

Swarm Optimization with Neural Networks for Effective Classification Techniques

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Abstract:Swarm intelligence is a cooperative behavior of collective systems like insects such as ant colony optimization(ACO), fish schooling, birds flocking, bee Colony Optimization (BCO) particle swarm optimization (PSO) and so on. In this paper, a hybrid performance for data organization and information extrapolation is recommended. The Honey Bee Mating Optimization (HBMO) algorithm and Artificial Neural Networks may also be considered as a distinctive swarm-based optimization, in which the exploration algorithm is encouraged by the development of real honey-bee marital and mimic the iterative mating process of honey bees and approaches to select applicable drones for mating progression through the fitness function enrichment for selection of superlative weights for hidden layers of Neural Network classifiers. Extended HBMO with Neural Network algorithm is now realistic to classify the data proficiently by training the neural network. Extended HBMO (EHBMO-NN) procedure is now realistic to categorize the data efficiently by teaching the neural network. The arrangement precision of EHBMO-NN is associated with several other procedures. In this paper, extended honey-bee coupling optimization process (EHBMO-NN) is presented and verified with few benchmark instances. A developed way of Honey Bee Mating Optimization performance is joined with Neural Network which increases exactitude and decrease time interruption in complication of numerous factual world datasets.

Key words: Swarm Intelligence, Extended Honey Bee Mating Optimization, Fitness function, Effective Classification, Neural Network.

I. INTRODUCTION: -

The Honey Bee Mating Optimization (HBMO) procedure is a group-founded kind optimization performance, in which the examining method imitates the coupling development in honey-bee associations. Thus, the HBMO process is interrelated to the universal field of crowd aptitude, but the coupling progression which is based on boundary and metamorphosis mechanisms, strongly relate this process to evolutionary calculating too [1]. Fundamentals on the HBMO procedure are concisely pronounced forward, based on general models offered. A honey bee colony community a single queen-bee, some thousands of murmurs and numerous tens of thousands of employee-bees. The queen-bee is focussed in egg positioning and lives up to 5 or 6 years, although murmurs and employee-bees live no longer than 6 months. Murmurs, considered as forebears of the association, companion with the queen-bee and then die. They are haploid and act to transmission the genome hereditary from their mother to the next generation without shifting its genetic configuration, except through transmutations. The evolutionary segment of the procedure starts with the reproducing journey of the colony [2]. Throughout the copulating journey the queen mates with murmurs to form a chromosomal pool, also called spermatheca, which comprises of chromosomes received by the queen from murmurs [3]. The second phase of the evolutionary progression starts afterwards the genomic pond was filled with genes, and comprises in breeding eggs with genetic evidence from the spermatheca, based on boundary procedures between chromosomes. The final stage of the evolutionary process comprises in floating the broods based on the fitness function generated during the second stage, and creating a new generation of bees, based on transmutation operators [4].

In this work, University of California, Irvine machine learning databank is used. Through the investigation of prevailing technique, based on the proficient consequences, that

it sources Scalability disputes, absenteeism of precision and time depletion in instance of outsized datasets [5]. To experiment this concern, our work is concentrated on numerous approaches deliberated and to progress the fitness function assessment in extended honey bee mating optimization (EHBMO). Our objectives are to resolve convolution and scalability concerns in real world datasets and to recover the proficiency in data cataloguing. The weights are optimized through the evaluation of enhanced fitness function. The results obtained from the Extended honey bee mating optimization with Neural Network is to avoid scalability issues in large datasets, reduce the timeconsumption and also provide better accuracy and performance.

In our work, the University of California, Irvine (UCI) Machine Learning database is used. Several more researchers are used this databank for honey bee reproducing optimization performances such as Iris flower, wine, heart disease, cancer, diabetes and soya bean etc. The UCI datasets are used here to calculate the intermission, exactness and adeptness in countless data and sources of the data are dissimilar from each other. Since that time, it has been broadly used by scholars, educationalists, and scientists all over the world as a topmost source of machine learning data sets and it is universally known as standard datasets.

II. NEURAL NETWORKS: -

Neural Networks are known vibrant and quick system for productivity forecasting. MLP neural system is measured as the session of forage system which comprises numerous layers of computational units [6]. Each unit is a demonstration of neuron in which involves of a linear or non-linear stimulation function. The construction of the system is a absorbed and manifold layer graph and neurons in each layer is fully associated with the neurons of the subsequent layer [7].

The loads of the network are usually prepared unsystematically and are increasingly transformed reiteration during the training process learn a goal function. Training in such networks means that the network has to learn the goal function. To do so, each input composed with its equivalent output is presented to the network [8]. Learning algorithm tries to adjust the weight in all layers in a way that the error between calculated output and accurate output become small. The most known and popular learning process is Back Propagation [9]. The Learning algorithm is used to reduce the overall error of the network based on optimization method called gradient descent.

After vaccination of any input to primary layer of the network defined error function in the output layer. To regulate those weights which modifications continue until the alteration of weights of the first layer [10]. This progression is done for all proceedings of training data in each time. After finding the best restrictions (kind of activation function in hidden layer and number of hidden layer neurons) of the NN model, in alongside with the preparation of the network in each repetition,

III. HBMO COMBINED WITH NEURAL NETWORK (NN)

HBMO-NN algorithm was exploited to elevate the masses of the system. The loads are optimized by the estimation of suitability function. The HBMO algorithm is functioned based on bee's auxiliary with crossover and mutation operator. The replacement of bees is done by fitness calculated after boundary and metamorphosis processes. So, the suitability function of HBMO is extended as EHBMO. The HBMO consists of F0 and F1 as Fitness Function whereas in EHBMO comprises additional two functions of F2, F3.

Fitness F_0 - is the sum of Euclidean remoteness of employee bees to its drone and drone to the queen bee. Fitness 1-is the ratio of the average dynamism sustainability of employee bees with its murmur. Fitness 2-is the ratio of the average Euclidean distance of the murmur to the queen with the sum of Euclidean distance of all the employee bees to the queen. Fitness 3-is the input particle which is strained with threshold significance of the employee bees and murmur, we use a better-quality way of Bee optimization technique known as Extended Honey Bee Mating Optimization is combined with Neural Network to form hybrid algorithms named as (EHBMO-NN) which improves accuracy and reduce time delay in complexity of numerous grounds.

- The HBMO process initiated with the coupling-flight, where a queen (best solution) chooses murmur probabilistically to arrangement the spermathecal.
- A murmur then nominated from the list at unsystematic for the formation of children. Creation of number of new broods by cross- overing the drones "genotypes with the empress's".
- Use of employees to conduct local examine on young bees. Reworking of employee's suitability based on the quantity of perfection accomplished on broods. Replacement of feebler queens by righter broods.

The procedure begins with three user-defined constraints and one predefined parameter. The predefined parameter is the number of employees, demonstrating the number of heuristics prearranged in the package. The three user-defined parameters are the quantity of monarchs, the queen's spermatheca size and the number of young's that is born by all queens.

HBMO-Neural Networks Algorithm

Step 1: The contribution neuron n is feed with teaching data xm with preferred target yn .

Step 2: The input preparation data xm undertakes recapitulation process. In each reiteration, the weightage of each node $w(n)$ is calculated.

Step 3: The bias or mistake amount of each node is considered as delta function.

Step 4: The heaviness of each node $w(n)$ is calculated based on the delta node and the input data xm , the $w(n)$ characterizes weights of connections between network input xm and neuron n in input layer, and the Symbols yn represents output signal of neuron n .

Step 5: The weight $w(n)$ is accustomed in the concealed layer by Extended Honey Bee Mating Optimization process.

Step 6: The target production is investigated and endures back dissemination system to reach bias reduced operative output.

Where f_0 is the sum of Euclidean distances of worker bees to its drone and drone to the Queen Bee, Br is a Replacement of Bees in the current round, αl ($l = 1, \beta$) is the number of worker bees, β is the number of clusters, $dw(b)$, d is the Euclidean distance from worker bee i in cluster j to its drone, $d_{d, Q(B)}$ is the Euclidean distance from j th drone to the Queen bee. Function f_1 is the ratio of the average energy of worker bees with its drone. Function f_2 is the ratio of the average Euclidean distance of the drone to the $Q(b)$ with the sum of Euclidean distance of all the worker bees to the Queen Bee. Function f_3 is the input particle is filtered with threshold value of the worker bees ($\alpha i \dots n$) and drone ($\beta 1 \dots n$), So that the worker bees are eliminated based on this threshold value which regains minimum iteration and Energy Efficiency.

The constants A, B, C, D are predefined constants used to weight the contribution of each of the sub-objectives and $A + B + C + D = 1$. The fitness function defined above has the objective of simultaneously minimizing the intra-cluster distance between worker bees and drone, as quantified by f_0 and of maximizing the cluster head's energy in its cluster as quantified by f_1 ; and of producing cluster with unequal size as quantified by f_2 ; and also, of optimizing the energy dissipation in the clusters as quantified by f_3 . According to the fitness function, a small value of f_0 , f_1 suggests compact clusters with the optimum set of worker bees that have sufficient energy to perform the drone tasks. A small value of f_2 means that the size of the clusters located closer to the base station is smaller. A small value of f_3 shows that the formed clusters are more energy efficient.

IV. EXPERIMENTAL RESULT AND DISCUSSION: -

Table-1 Input Dataset Statistics:

Dataset Used	Iris	Liver	Cancer	Diabetes	Arrhythmia
No of Instances	150	345	32	768	452
No of Classes	3	2	2	2	16
No of Attribute	5	78	57	9	280

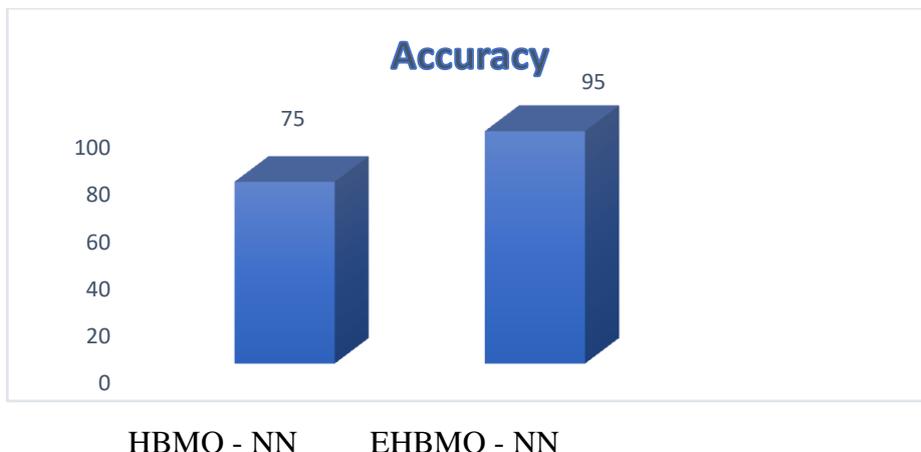
Table-2 : This Table Represents the Data Analysis using HBMO-ANN Algorithm with the Standard Repository Dataset.

HBMO-NN Statistical Data Analysis									
Dataset Used	TP Rate	FP Rate	Precision	Recall	F-Measure	MCC	ROC Area	Accuracy	No. of Classes
Iris	0.987	0.011	0.968	0.944	0.881	0.91	1.005	1.000547	3
Liver	0.739	0.281	0.77	0.739	0.711	0.405	0.751	1.052125	2
Cancer	0.69	0.24	0.738	0.5	0.726	0.315	0.66	1.028441	2

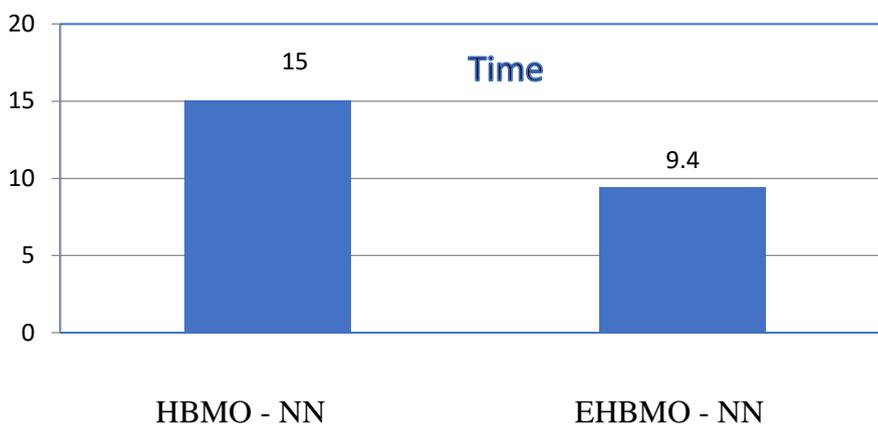
Table-3 : This Table Represents the Data Analysis using EHBMO-NN Algorithm with the Standard Repository Dataset.

EHBMO- NN Statistical Data Analysis									
Dataset Used	TP Rate	FP Rate	Precision	Recall	Measure	MCC	ROC Area	Accuracy	No. of Classes
Iris	0.989	0.004	1.008	0.987	0.989	0.97	1.087	1.001585	3
Liver	0.797	0.214	0.89	0.799	0.788	0.469	0.784	1.065278	2

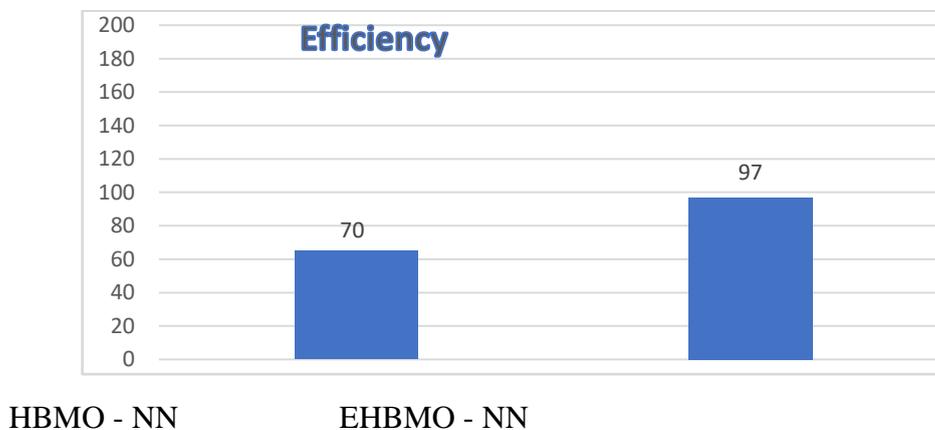
Cancer	0.9	0.4	0.789	0.8	0.768	0.367	0.81	1.050737	2
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The Figure-1 illustrates the Accuracy of HBMO-NN is 75% and EHBMO-NN is 95%



The Figure-2 illustrates the Time of HBMO-NN is 15 min and EHBMO-NN is 9.4 min



The Figure-3 illustrates the Performance Efficiency of HBMO-NN is 70% and EHBMO-NN is 97 %

V. CONCLUSION:

In this research, first check the efficiency of HBMO-NN in Data Cataloguing tasks, based on the attained results, the HBMO-NN causes Scalability disputes in case of outsized Datasets, to challenge this Scalability issue, the research is focus based on alternates scrutinised, and committed to progress the fitness function assessment in ExtendedHBMO-NN. Our objectives are to answer complication and scalability issues in real world datasets and to expand competence in data extrapolation. We executed and matched this EHBMO-NN with the HBMO-NN, from the results, it concludes that EHBMO-NN can obtain competitive results against the real-world data sets used, although there is some growth in the computational effort needed. Guidelines for forthcoming work include examination and manipulation by relating this tool to more challenging data sources comprising continuous qualities.

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