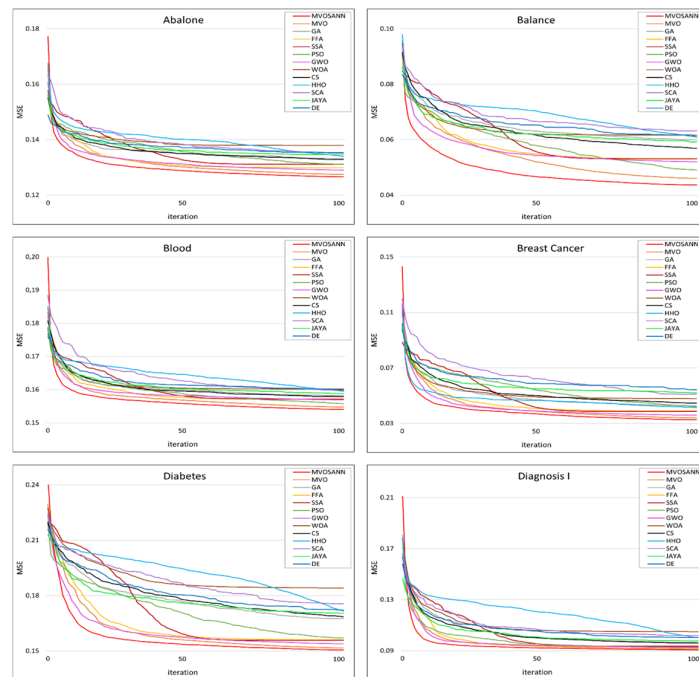


Fig. 7. Convergence curves of Abalone, Balance, Blood, Breast Cancer, Diabetes, Diagnosis I datasets.

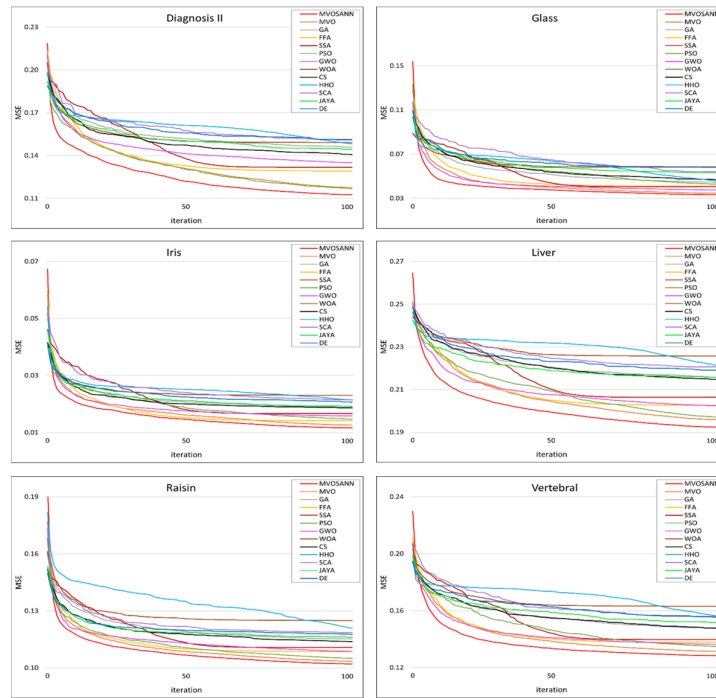


Fig. 8. Convergence curves of Diagnosis II, Glass, Iris, Liver, Raisin, Vertebral datasets.

6. Conclusion remarks

One of the problems met in implementations of ANNs is related to the training of the ANNs model. Training in ANNs is one of the determinants that directly affects the achievement of the model. In the studies, it is seen that many different techniques are used in ANNs training. Today, meta-heuristic algorithms are preferred over traditional algorithms during the training stage in ANNs models. Meta-heuristic algorithms appear to produce more effective results than traditional algorithms. An algorithm cannot be expected to perform highly against every problem. Therefore, it may be necessary to work on different algorithms for different problems or to develop existing algorithms.

In this article, the MVOSANN algorithm, which was developed based on the MVO and SA algorithms, is tested for the training of the feed forward MLP. The suggested algorithm (MVOSANN) has been used to optimize biases and weights in MLP for better results. The algorithm has been experimented on 12 datasets with different characteristics. The outcomes were compared with the outcomes of 12 various meta-heuristic algorithms. Experimental results suggest that the MVOSANN algorithm significantly enhances the performance of the MVO algorithm and carries out better than other algorithms. As a result, MVOSANN is an effective alternative for training ANNs.

In the future, larger and different datasets can be studied with the proposed MVOSANN algorithm. In addition, the MVOSANN algorithm can be experimented for the training of different model neural networks or real-world problems.

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