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DBSCAN with cuckoo search algorithm for social networks community detection

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ABSTRACT

The discovery of community structures in large-scale, complex networks is of fundamental importance to social network analysis. This task has enormous implications for understanding information flow and the behavioral characteristics of groups. This paper introduces a new hybrid model that combines the DBSCAN algorithm with a modified Cuckoo Search Optimization (CSO)algorithm of L'evy flight. This synergy leverages DBSCAN's robust capability to isolate noise and identify dense core nodes, while the optimized CSO performs a global search for the most effective community partitions. The proposed model was rigorously evaluated against a suite of established algorithms, including the Bat Algorithm, AFSA, Multilevel, Walktrap, AKHSO, Ant-Lion Optimizer, Lion Optimization Algorithm, and standard Cuckoo Search. Experimental values of four standard social networks substantiate the better performance of our method, which obtained the top recorded results for both Modularity and NMI. These findings indeed validate that the approach not only detects communities with higher internal cohesion but also more properly mirrors the observed ground-truth structures, verifying its effectiveness and strength.

1. Introduction

Internet pervasiveness has radically changed global communications with seamless social interactions by virtue of various applications being made possible by smart electronic equipment like smart phones, smart televisions, and computers. These online spaces can themselves be envisioned as intricate networks with individual elements like users represented by nodes, while interactions or relationships amongst them are

characterized by edges with sites like Twitter, LinkedIn, and Facebook being a few illustrations [1]. One critical factor to a comprehension of the functionality and architecture of these networks is community detection—sets of nodes with a larger number of connectors to one another relative to the full network and frequently with mutual attributes or functionalities [1–3].

The community discovery task has stimulated various computational techniques. These include

algorithms based on random walks [4, 5], clustering techniques [6], and optimization strategies aimed at maximizing quality metrics like modularity [7]. While these new developments, it is still challenging to accurately partition communities under noise and outlying nodes. More recent surveys and advancements, including those leveraging deep learning approaches, are discussed in [34, 35].

Noting this issue, we propose a novel two-stage hybrid framework as a solution for it. At first stage, DBSCAN is applied as a preprocessing step to efficiently identify and filter out noise nodes potentially interfering with clear boundaries between communities. At the second stage, an optimized version of the Cuckoo Search Algorithm with L'evy flight for efficient global search is applied to further improve the community structure optimization in the cleaned network. The proposed framework was evaluated on four well-known social network datasets and compared against several state-of-the-art algorithms using Precision, Recall, Accuracy, and F-measure metrics.

The structure of this work is defined as follows: Section 2 provides a literature review regarding social networks, community detection measures, and preliminary algorithms applied. Section 3 provides the architecture of the new DBSCAN-CSO model. Section 4 includes experimental outcomes with comparative analysis. Section 5 discusses the findings and explores the limitations of the proposed approach. Section 6 concludes the work and briefly discusses directions for further research.

2. Literature Review

2.1. Social Network and Community Discovery Introduction

Social Network consists of sets of nodes forming interrelated objects that are joined by edges. They can mathematically be represented as G(N, E), with N being the set of nodes and denoted by |N| = n, and E denoted by the set of relations connecting the nodes, $E \subseteq N \times N, |E| = m$.

Communities can be expressed as sets of non-empty node sets [8]. The task is to partition a set of nodes $\{x_1, x_2, \dots, x_n\}$, into disjoint sets (communities C), expressed as $C = \{N_1, \dots, N_{cn}\}$, where cn is the number of all the communities. The community's size is to meet the objective quality function.

$$\min F(S), S \in \Omega., \tag{1}$$

where F(S) is an objective quality function and S is the quality measure minimized without loss in the context.

2.2. Social Network Analysis Measures.

Social Networks has quite a few of these measures that are used for analysis, for example, Association Index, Strength, Closeness Centrality, Eigenvector centrality, Node Degree, Affinity, and Reach [9], [10].

Modularity is yet another quantity built to validate the quality and strength of partitions of networks. Modularity measure by Girvan and Newman [11], [12]. Mathematically, the Modularity can be expressed as in equation (2). Suppose ci is the community where node i is assigned. Then the fraction of graph edges that are inside the communities, i.e., that connect the nodes belonging to the same community, is:

$$\frac{\sum_{ij} A_{ij} \delta(c_i, c_j)}{\sum_{ij} A_{ij}} = \frac{1}{2m} \sum_{ij} A_{ij} \delta(c_i, c_j). \tag{2}$$

The adjacency matrix (A) with (n) nodes:

$$A_{ij} = \begin{cases} 1, & where \ i \ and \ j \ connected \\ 0. & otherwise \end{cases}$$
 (3)

The δ -function $\delta(u, v)$ has two values: 1 where u = v and 0 otherwise, while the term $\frac{1}{2m}\sum_{ij}A_{ij}$ represents the number of edges. The modularity Q defined as:

$$Q = \left(\frac{1}{2m}\right) \sum_{ij} \left[A_{ij} - \frac{k_i k_j}{2m} \right] \delta(c_i, c_j)$$
 (4)

Normalized Mutual Information (NMI) is a measure of performance that measures the similarity between the detected communities in networks and the ground truth [13] [1]. NMI presented by A. Lancichinetti [14]. NMI calculated by equation (5).

$$I(X,Y) = -\frac{2\sum_{i=1}^{C_X} \sum_{j=1}^{C_Y} c_{ij} \cdot log(\frac{c_{ij}N}{c_i} \cdot c_j)}{\sum_{i=1}^{C_X} c_i \cdot log(\frac{c_i}{N}) + \sum_{j=1}^{C_Y} c_j \cdot log(\frac{c_j}{N})}$$
(5)

X and Y represents two networks, C is the confusion matrix, C_{ij} is the number of individuals in community i in X and in community j in Y, C_X and C_Y are the number of groups in part X and in part Y, C_i and C_j are the elements of row i and column j of C and N is the total number of individuals in the networks. The output of the equation value will be 1 if two communities are similar and 0 if there are various communities.

2.3. Density-Based Spatial Clustering of Applications with Noise

To enhance the quality of found communities, we use the DBSCAN algorithm [15] due to its established efficiency for finding clusters of arbitrary shape from noise-contaminated spatial data. Although partition-based methods, DBSCAN clusters are based on the concept of density connectivity, hence emerging successful in partitioning of core, borderline, and outlying nodes.

The algorithm takes two parameters: a number of points (MinPts) and radius (ε) to be at least. Mark a node as a core node if at least MinPts nodes can be seen within its ε -neighborhood. A border node is within the ε -neighborhood of a core-node but does not itself qualify for MinPts. Each of those nodes that is not a core node nor a borderline node is labeled as noise or an outlier. Its natural capability to exclude non-core elements during data preprocessing makes DBSCAN particularly for social network analysis, for which outliers are the typical case. It is thus desirable to observe that the algorithm's success depends upon the choice of ε and MinPts judiciously [18].

2.4. Cuckoo Search Algorithm

The Cuckoo Search Algorithm, pioneered by Yang and Deb [19], is a metaheuristic optimization algorithm inspired by the brood parasitism of certain cuckoo species. The algorithm is governed by three idealized rules:

- Every bird drops off only one egg (solution) at a random nest.
- The nests containing superior solution eggs are kept for later generations.
- 3. The number of nests of the host is constant, and the host can find the alien egg with the probability $pa \in [0,1]$. When finding it, the host can abandon the nest or refuse to hold the egg.

The L'evy flight creates new candidate solutions by a random-walk process whose step sizes are selected from a heavy-tailed distribution. It balances extensive exploitation of the search space to the same extent that it balances aggressive searching. The new solution is computed by the following equation [21]:

$$x_c^{(t+1)} = x_c^t + \alpha \oplus Levy(\lambda)$$
 (6)

The new solution is expressed $asx_c^{(t+1)}$, x_c^t is the detected solution, and α is the step size where $\alpha > 0$.

The probability of distribution that L'evy flight step is governed by:

$$Levy = t^{-\lambda}, 1 < \lambda \le 3 \tag{7}$$

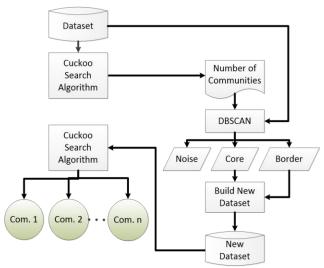


Figure 1: The proposed two-stage hybrid DBSCAN-CSO model. Stage 1: The Cuckoo Search Algorithm (CSA) provides an initial estimate of the number of communities (k). Stage 2: DBSCAN uses k to guide its parameters (ε, MinPts) to identify and remove noise/outlier nodes. Stage 3: The purified network is then processed by CSA again to obtain the final, optimized community partition.

3. The Proposed Model

The CSA employed in this research to identify the quantity of communities from the utilized datasets. Subsequently, the DBSCAN algorithm uses this estimated community count as an input parameter to cluster individuals engaged in Social Network datasets to core, border, and noise individuals by changing Eps. value to eliminate the outliers individuals in the datasets. After that, the datasets will be free of noise nodes and ready to implement CSA on it again to detect communities. The model of this research is shown in Fig. [1]. The DBSCAN Cuckoo Search Algorithm based on Algorithm 1. The locus-based adjacency encoding scheme employed in order to present individuals in CSA who are based on the genetic algorithm [22] [23].

4. Experimental Results

The developed approach was implemented for four data sets of social networks to discover the communities there. The number of communities in each data set is already defined. NMI was used [24] to compare the accuracy of the quality of the resulting groups is measured by modularity.

Algorithm 1: Cuckoo Search Algorithm and DBSCAN For Detecting Communities in Social Networks

- 1: **Input:** A network G = (N, E).
- 2: Output: Group of similar nodes.
- 3: procedure Initial value for the parameters:
- 4: ϵ , MinPts, α , s, Max no. of iteration and #trails
- 5: Label all nodes as core, border and outlier.
- 6: Eliminate *outlier* nodes.
- 7: Generate a random node of *n* host nests $\overrightarrow{x_h}$
- 8: For each nest, calculate the fitness F_h .
- Get a solution randomly by L'evy flight with fitness F_c.
- 10: Detect the best-solution
- 11: for each solution do
- 12: if $F_c > F_h$ then
- 13: Replace h by new-solution c
- 14: end if
- 15: end for
- 16: Update the best solution.
- 17: $t \leftarrow t + 1$.
- 18: Until t > Max no. ofIteration.
- 19: return The best-solution.
- 20: end procedure

The model was applied to four benchmark datasets: Zachary Karate Club [25], The Bottlenose Dolphin Network [26], American College Football Network [27], and the Facebook dataset [28]. The algorithm given is run with 10 iterations, and NMI and modularity are processed. The algorithm's iteration number is 10 times, and average NMI and Modularity are maintained. The parameters of CSA were as follows: maximum number of iterations was 100, trials were 10, α was 1.5, and s = 1000.

To compare the performance of proposed model, results are compared with different algorithms used to discover communities in social networks such as discrete Bat Algorithm [24], Artificial Fish Swarm Algorithm [29], Multilevel [30], Walktrap [31], A Discrete Krill Herd Swarm Optimization (AKHSO) [32], Ant Lion Algorithm [33], Lion Optimization Algorithm [34] and Cuckoo Search Algorithm.

Fig. [2] shows the average Modularity, Fitness and NMI of the proposed model before and after applying

DBSCAN on the social networks we used to remove the noises nodes, as shown after eliminating the noise nodes from the datasets the Modularity, Fitness and NMI values are increased. Fig. [3] shows the Modularity of some previous algorithms used to detect communities in Social Network and the proposed model was superior. Also, Fig. [4] shows the comparison between the NMI values of some previous algorithms and the proposed one. The proposed model gets the highest values for NMI. Fig. [5] reports that the accuracy, precision, recall, and F-measure of the proposed model are neither improved nor changed by using DBSCAN to eliminate noise nodes from the datasets.

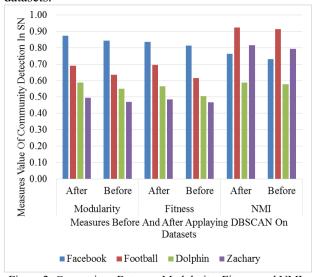


Figure 2: Comparison Between Modularity, Fitness and NMI Values of Social Networks Datasets Before and After Eliminate Noise Nodes.

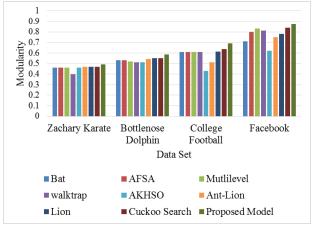


Figure 3: Comparison Between Modularity Values of Different Algorithms and The Proposed Model.

υı	e i: The Num	der of Noa	ies, Euges ai	na Commu	mues befor	e and Aner A	ppryring Dosc	AN OII	Tested Datase	is and Eps. van
	Dataset	Before DBSCAN		After DBSCAN					Number of	
				Alter DBSCAN					Communities	
		Nodes	Edges	Nodes	Edges	% Nodes	% Edges	Eps	Before	After
						Removed	Removed		DBSCAN	DBSCAN
	Facebook	3959	168486	3582	127516	9.52	24.32	5.70	156	139
	Football	115	1226	106	1036	7.83	15.50	3.75	11	13
	Dolphin	62	318	54	236	12.90	25.79	2.80	5	5
	Zachary	34	156	30	140	11.76	10.26	1.00	4	4

Table 1: The Number of Nodes, Edges and Communities Before and After Applying DBSCAN on Tested Datasets and Eps. Values.

Table 2: Average Number of Communities, Time, Mean, and Standard Deviation of The Tested Datasets According to Modularity and NMI Measures.

Dataset	Eps.	Average	Modularity			NMI			
		Number of	Time (ms)	Mean	Standard	Time (ms)	Mean	Standard	
		Communities			Deviation			Deviation	
Facebook	5.70	144.20	183860.731	0.64684	0.00758	184287.010	0.70325	0.01083	
Football	3.75	8.70	4867.701	0.84845	0.02548	4895.620	0.60957	0.00429	
Dolphin	2.80	4.60	3277.562	0.57915	0.00766	3332.668	0.54896	0.00080	
Zachary	1.00	4.00	2579.918	0.68730	0.00000	2562.486	0.46960	0.00000	

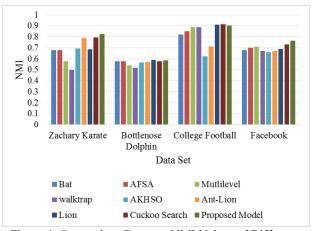


Figure 4: Comparison Between NMI Values of Different Algorithms and The Proposed Model.

Table [1] shows the number of nodes and edges of each benchmark dataset before and after using DBSCAN to remove the noise nodes from the datasets. By removing those nodes, the processing time will decrease and the accuracy of communities will increase. The table shows that the number of communities detected of American College Football Network dataset increased after applying DBSCAN, which means that CSA became able to detect more communities after removing noise nodes.

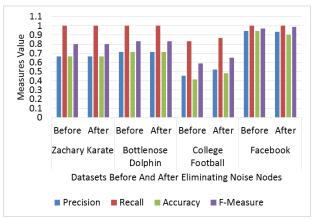


Figure 5: Recall, Accuracy and F-Measure Values of Proposed Model Before and After Applying DBSCAN on Datasets.

The new approach is run 10 times for all tested data sets. The mean and standard deviation are obtained and shown in Table [2]. It is observed from the table that the new approach 1 is stable for all data sets tested here; the standard deviation values are zero for the smallest data set and grow when the dataset size grows, but are all closest to the zero value, thus the data sets' distribution is normally distributed.

5. Discussion and Limitations

Even as the new proposed model is demonstrated to be better than the benchmark datasets, its limitations need to be noted so as to provide a balanced perspective and guide future work.

5.1. Parameter Sensitivity:

One of its primary limitations is that the DBSCAN algorithm is too sensitive to its parameters, ε (eps) and MinPts. The noise-removal preprocessing step depends much on the selection of these parameters' values. In this study, these parameters were empirically determined for each dataset. However, for totally new or extremely large networks with no visible ground truth, such empirical tuning might be challenging and can even not lead to the best result. As mentioned in future work, this parameter choice can be made automatic, e.g., within a meta-heuristic optimization cycle, which is a crucial means of enhancing model's generalizability the applicability.

5.2. Computational Complexity:

The computational cost of the model should also be taken into account. The Cuckoo Search Algorithm, as most population-based metaheuristics, calculates the fitness of numerous candidate solutions over numerous iterations. Node removal through DBSCAN reduces the size of the problem, but the CSO complexity may become a bottleneck in extremely large-scale networks. Alternative, more efficient encoding schemes or parallel computing schemes might be explored in future deployments to circumvent this issue.

5.3. Network Type Scope:

Static unweighted networks were the scope of the model being tested. Most social networks in the real world are dynamic networks whose composition evolves with time, with weighted edges representing the strength of relationships. These aspects are not considered in the proposed framework explicitly. Extending the model to operate in dynamic networks using incremental clustering techniques, or in weighted networks through the modification of the density in DBSCAN and the fitness function in CSO, is a significant and valuable area of future research.

Conclusion

This research has introduced and validated a novel

hybrid model for community detection in SN. The model innovatively combines the Cuckoo Search Algorithm (CSA) with DBSCAN preprocessing. An initial application of CSA identifies a preliminary community structure, the count of which informs the DBSCAN parameterization to effectively filter out noise and outlier nodes. A subsequent execution of CSA on the purified network then determines the final, optimized community partition.

The proposed model was subjected to rigorous comparative analysis against a range of established algorithms, including the Bat Algorithm, AFSA, Multilevel, Walktrap, AKHSO, Ant-Lion Optimizer, Lion Optimization Algorithm, and standard Cuckoo Search across four benchmark datasets. The results, tested by Modularity and NMI, consistently demonstrated the superiority of our approach. The significant finding is that the elimination of noise individuals through DBSCAN directly contributes to more accurate and robust community detection, thereby enhancing the performance of swarm intelligence algorithms in the complex domain of social network analysis. As explained in Section 5, future research will address the automatic optimization of parameter DBSCAN, scalability for extensive networks, and model extension to weighted and dynamic network structures.

Conflict of interest

The author confirms that there are no conflict of interests.

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