



A Discrete Moth Flame Algorithm for Feature Selection

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Abstract: A crucial and important task in machine learning is feature selection (FS). The primary goal of the feature selection task is to minimize the dimension of the feature set while preserving performance accuracy. In order to address the FS task, a discrete moth flame algorithm that is combined with levy flights (DL-MFA) is presented in this research. The proposed DL-MFA imitates the natural navigational patterns of moths. The moths move along a straight line at a constant angle in the direction of the true light source (the moon) known as transverse orientation. Additionally, moths are drawn to artificial lights like fires and because of the close proximity; they constantly adjust their flying angles, creating a spiral path. In order to maintain healthy population diversity and increase the global search capabilities of the algorithm, the levy flight search technique is also used as a regulator of the moth position updating mechanism. The five swarm intelligence algorithms (SIAs) are contrasted with the proposed algorithm using measures such as entropy, purity, completeness score (CS), and homogeneity score (HS). For evaluating fitness, the SSE fitness function is utilised. The outcomes have shown that the proposed algorithm achieved purity values in the range 90% to 100%, and entropy 10% to 50%. Proposed DL-MFA has also achieved homogeneity score and completeness score up to 50%. These results prove that the proposed algorithm is better than its state-of-the-art competitors.

Keywords: Feature selection, Levy flight, Moth flame algorithm, Swarm intelligence, Classification.

1. Introduction

The process of choosing the most crucial features to feed into machine learning algorithms is known as feature selection (FS) [1]. Feature selection (FS) eliminates the majority of the dataset's pointless features, which simplifies our machine learning model. Additionally, it decreases the problem's dimensionality while increasing prediction accuracy and processing efficiency. Filter methods, wrapper methods, and embedded methods are the three categories of feature selection techniques, and they are detailed below.

Filter methods: These methods forecast the relationship between the features and our aim using some statistical techniques. Each feature receives a score based on how essential it is. This score is used by filter algorithms to determine if a characteristic is

pertinent or not. There is no machine learning algorithm used in this strategy.

Wrapper Methods: Using the classifier's performance evaluation, wrapper methods rate the subset of features' quality. Here, a feature subset selection machine learning approach on a given dataset is applied. A greedy technique is used to assess all potential feature combinations against the evaluation criterion. When compared to filter approaches, the wrapper feature selection typically produces good prediction accuracy.

Embedded Methods: Methods that incorporate feature selection into the learning algorithm are referred to as embedded methods. The feature selection and classification operations are carried out simultaneously by embedded algorithms. Typically, during training, the characteristics that have contributed the most across all iterations are extracted. Examples of embedded approaches

include feature selection using decision trees and random forests.

The filter methods ignore the interaction with the classifier and each feature is considered independently thus ignoring feature dependencies. In addition, it is not clear how to determine the threshold point for rankings. Wrapper methods are computationally expensive as most of the execution time is spent in training the predictor. They are also prone to overfitting. Embedded methods suffer from the loss of a large part of the information contained in the data set due to the elimination of most of the features. These methods also ignore interactions and correlations between variables.

Generally speaking, any FS problem is an NP-hard problem. The solutions to this problem are found via optimization algorithms. Most of the optimization algorithms have the ability to address the problems faced by the different FS methods mentioned above.

Swarm intelligence algorithms (SIAs) are one of the types of the optimization algorithms. SIAs are based on the collective intelligence shown by the objects like animals, insects and others. A number of SIAs are proposed to solve the optimization algorithms. An SIA that is based on the natural navigational patterns of moths is called the moth-flame algorithm (MFA) [2]. We have proposed a discrete version of the moth flame algorithm (DL-MFA) hybridized with levy flights in this paper. We have applied this discrete MFA to feature selection (FS) problem.

Below we emphasize this study's contribution.

1. The introduction of a swarm intelligence algorithm for feature selection based on MFA.
2. The newly presented algorithm incorporates the ideas of levy flights.
3. Twelve medical datasets that are available at UCI are used to evaluate the introduced algorithm.
4. Five cutting-edge SIAs are contrasted with the outcomes produced by the newly introduced MFA.
5. The effectiveness of the newly implemented algorithm is assessed using four external evaluation measures.
6. Based on the evaluation, it is determined that the introduced algorithm's time, convergence, and solution quality are acceptable.

There are seven sections in the paper. Section two presents a review of the literature. Section three provides an overview of the fundamental moth flame algorithm. Section four and five presents levy flights and Proposed DL-MFA. Fitness function is discussed in section 6. The results and their

comparison with other known SIAs are covered in section seven. Conclusion is followed section seven.

2. Related work

In this research, we propose a discrete moth flame algorithm (DL-MFA) that is integrated with levy flights. DL-MFA is a swarm based algorithm (SIA). A brief review of the literature based on the SIAs for feature selection (FS) is as follows.

Xie et al. [3] tried to overcome the poor exploitation and premature convergence of particle swarm optimization (PSO) by presenting two new variants of PSO. In the first variant they integrated global best signals, rectified personal, swarm leader enhancement with Gaussian distribution, local exploitation using spiral search, mutation operations, and mirroring for solution improvement. The authors tried to improve the first approach by using search coefficients, scattering schemes, adaptive breeding mechanism, and multiple optimal signals. A unique binary gray wolf algorithm (BGWO) is introduced by Emery et al. [4] for FS. Hu et al. [5] introduced an enhanced update equation for balancing the exploration and exploitation in the searching. The introduced method gave some new transfer functions to be used in the new update equation. A grey wolf algorithm with multiple search strategies (MEGWO) was introduced by Q. Tu et al. [6]. Guo et al. hybridized the whale optimization algorithm (WOA) with new mutation and adaptive neighborhoods strategies for the FS problem. Using this, the algorithm chooses solutions from the neighborhoods of the current best solution. The new mutation strategy helps the algorithm balance exploration and exploitation to overcome the local optima problem. R. K. Agrawal et al. [7] integrated the quantum concept with a whale optimization algorithm (QWOA) to improve the convergence of the basic WOA for the FS problem. Chantar et al. [8] have integrated a simulated annealing (SA) algorithm with a binary dragonfly algorithm (BDA) to improve the classification accuracy of BDA in feature selection. The best solution found with the BDA was given to SA for further improvement of the search results. Ibrahim et al. [9] improved the social spider algorithm (SSO) by using opposition-based learning to increase the search area's exploration. They did this to save the SSO from falling into the local optima.

Hichem et al. [10] presented a binary grasshopper algorithm (BGHO) for the feature selection problem. The results obtained are compared with the other five known approaches, and BGHO has proved itself the best among all of

them. Ahmed and colleagues [11] improved SSA by integrating a new local search and a method for re-positioning the search agents (Sparrows) into the search space that are wandering beyond the search space. This improvement was carried out to enhance the searching efficiency of the original SSA. Arora et al. presented [12] two variants of the butterfly algorithm (BBOA) by using V-shaped and S-shaped transfer functions, respectively. The authors tested both variants with the 21 different UCI datasets.

Wang et al. [13] improved the cuckoo search algorithm by integrating the chaotic maps, two population preservation strategies, levy flights, and a mutation strategy. Initially, chaotic maps are used to improve the initialization diversity of the algorithm to avoid local optima. Population preservation strategies are used in the next step to select the fittest feature from each iteration. Finally, levy flight is applied with the new mutation strategy to avoid convergence issues while working with a large search space. Naseer et al. [14] presented a hybridized filter-based feature selection algorithm in which ACO is combined with the gain ratio. A gain ratio is used here to normalize preferences between information gain and mutual information. The gain ratio penalizes some high-split information as a part of the classifier and uses it over different convergence thresholds for final feature subset selection. Hu et al. presented [15] an improved grey wolf optimizer for feature selection. They have analyzed the A and D parameters, which are controlled by parameter a in the position updating. These parameters influence the exploration and exploitation process. A new position update function for balancing global and local search is proposed by analyzing the range of values of A and D under binary conditions.

Zawbaa et al. [16] presented a chaotic antlion optimizer integrated with random walks and a new controlling parameter for balancing exploration of the search space and exploitation of the best solutions. The parameter I is used to control the range of the random walks. The chaotic approach is used in this algorithm to improve the tradeoff between exploitation and exploration. Too et al. [17] presented two variants of binary harris hawk algorithms which use different types of transfer functions for converting a continuous version into a discrete one. Hegazy et al. [18] improved the convergence rate, consistency, and accuracy of the salp swarm algorithm by using a new control parameter, inertia weight. They have combined this new algorithm with the KNN classifier for feature selection. Sureja et al. [19] improved the shuffle frog leaping algorithm (SFLA) by integrating

simulating annealing (SA) with it. SA is used to exploit more and more near-optimal solutions for the enhancement of the solution quality. Uzer et al. [20] improved classification accuracy by combining the artificial bee colony (ABC) algorithm with support vector machines. A 10-fold cross-validation is applied to obtain the classification accuracy of the proposed approach. Salima et al. [21] presented an improved version of a wrapper-based crow search algorithm (CSA) to extract the finest feature subsets. They made improvements by integrating adaptive awareness probability to enhance the balance between exploration and exploitation, selecting crow by the dynamic local neighborhoods to follow, and by developing new searching techniques to improve global exploration.

All of the above work suffers from the one or more drawbacks like computational complexity, execution time, stuck in local optima, loss of information, and noise.

There is a scope to propose a new, modified, and hybrid algorithm for feature selection (FS) problem based on no free lunch (NFL) theorem. This motivates us to modify and present a discrete version of the moth flame algorithm (DL-MFA) for feature selection (FS). The performance of any optimization algorithm depends on how efficiently it explores and exploit the search space and how quickly finds global optimal solution without getting trapped in local optimal solution. So, we have hybridized levy flights with DL-MFA to improve the exploration and exploitation of the search space.

3. Basic moth flame algorithm

The moth flame algorithm (MFA) is a population based approach [2]. MFA imitates moths' natural navigational strategies. The moths move in a straight line at a constant angle in the direction of the moon. Transverse orientation is the term used to describe this navigation. Typically, artificial lighting, such as flames, has a strong attraction to moths. Due to the near proximity, moths continuously alter their flight angle, which causes them to spiral. In order to solve NP-hard problems, the MFA algorithm simulates the aforementioned moth behaviors.

The set of moths is represented by a matrix M in the fundamental moth flame algorithm [2]. The fitness value of each moth in the population is kept in an array called OM . The flames are represented by a matrix F , which is very similar to the moths. The fitness value of each flame is kept in an array OF [2, 22].

The Moth Flame algorithm generates a three-tuple that represents an approximation of the global

ideal for every problem. It is made in accordance with Eq (1).

$$MFO = (I, P, T) \quad (1)$$

Here I represent a function that is used to generate a population of moths randomly with the respective values of fitness. The function can be modeled methodically as under.

$$I: \emptyset \rightarrow \{M, OM\} \quad (2)$$

The P function is developed for moving the moths about the search region. The updated matrix M of moths is returned by the P function in the output.

$$P: M \rightarrow M \quad (3)$$

If the termination criterion is satisfied then T function returns true and false otherwise.

$$T: M \rightarrow \{true, false\} \quad (4)$$

The function P performs itself repeatedly until it produces a true value. We update each moth's position in relation to the flames for mathematically simulating moth behavior using Eq. (5).

$$M_i = S(M_i, F_j) \quad (5)$$

Here, M_i and F_j represents i th moth and j th flame respectively. S is the spiral function.

We follow the following conditions to use any type of spirals.

1. Both, starting and ending points for the spiral would be a moth.
2. The position of the flame should be the ending point of the spiral.
3. Variation in the range of the spiral would be restricted to the search area.

For a moth flame algorithm, we can define a logarithmic spiral by considering the above conditions as per Eq. (6).

$$S(M_i, F_j) = D_i \cdot e^{bt} \cdot \cos(2\pi t) + F_j \quad (6)$$

Here, D_i is the distance between i th moth and j th flame, b is a constant defines spiral's shape, and t represents a number (random) in $[-1, 1]$.

Now we calculate D using Eq. (7).

$$D_i = |F_j - M_i| \quad (7)$$

Here, M_i and F_j are i th moth and j th flame, and D_i is the distance between i th moth and j th flame.

Eq. (6) is used to determine a moth's next position in relation to a flame. The t parameter in Eq. (6) determines how close the moth will be to the flame in its subsequent position. (The closest is represented by $t = -1$, and the farthest by $t = 1$).

Only moths travelling in the direction of the flame are considered position updates according to Eq. (6) MFA could become swiftly trapped in local optima under this circumstance. The exploitation of the most useful solutions is equally decreased when n distinct locations are used for moths' position updates. Eq. (8) is used to solve this problem.

$$flame\ no = round\left(N - l \times \frac{N-1}{T}\right) \quad (8)$$

Here, l is a current iteration, N is maximum flames and T represents maximum iterations. The basic moth flame algorithm is given in algorithm 1.

4. Levy flight

The concepts of Levy-flight were introduced by Paul Levy in 1937 [23]. Levy-flight is a random walk with the particular heavy jumps using step lengths which are derived from a probability distribution. We can define term flight as maximum distance in a straight line between two points that an entity in motion covers without directional variation.

Levy flight is integrated with basic MFA to improve the diversity of the population of moths to jump out of the local optima. Specifically, levy

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Initialize moths ( $M$ ) randomly in dimension ( $D$ )
While ( $Iter \leq Maxiter$ )
  Perform Updating of flame number using Eq. (8)
   $OM =$  Fitness Function ( $M$ );
  if  $Iter == 1$ 
     $F =$  sorting ( $M$ );
     $OF =$  sorting( $OM$ );
  else
     $F =$  sorting( $M_t - 1, M_t$ );
     $OF =$  sorting ( $M_{t-1}, M_t$ );
  end
  for  $i = 1 : n$ 
    for  $j = 1 : d$ 
      • Perform Updating of  $r$  and  $t$ 
      • Compute  $D$  using Eq. (7) concerning the respective moth
      • Updating  $M(i, j)$  using Eq. (5) and Eq. (6) concerning the respective moth
    end
  end

```

Figure. 1 Basic MFA algorithm

flights are composed of clusters of multiple short steps connected by longer relocations.

5. Proposed moth flame algorithm

To solve the local optima problem, this research suggests the discrete levy flight moth-flame algorithm (DL-MFA). Additionally, we work to increase the diversity of the moth population. Excellent qualities of Levy flight contribute to increasing population diversity. Due to this, the suggested method can very easily jump out from the local optimum. It is also possible to achieve a good balance between the MFA's exploitation and exploration capabilities. Therefore, following the update of positions [24, 25], we permit each moth to perform its flight in accordance with Eq. (9).

$$X_i^{t+1} = X_i^t + u \text{ sign}[\text{rand} - 0.5] \oplus \text{levy}(\beta) \quad (9)$$

Here, X_i^t is a solution vector, u and rand are uniformly distributed random numbers, and \oplus represents entry-wise multiplications.

There are only three possible values for the signed random integer u (rand): 1, 0, and -1. Levy-flight and u are combined to improve the random walks of the moth, which then aids the MFA in avoiding local maxima. According to Eq. (10) [24, 25], levy flight is considered to be a random walk in which the step lengths determine the steps and a levy distribution determines the leaps. Levy random numbers are provided by Eq. (11).

$$\text{levy}(\beta) \sim \mu = t^{-1-\beta}, (0 \leq 2) \quad (10)$$

$$\text{levy}(\beta) \sim \frac{\emptyset \times \mu}{|v|^{1/\beta}} \quad (11)$$

Here, u and v are normal distributions, Γ is a gamma function, and $\beta=1.5$. \emptyset is defined using Eq. (12).

$$\emptyset = \left[\frac{\Gamma(1+\beta) \times \sin(\pi \times \frac{\beta}{2})}{\Gamma(\frac{1+\beta}{2}) \times \beta \times 2^{\frac{\beta-1}{2}}} \right]^{1/\beta} \quad (12)$$

6. Fitness function

To assess the fitness of the suggested DL MFA solutions, we employ a fitness function. In this study, evaluation is conducted using the Sum of Squared Error (SSE) [26-28] fitness function provided in Eq. (13).

$$SSE = \sum_{n=1}^N \text{dist}_{nc}^2 \quad (13)$$

```

Initialize moths ( $M$ ) randomly in dimension ( $D$ )
While ( $Iter <= Maxiter$ )
  Perform Updating of flame number using Eq. (8)
   $OM =$  Fitness Function ( $M$ );
  If  $Iter == 1$ 
     $F =$  sorting ( $M$ );
     $OF =$  sorting ( $OM$ );
  else
     $F =$  sorting ( $M_{t-1}, M_t$ );
     $OF =$  sorting ( $M_{t-1}, M_t$ );
  end
  for  $i = 1 : n$ 
    for  $j = 1 : d$ 
      • Perform Updating of  $r$  and  $t$ , and Compute  $D$ 
        using Eq. (7) concerning the respective moth
      • Update ( $i, j$ ) using Eq. (5) and Eq. (6) concerning
        the respective moth
    end
  for each moth (search agent)
    • Perform Updating the position of the current
      moth (search agent) using Levy-flight
  end
   $Iter = Iter + 1$ ;
End

```

Figure. 2 Proposed DL-MFA algorithm

$Dist_{nc}$ in this case refers to the (Euclidean) distance between a point and the centroid. By reducing the sum of the squared error (SSE) function, we can get the improved outcomes.

7. Results and discussions

The effectiveness of the DL-MFA algorithm in feature selection is examined in this section. To make an accurate comparison, we present the outcomes of utilizing the DL-MFA method on 12 well-known medical data sets. A fitness function sum of squared error (SSE) is also employed to demonstrate the algorithm's strength. We also compare the DL-MFA to five other popular algorithms that have been applied to feature selection in the past: BDASA [8], OBSSO [9], BGHO [10], IBSSA [11], and BBOA [12].

7.1 Datasets and environment

A computer system with an Intel i3 processor, 8 GB of RAM, and the 64-bit Windows 10 operating system is used to carry out the research. Table 2 displays the parameter settings for the proposed DL-MFA. Twelve medical data sets were utilized [29] to evaluate the proposed MFA in Table 1 based on their characteristics.

7.2 Evaluation measures

Homogeneity score (HS), purity, completeness

Table 1. Data set properties

| Dataset | Instances | Attributes |
|---------------|-----------|------------|
| Exactly | 1000 | 13 |
| KrVsKpEW | 3196 | 36 |
| M-of-N | 1000 | 13 |
| Vote | 300 | 16 |
| BrestEW | 569 | 30 |
| CongressEW | 435 | 16 |
| Lymphography | 148 | 18 |
| Tic-tac-Toe | 958 | 9 |
| PenglungEW | 73 | 325 |
| Breast cancer | 569 | 569 |
| SpectEW | 267 | 22 |
| WaveformEW | 5000 | 40 |

score (CS), and entropy measures are used to evaluate the performance of the DL-MFA [26, 30, 31]. The entropy is calculated Using Eq. (14) [26, 31], and provides information about the distribution of the semantic classes inside the cluster.

$$Entropy = \sum_{j=1}^k \frac{|P_j|}{n} E(P_j) \tag{14}$$

Here, $E(P_j)$ represents the individual cluster entropy.

$$E(P_j) = \frac{1}{\log k} \sum_{i=1}^k \frac{|P_j \cap T_i|}{P_j} \log \left(\frac{|P_j \cap T_i|}{P_j} \right) \tag{15}$$

The purity is computed using Eq. (16) [26, 31].

$$Purity = \frac{1}{n} \sum_{j=1}^k \max_i (|T_i \cap P_j|) \tag{16}$$

Here, P_j represents points allotted to cluster j , k represents all clusters, and T_i represents points actually allotted to cluster I .

We compute homogeneity score (HS) using Eq. (17) [26, 30, 31].

$$HS = 1 - \frac{H(T|P)}{H(T)} \tag{17}$$

Here $H(T)$ and $H(T|P)$ are entropy and conditional entropy of the classes and computed using Eqs. (18) and (19).

$$H(T) = - \sum_{t=1}^{|T|} \frac{n_t}{N} \cdot \log \left(\frac{n_t}{N} \right) \tag{18}$$

$$H(T|P) = - \sum_{p=1}^{|P|} \sum_{t=1}^{|T|} \frac{n_{pt}}{N} \log \left(\frac{n_{pt}}{n_p} \right) \tag{19}$$

Here, p is the anticipated cluster, while n_t and n_p stand for the points that a true class t holds. Additionally, n_{pt} denotes the quantity of points that

are grouped together into a true class (t) of a predicted cluster (p). Eq. (20) is used to get the completeness score (CS) [26, 30, 31].

$$CS = 1 - \frac{H(P/T)}{H(P)} \tag{20}$$

Table 2. Experimental settings of DL-MFA

| Parameters | Value |
|--------------------------|-------|
| Population size of moths | 40 |
| Maximum Iterations | 500 |
| Search agents | 50 |

Table 3. Comparison of results (Brest Cancer)

| Criteria | DL-MFA | OB SSO | BDA -SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|---------|-------|-------|--------|
| SSE | | | | | | |
| Best | 209 | 374 | 216 | 331 | 274 | 359 |
| Worst | 699 | 695 | 670 | 653 | 647 | 742 |
| Avg | 321 | 429 | 253 | 378 | 384 | 496 |
| Std. | 0.3 | 0.3 | 0.3 | 0.3 | 0.3 | 0.3 |
| Evaluation Measures | | | | | | |
| Purity | 1.0 | 0.8 | 0.9 | 0.8 | 0.9 | 0.8 |
| Entropy | 0.1 | 0.2 | 0.1 | 0.1 | 0.1 | 0.1 |
| HS | 0.8 | 0.6 | 0.7 | 0.8 | 0.7 | 0.7 |
| CS | 0.8 | 0.6 | 0.7 | 0.8 | 0.7 | 0.7 |
| Time | 30.0 | 37.0 | 31.4 | 31.2 | 55.4 | 69.9 |

Table 4. Comparison of results (BrestEW)

| Criteria | DL-MFA | OB SSO | BDA -SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|---------|-------|-------|--------|
| SSE | | | | | | |
| Best | 292 | 578 | 400 | 630 | 610 | 887 |
| Worst | 1622 | 1773 | 1187 | 1537 | 1610 | 1716 |
| Avg | 653 | 675 | 500 | 700 | 1094 | 1178 |
| Std. | 0.3 | 0.2 | 0.2 | 0.2 | 0.3 | 0.2 |
| Evaluation Measures | | | | | | |
| Purity | 0.9 | 0.8 | 0.6 | 0.7 | 0.6 | 0.6 |
| Entropy | 0.5 | 0.6 | 0.6 | 0.5 | 0.6 | 0.6 |
| HS | 0.2 | 0.0 | 0.0 | 0.2 | 0.0 | 0.0 |
| CS | 0.4 | 0.1 | 0.1 | 0.3 | 0.2 | 0.1 |
| Time | 34.2 | 43.8 | 38.3 | 37.4 | 60.7 | 70.4 |

Table 5. Comparison of results (CongressEW)

| Criteria | DL-MFA | OB SSO | BDA -SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|---------|-------|-------|--------|
| SSE | | | | | | |
| Best | 1010 | 1222 | 1116 | 1429 | 1067 | 1254 |
| Worst | 1685 | 1756 | 1620 | 1685 | 1767 | 1743 |
| Avg | 1181 | 1296 | 1171 | 1477 | 1300 | 1469 |
| Std. | 0.4 | 0.3 | 0.3 | 0.3 | 0.4 | 0.3 |
| Evaluation Measures | | | | | | |
| Purity | 0.9 | 0.8 | 0.9 | 0.7 | 0.9 | 0.8 |
| Entropy | 0.2 | 0.3 | 0.3 | 0.3 | 0.3 | 0.3 |
| HS | 0.6 | 0.5 | 0.5 | 0.5 | 0.5 | 0.5 |
| CS | 0.7 | 0.5 | 0.5 | 0.5 | 0.5 | 0.5 |
| Time | 17.9 | 20.5 | 17.8 | 17.7 | 29.2 | 34.9 |

Here, $H(P)$ and $H(P|T)$ stand for the entropy and the conditional entropy of the clusters and computed using Eqs. (21) and (22) [26, 30, 31].

$$H(P) = -\sum_{p=1}^{|P|} \frac{n_p}{N} \cdot \log\left(\frac{n_p}{N}\right) \quad (21)$$

$$H(P|T) = -\sum_{t=1}^{|T|} \sum_{p=1}^{|P|} \frac{n_{pt}}{N} \log\left(\frac{n_{pt}}{n_p}\right) \quad (22)$$

Table 6. Comparison of results (Exactly)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| SSE | | | | | | |
| Best | 3044 | 3097 | 3123 | 3126 | 3051 | 3147 |
| Worst | 3191 | 3437 | 3432 | 3373 | 3420 | 3437 |
| Avg | 3009 | 3131 | 3163 | 3207 | 3166 | 3212 |
| Std. | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 |
| Evaluation Measures | | | | | | |
| Purity | 1.0 | 0.9 | 0.9 | 0.9 | 1.0 | 0.9 |
| Entropy | 0.5 | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 |
| HS | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 |
| CS | 0.7 | 0.7 | 0.6 | 0.6 | 0.7 | 0.6 |
| Time | 51.9 | 51.8 | 59.6 | 53.3 | 52.6 | 60.7 |

Table 7. Comparison of results (KrVsKpEW)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| SSE | | | | | | |
| Best | 4083 | 6839 | 5097 | 5671 | 4565 | 7099 |
| Worst | 1172 | 1044 | 1145 | 1042 | 7605 | 1106 |
| | 1 | 9 | 2 | 5 | | 7 |
| Avg | 6714 | 8463 | 5784 | 6318 | 4907 | 8477 |
| Std. | 0.2 | 0.2 | 0.1 | 0.1 | 0.2 | 0.2 |
| Evaluation Measures | | | | | | |
| Purity | 0.9 | 0.7 | 0.8 | 0.8 | 0.9 | 0.7 |
| Entropy | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 |
| HS | 0.2 | 0.1 | 0.1 | 0.1 | 0.2 | 0.3 |
| CS | 0.2 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 |
| Time | 703 | 2719 | 781 | 609 | 1120 | 1297 |

Table 8. Comparison of results (Lymphography)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| Best | 127 | 153 | 157 | 159 | 131 | 200 |
| Worst | 290 | 278 | 276 | 260 | 232 | 288 |
| Avg | 168 | 168 | 202 | 169 | 141 | 235 |
| Std. | 0.2 | 0.2 | 0.2 | 0.1 | 0.2 | 0.2 |
| Evaluation Measures | | | | | | |
| Purity | 0.9 | 0.8 | 0.8 | 0.8 | 0.8 | 0.7 |
| Entropy | 0.5 | 0.5 | 0.5 | 0.5 | 0.5 | 0.5 |
| HS | 0.3 | 0.1 | 0.1 | 0.2 | 0.1 | 0.1 |
| CS | 0.5 | 0.3 | 0.4 | 0.3 | 0.4 | 0.2 |
| Time | 6.1 | 6.6 | 6.2 | 7.2 | 11.3 | 11.9 |

Table 9. Comparison of results (M-of-N)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| SSE | | | | | | |
| Best | 3033 | 3127 | 3107 | 3143 | 3074 | 3162 |
| Worst | 3474 | 3406 | 3414 | 3462 | 3161 | 3379 |
| Avg | 3170 | 3201 | 3141 | 3171 | 3027 | 3224 |
| Std. | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 |
| Evaluation Measures | | | | | | |
| Purity | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 |
| Entropy | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 |
| HS | 0.2 | 0.1 | 0.2 | 0.1 | 0.2 | 0.0 |
| CS | 0.3 | 0.1 | 0.2 | 0.1 | 0.3 | 0.1 |
| Time | 52.4 | 52.7 | 52.0 | 52.7 | 87.9 | 105.2 |

Table 10. Comparison of results (PenglungEW)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| SSE | | | | | | |
| Best | 2457 | 3337 | 3096 | 3755 | 2612 | 4050 |
| Worst | 4137 | 4095 | 3995 | 4131 | 3004 | 4090 |
| Avg | 2620 | 3387 | 3366 | 4008 | 2585 | 3996 |
| Std. | 0.4 | 0.4 | 0.5 | 0.3 | 0.5 | 0.3 |
| Evaluation Measures | | | | | | |
| Purity | 0.6 | 0.5 | 0.5 | 0.5 | 0.5 | 0.5 |
| Entropy | 0.5 | 0.6 | 0.5 | 0.6 | 0.6 | 0.6 |
| HS | 0.4 | 0.3 | 0.4 | 0.3 | 0.3 | 0.3 |
| CS | 0.4 | 0.5 | 0.5 | 0.4 | 0.5 | 0.3 |
| Time | 11.9 | 7.1 | 6.6 | 22.9 | 9.6 | 11.3 |

Table 11. Comparison of results (SpectEW)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| SSE | | | | | | |
| Best | 1035 | 1193 | 1162 | 1198 | 1149 | 1244 |
| Worst | 1471 | 1461 | 1485 | 1467 | 1412 | 1480 |
| Avg | 1170 | 1228 | 1270 | 1222 | 1164 | 1345 |
| Std. | 0.5 | 0.4 | 0.5 | 0.5 | 0.4 | 0.3 |
| Evaluation Measures | | | | | | |
| Purity | 0.8 | 0.8 | 0.8 | 0.8 | 0.8 | 0.8 |
| Entropy | 0.4 | 0.4 | 0.4 | 0.4 | 0.4 | 0.4 |
| HS | 0.1 | 0.1 | 0.1 | 0.2 | 0.1 | 0.1 |
| CS | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 |
| Time | 12.2 | 12.8 | 11.7 | 11.9 | 17.9 | 20.8 |

Table 12. Comparison of results (Tic-Tac-Toe)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| Best | 538 | 581 | 551 | 579 | 543 | 585 |
| Worst | 823 | 775 | 720 | 817 | 667 | 816 |
| Avg | 603 | 604 | 606 | 601 | 545 | 642 |
| Std. | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 |
| Evaluation Measures | | | | | | |
| Purity | 0.7 | 0.7 | 0.7 | 0.7 | 0.7 | 0.7 |
| Entropy | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 | 0.6 |
| HS | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| CS | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 | 0.0 |
| Time | 47.6 | 54.9 | 46.5 | 51.3 | 83.0 | 102.2 |

Table 13. Comparison of results (Vote)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| SSE | | | | | | |
| Best | 376 | 429 | 410 | 396 | 398 | 468 |
| Worst | 567 | 659 | 682 | 608 | 664 | 653 |
| Avg | 390 | 455 | 481 | 419 | 459 | 533 |
| Std. | 0.2 | 0.2 | 0.2 | 0.3 | 0.2 | 0.2 |
| Evaluation Measures | | | | | | |
| Purity | 0.7 | 0.6 | 0.6 | 0.6 | 0.8 | 0.7 |
| Entropy | 0.5 | 0.6 | 0.6 | 0.6 | 0.4 | 0.5 |
| HS | 0.4 | 0.1 | 0.1 | 0.2 | 0.3 | 0.2 |
| CS | 0.3 | 0.1 | 0.0 | 0.1 | 0.3 | 0.2 |
| Time | 12.2 | 13.0 | 13.6 | 22.2 | 12.2 | 23.7 |

Table 14. Comparison of results (WaveformEW)

| Criteria | DL-MFA | OB-SSO | BDA-SA | B-BOA | B-GHO | IB-SSA |
|----------------------------|--------|--------|--------|-------|-------|--------|
| SSE | | | | | | |
| Best | 1582 | 2696 | 2824 | 2480 | 2069 | 3364 |
| Worst | 3 | 0 | 2 | 8 | 1 | 7 |
| Avg | 1696 | 2897 | 2964 | 3224 | 2926 | 3826 |
| Std. | 0.2 | 0.2 | 0.2 | 0.3 | 0.2 | 0.2 |
| Evaluation Measures | | | | | | |
| Purity | 0.5 | 0.3 | 0.3 | 0.3 | 0.3 | 0.4 |
| Entropy | 0.7 | 0.8 | 0.8 | 0.8 | 0.8 | 0.8 |
| HS | 0.3 | 0.2 | 0.2 | 0.3 | 0.3 | 0.2 |
| CS | 0.2 | 0.1 | 0.1 | 0.1 | 0.1 | 0.1 |
| Time | 1194 | 2239 | 1709 | 1820 | 2069 | 2922 |

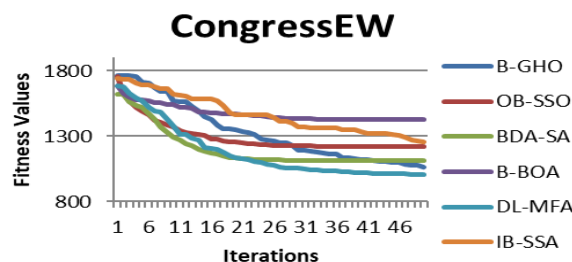


Figure. 5 Result for fitness function SSE (CongressEW dataset)

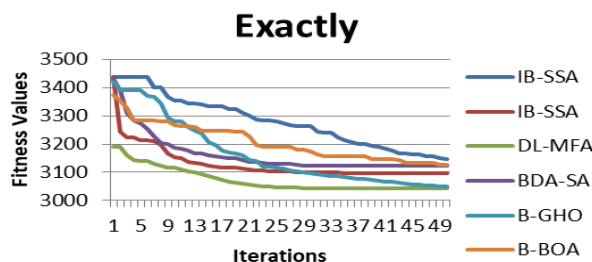


Figure. 6 Result for fitness function SSE (Exactly dataset)

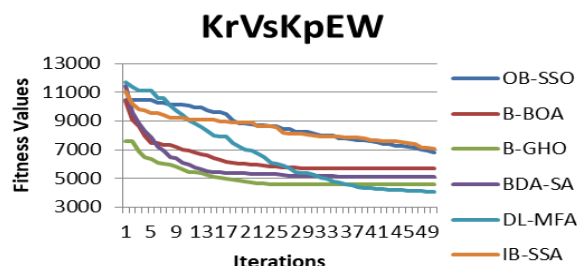


Figure. 7 Result for fitness function SSE (KrVsKpEW dataset)

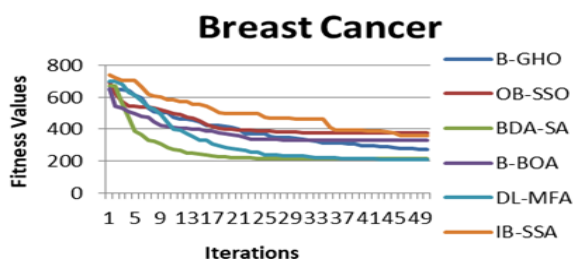


Figure. 3 Result for fitness function SSE (Breast Cancer dataset)

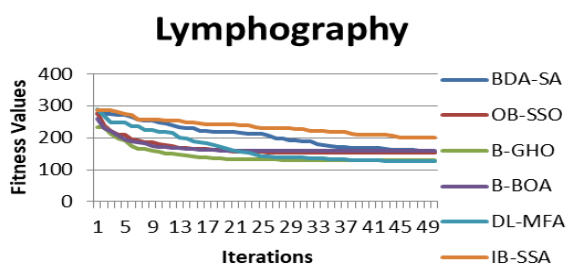


Figure. 8 Result for fitness function SSE (Lymphography dataset)

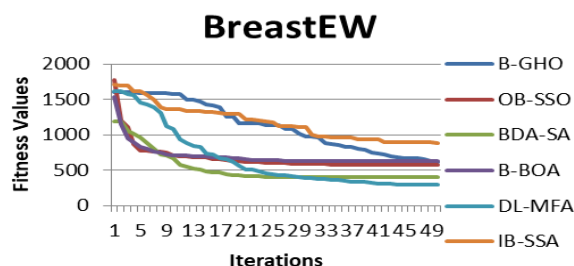


Figure. 4 Result for fitness function SSE (BreastEW dataset)

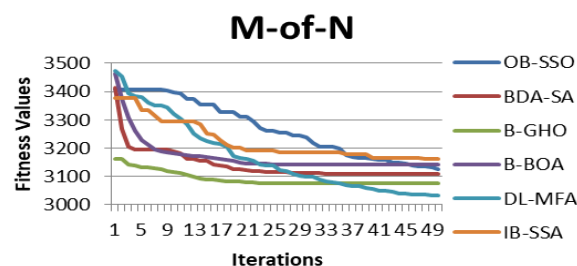


Figure. 9 Result for fitness function SSE (M-of-N dataset)

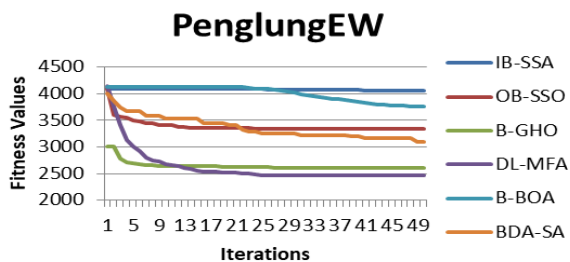


Figure. 10 Result for fitness function SSE (PenglungEW dataset)

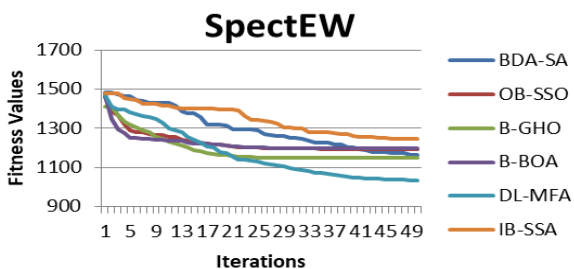


Figure. 11 Result for fitness function SSE (SpectEW dataset)

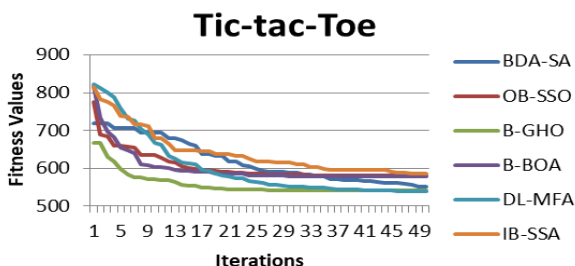


Figure. 12 Result for fitness function SSE (Tic-tac-Toe dataset)

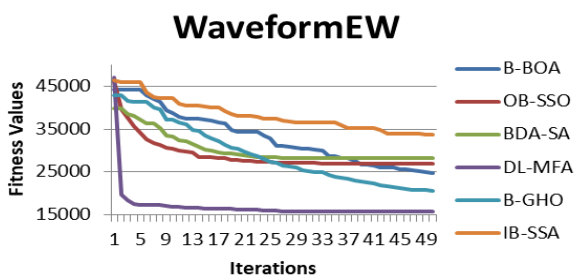


Figure. 13 Result for fitness function SSE (Vote dataset)

8. Conclusion

This paper proposes a discrete moth flame algorithm (DL-MFA) for feature selection that has been improved with levy flights. Levy flights are used in this algorithm to further balance the exploration and exploitation of MFA. Along with presenting the DL-MFA, the experimental analysis and outcomes were also covered. Purity, entropy, homogeneity score, and completeness score

evaluation measures are used to evaluate the performance of the algorithm. This algorithm (DL-MFA) produces purity in the range of 90% to 100%, very less entropy values up to 10%, good homogeneity score up to 70%, and completeness score up to 70%. Based on the results achieved, it was determined that DL-MFA produces good performance in terms of quality, consistency, and convergence when compared to the other SIA algorithms. Future applications of DL-MFA to various real-world issues can be made by combining classifiers such as neural networks (NN) and support vector machines (SVM).

Conflicts of Interest

The authors declare no conflict of interest

Author Contributions

Conceptualization, NMS, PDP, MJR and ML; methodology and software, NMS, and PDP; validation, and formal analysis, NMS, PDP and MJR; writing—review and editing, NMS; visualization, NMS, PDP, MJR; supervision, NMS investigation, NMS; resources, PDP, MJR and ML; data collection, PDP, MJR and ML; writing—original draft preparation, NMS, PDP, MJR and ML.

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