Binary rat swarm optimizer algorithm for computing independent domination metric dimension problem

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Abstract. In this article, we look at the NP-hard problem of determining the minimum independent domination metric dimension of graphs. A vertex set B of a connected graph G(V, E)resolves G if every vertex of G is uniquely recognized by its vector of distances to the vertices in B. If there are no neighboring vertices in a resolving set B of G, then B is independent. Every vertex of G that does not belong to B must be a neighbor of at least one vertex in B for a resolving set to be dominant. The metric dimension of G, independent metric dimension of G, and independent dominant metric dimension of G are, respectively, the cardinality of the smallest resolving set of G, the minimal independent resolving set, and the minimal independent domination resolving set. We propose the first attempt to use a binary version of the Rat Swarm Optimizer Algorithm (BRSOA) to heuristically calculate the smallest independent dominant resolving set of graphs. The search agent of BRSOA are binary-encoded and used to identify which one of the vertices of the graph belongs to the independent domination resolving set. The feasibility is enforced by repairing search agent such that an additional vertex created from vertices of G is added to B, and this repairing process is repeated until B becomes the independent domination resolving set. Using theoretically computed graph findings and comparisons to competing methods, the proposed BRSOA is put to the test. BRSOA surpasses the binary Grey Wolf Optimizer (BGWO), the binary Particle Swarm Optimizer (BPSO), the binary Whale Optimizer (BWOA), the binary Gravitational Search Algorithm (BGSA), and the binary Moth-Flame Optimization (BMFO), according to computational results and their analysis.

Keywords: optimization, metaheuristics, swarm-intelligence, rat swarm optimizer.

1. Introduction

Independent graph dominance numbers have recently been introduced in [1]. Robot navigation [2-4], network discovery and verification [5], localization of wireless sensor networks [6], combinatorial optimization [7], and applications to pharmaceutical chemistry [8] are only a few areas where metric dimension is used. Domination theory is applied in wireless communication networks [9], electrical networks [10], Backbone based routing [11], or spine based routing [12, 13] and chemical structures [14]. The independent domination set with the minimum cardinality is a logical choice for usage in any network type for information transmission.

2. Problem description

Let d(u, v) be the shortest path between two vertices $u, v \in V(G)$ in the connected graph G = (V, E). If the representation $r(v|B) = (d(v, x_1), d(v, x_2), \dots, d(v, x_k))$ is unique for every $v \in V(G)$, then the ordered vertex set $B = \{x_1, x_2, \dots, x_k\} \subseteq V(G)$ is a resolving set of G. If every vertex of $V \setminus B$ has at least one neighbor that belongs to G, then G is a dominating resolving set of G. A dominating resolving set G is independent if no two vertices in G are adjacent.

Let Card(X) stand for the cardinality of a set X. The metric dimension of G, denoted as

dim(G), the domination metric dimension of G, denoted as Ddim(G), and the independent domination metric dimension of G, denoted as $\gamma_{ir}(G)$, are as follows:

- $-dim(G) = min\{Card(B): B \text{ is a resolving set of } G\},$
- $-Ddim(G) = min\{Card(B): B \text{ is a domination resolving set of } G\},$
- $-\gamma_{ir}(G) = min\{Card(B): B \text{ is an independent dominating resolving set of } G\}.$

Example 2.1. The set $B = \{v_2, v_4, v_6\}$ is a minimal resolving set for the friendship graph F_{r_7} given in Fig. 1, and hence $dim(F_{r_7}) = 3$. Here, B is also a minimal domination resolving set since every vertex of $V \setminus B$ has at least one neighbor that belongs to B. For example, v_3 is adjacent to v_1 and v_2 . Also, v_5 adjacent to v_1 and v_4 . In the same regard, v_7 adjacent to v_1 and v_6 . Also, v_1 adjacent to v_2 , v_3 , v_4 , v_5 , v_6 and v_7 , and so $Ddim(F_{r_7}) = 3$. On the other hand, B is also independent domination resolving set of F_{r_7} . The set $B = \{v_2, v_4, v_6\}$ is a minimal independent domination resolving set of F_{r_7} , so v_{ir} (F_{r_7}) = 3.

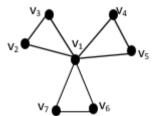


Fig. 1. Friendship graph F_{r_2}

Three elements are combined in the independent domination metric dimension problem: independent, dominance, and metric dimension of graphs. Integer programming is used to discuss the difficulty of determining the metric dimension of a graph G [8]. Let $D = [d_{ij}]$ be the distance matrix of G, $d_{ij} = d(v_i, v_j)$ for $1 \le i$, $j \le n$. For $x_i \in \{0, 1\}$, $1 \le i \le n$, the function F is defined by $F(x_1, x_2, \dots, x_n) = x_1 + x_2 + \dots + x_n$.

defined by $F(x_1, x_2, ..., x_n) = x_1 + x_2 + \cdots + x_n$. Minimizing F subject to the $\binom{n}{2}$ constraints $|d_{i1} - d_{j1}|x_1 + |d_{i2} - d_{j2}|x_2 + \cdots + |d_{in} - d_{jn}|x_n > 0$ is equivalent to finding a basis in the sense that if $x'_1, x'_2, ..., x'_n$ is a set of values for which F reaches its minimum, for $1 \le i < j \le n$. Then $B = \{v_i, x'_i = 1\}$ is a basis for G and conversely, if $B = \{v_{i1}, v_{i2}, ..., v_{in}\}$ is a basis for G and if we define:

$$x'_s = \begin{cases} 1, & s = i_j, j, & (1 \le j \le k), \\ 0, & \text{otherwise,} \end{cases}$$

then $F(x'_1, x'_2, ..., x'_n)$ is a minimum subject to the given constraints.

Both the metric dimension problem and the dominant set problem are NP-complete [15, 16]. As a result, the independent domination metric dimension $\gamma_{ir}(G)$ is a typical NP-complete problem that involves determining if $\gamma_{ir}(G) \leq K$ for a given graph G and input K. The remaining part of the paper is structured as follows: A literature review is presented in Section 3. In Section 4, the Rat Swarm Optimizer Algorithm is introduced. The BRSOA for calculating the independent domination metric dimension is provided in Section 5. Results of calculations are reported in Section 6. Section 7 presents the conclusion and recommendations for future work.

3. Literature review

For a number of graphs in the literature, the metric dimension, domination metric dimension, and independent domination metric dimension are all theoretically specified. Following is a brief summary of the key recently discovered theoretical metric dimension results [17-25]. The metric dimension of subdivisions of several graphs, including the Lilly graph, the Tadpole graph, and the

special trees star tree, bistar tree, and coconut tree is determined theoretically in [17], the bary centric subdivision of Möbius ladders and the generalized Petersen multigraphs in [18], trapezoid network, $Z - (P_n)$ network, open ladder network, tortoise network in [19], French windmill graph and Dutch windmill graph in [20], total graph of path power three and four in [21], two types of bicyclic graphs in [22], Mobius Ladder in [23], power of paths and complement of paths in [24], and Kayak Paddles graph and Cycles with chord in [25].

Theoretically, the independent metric dimensions are identified in [26, 27]. Cartesian product and corona of graphs are determined in [26], finite projective planes, finite biplanes in [27] and Titanium dioxide nanotube in [28]. The independent domination metric dimensions of the path graph, cycle graph, friendship graph, helm graph, and fan graph are determined theoretically in [1]. In [29, 30], the dominant metric dimension is found theoretically. The Path graph, cycle graph, star graph, complete graph, and complete bipartite graph are determined in [29], the corona product graph of G and H is studied whenever H is a path graph, and the cycle graph, complete bipartite graph, complete graph, and star graph are determined in [30]. The connected domination metric dimensions of the complete graph, path graph, and cycle graph are determined theoretically in [31]. To compute the metric dimension problem heuristically, however, only a small number of algorithms have been presented [32-34]. For determining the metric dimension of numerous classes of graph instances, such as pseudo-Boolean, crew scheduling, and graph coloring, a genetic algorithm has been proposed in [32]. A limited number of distinct individuals with the same objective value, the binary representation, frozen gene mutation, and the caching technique were all used. Infeasible individuals are changed by the addition of the required nodes in order to become feasible. In [33], Particle Swarm Optimization is used for determining the metric dimension where an infeasible particle is mended by adding some vertices until the particle becomes feasible and a real-valued vector of vertices is converted to a binary-valued vector using a linear function. The Particle Swarm Optimization is tested by computing the metric dimension of hypercube graphs. In order to enhance the current upper bounds in [34], a variable neighborhood search method has been suggested for tackling metric dimension and minimal doubly resolving set problems. The metric dimension problem and the minimal doubly resolving set problem are divided into a series of sub problems with an auxiliary objective function as the foundation for the variable neighborhood search method. Additionally, the equivalent new integer linear programming formulations for both problems are suggested. The connected domination metric dimension problem is resolved here by encoding and adapting the operations of the equilibrium optimization algorithm. Using theoretically computed graph results, the proposed binary equilibrium optimization algorithm is put to the test and contrasted with competing techniques. Also see more details in the literature [35-38], and some future notions may be applied to some applications like [39-41]. In the binary forms of the metaheuristics, a transfer function plays a crucial role, according to Mirjalili and Lewis in [42]. It has a major effect on both the balance between exploration and exploitation and the avoidance of local optima. In 2013, Sharafi et al. [43] altered the definition of the velocity vector to the probability of mutation in each cat dimension, introduced a transfer function to the tracking mode of the cat swarm algorithm, and converted the continuous cat swarm algorithm into a discrete binary cat swarm algorithm. A binary variant of the bat technique, which is likewise a probability value that uses a transfer function to convert velocity data to updated positions, was proposed by Mirjalili et al. [44] in 2014. A discrete binary bat method (BINBA) was proposed by Sabba and Chikhi [45] in the same year to solve binary space optimization problems.

4. Rat swarm optimizer (RSO) algorithm

4.1. Inspiration

Long-tailed, medium-sized rodents with different sizes and weights include rats. Black rats and brown rats are the two main species of rat. In the family of rats, male rats are referred to as

bucks and female rats as does. In general, rats are naturally socially intelligent. They train one another and engage in a variety of sports, including boxing, chasing, jumping, and tumbling. Rats are social, territorial creatures that coexist in households with both males and females. Rats are frequently quite aggressive in their behavior, which may cause the demise of some animals. This work's primary motive for pursuing and fighting with prey is this aggressive behavior. Rat chasing and hunting behaviors are mathematically modeled in this study in order to create the RSO algorithm and carry out optimization.

4.2. Mathematical model and optimization algorithm

This section explains the chasing and fighting behaviors of rats. The suggested RSO algorithm is then described.

4.2.1. Chasing the prev

Rats are typically sociable creatures that hunt their prey in packs using social agonistic behavior. The position of the prey must be known by the optimal search agent in order to mathematically define this behavior. With regard to the best search agent found thus far, the other search agents can update their places. In this mechanism, the following equations are proposed:

$$\vec{P} = A.\vec{P}_i(x) + C.\left(\vec{P}_r(x) - \vec{P}_i(x)\right),\tag{1}$$

where $\vec{P}_i(x)$ is the best optimal solution and $\vec{P}_r(x)$ defines the placements of the rats. But here's how the A and C parameters are determined:

$$A = R - x \times \left(\frac{R}{Max_{Iteration}}\right),\tag{2}$$

where $x = 0,1,2,...,Max_{Iteration}$, and:

$$C = 2. rand (\cdot). \tag{3}$$

Thus, R and C are independent random variables with ranges of [1, 5] and [0, 2], respectively. Over the course of iterations, the parameters A and C lead to better exploration and exploitation.

4.2.2. Fighting with prev

The following equation has been developed to quantitatively define the interaction of rats with prey:

$$\vec{P}_i(x+1) = |\vec{P}_r(x) - \vec{P}|,$$
 (4)

where the revised next position of the rat is defined by $\vec{P}_i(x+1)$. It changes other solutions' positions and saves the best solution and comparing search agents based on which one is the best. The parameters can be changed to obtain a different number of locations relative to the current position, as indicated in Eqs. (2) and (3). But this idea can also be expanded in an n-dimensional setting. As a result, the altered value of parameters A and C ensures exploration and exploitation. The best response is saved using the fewest operators via the suggested RSO technique.

5. Binary rat swarm optimizer algorithm for independent dominant metric dimension

Because it maintains a population of solutions and explores a wide region to find the optimum

global solution, the Rat Swarm Optimizer Algorithm can solve difficult optimization problems with several locally optimal solutions. This benefit enables a binary variant of the approach to be applied to the dominant independent metric dimension problem. Using position vectors located within the continuous real domain, search agents can navigate the search space in the continuous form of RSOA. By using an S-shaped transfer function to turn the continuous variable RSOA into a binary one, we may convert it to binary values. Position changes in discrete binary search space necessitate flipping between 0 and 1. The initialization phase makes use of the subsequent equation. Transfer function merits particular consideration and investigation since it plays a significant role in the discrete RSO algorithm:

$$Obinary_{ij} = \begin{cases} 1, & rand(\cdot) > 0.5, \\ 0, & \text{else,} \end{cases}$$
 (5)

where $rand(\cdot)$ refers to a random number between 0 and 1. To convert continuous values to binary ones, a transfer function is used. The sigmoid function (S) is applied as follows in this study:

$$S = \frac{1}{1 + e^{-10x^d}},\tag{6}$$

where S is the function output and x^d denotes the continuous-valued location at dimension d. To create a binary value, use the equation below:

$$SAbinary_{ij} = \begin{cases} 1, & rand(\cdot) < S, \\ 0, & \text{otherwise.} \end{cases}$$
 (7)

Table 1. Parameter setting with search agents 30 for all algorithms

		aren agents 50 for an argorithms		
Algorithms	Parameter name	Value		
	Number of generations	1000		
BRSOA	Control parameter (R)	[1, 5]		
DKSOA	Constant parameter C	[0, 2]		
	Number of runs	20		
	a_1	Decreasing from 2 to 0		
BWOA	a_2	Decreasing from –1 to –2		
	Number of runs	20		
	C_1	Increasing linearly from 0.5 to 2.5		
BPSO	C_2	Decreasing linearly from 2.5 to 0.5		
	Inertia weight (w)	0.8		
	Number of runs	20		
BGSA	Gravitational constant	100		
DUSA	Alpha coefficient	20		
BGWO	Control parameter (\vec{a})	[2, 0]		
	Convergence constant	[-1, -2]		
BMFO	Logarithmic spiral	0.75		
	Number of runs	20		

The proposed algorithm deals with the dominant independent resolving set problem as an optimization problem where it searches for the best solution, so each search agent can be represented as a one-dimensional vector $SAbinary_{ij} = (SA_{i1}, SA_{i2}, ..., SA_{ij})$, for which $SA\ binary_{ij}$ is a binary-valued position vector if the j-th element of the vector has a value of 1, it means that vertex j belongs to B. If every $v \in V(G)$ has a distinct representation r(v|B), then B is an independent dominant resolving set. The value of a binary-valued position vector is produced by computing the value of the S-shaped transfer function. In the BRSOA algorithm, when a search agent is not feasible as an independent dominant resolving set, that search agent is repaired by

adding a vertex from $V \setminus B$. This repair is applied until that object becomes an independent dominant resolving set.

The algorithm represents each solution (individual) in the population as a string of binary in which 1 means that the independent dominant resolving set will be chosen, then the corresponding value will be "1", and if the independent dominant resolving set is not selected, then the corresponding value will be "0". The pseudo-code in Algorithm 1.

Table 2. Algorithm 1: Pseudo-code of BRSOA

Input: The rat population P_i (i = 1, 2, ..., n)

Output: The optimal solution agent

- 1: Procedure RSOA
- 2: Initialize the parameters A, C and R
- 3: Evaluate the initial population and select the one with the best fitness value.
- 4: $P_r(x)$ ← The best search agent
- 5: while $(x \le Max_{Iteration})$ do
- 6: for each search agent do
- 7: Update the position of current search agent using Eq. (4)
- 8: Convert each $\overrightarrow{SA_i}$ into binary using the S-shaped transfer function in SAbinary_{ii}
- 9: Calculate the fitness of each SAbinaryii
- 10: Update the new position of the search agent using Eq. (7)
- 11: end for
- 12: Update parameters A, C and R
- 13: Verify if any search agents exist that extend beyond the specified search space, and then modify them
- 14: Evaluate the fitness of each search agent
- 15: Update P_r if a better solution becomes available than the previously optimal option
- 16: Convert each $\overrightarrow{SA_i}$ into binary using the S-shaped transfer function in SAbinary_{ij}
- 17: Calculate the fitness of each SAbinaryii
- 18: Update the new position of the search agent using Eq. (7)
- 19: Compare the fitness values of each search agent and choose the best candidate
- 20: Set x = x + 1
- 21: end while
- 22: return search agent with best fitness value
- 23: end procedure

5.1. Experimental results

This section uses theoretically generated graph findings to evaluate the proposed BRSOA. On a path graph, a cycle graph, a fan graph, a ladder graph and a circular ladder graph, the proposed BRSOA is compared to the BWOA, BPSO, BGSA, BGWO and BMFO. The code was implemented in MATLAB 2021b, and the algorithm tests and comparisons were carried out using a Windows 10 Ultimate 64-bit operating system with an Intel Core i7 running at 16 GB of RAM, a 1TBHDD + 1TBSSD hard drive. Table 1 displays the parameter setting values.

All algorithms have been run 20 times for each graph and the results are summarized in Tables 3-6. The tables are organized as follows: The first three columns contain the number of nodes N, the number of edges M, the independent domination resolving number γ_{cr} , the CPU time (t) used to indicate the exactly independent domination resolving number and iteration: The average number of iterations for finishing the algorithms to achieve the best solution, respectively.

It should be noted, based on Table 3, that when computing connected domination resolving number for path graph P_n , $4 \le n \le 19$, then the BRSOA has reached an optimal solution.

Table 3. Comparison between BRSOA, BWOA, BPSO, BGSA, BGWO and BMFO for computing connected domination resolving number for path graph P_0 , 4 < n < 19

	connected domination resolving number for path graph P_n , $4 \le n \le 19$								
N	Μ		BRSOA	BWOA	BPSO	BGSA	BGWO	BMFO	
		γ_{ir}	2	2	2	2	2	2	
4	3	t (sec)	16.3	52.7	36.4	24.5	29.2	46.7	
		Iteration	1	2	1	1	1	1	
		γ_{ir}	3	3	3	3	3	3	
5	4	t (sec)	35.8	87.3	51.9	31.2	49.1	73.5	
		Iteration	2	8	5	3	4	6	
		γ_{ir}	3	3	3	3	3	3	
6	5	t (sec)	58.4	132.9	86.1	72.8	83.4	109.6	
		Iteration	5	22	13	9	11	18	
			4	4	4	4	4	4	
7		γ_{ir}	83.7	175.2	155.9	126.9	68.3	148.1	
7	6	t (sec)	7	35	25	16		26	
		Iteration					19		
		γ_{ir}	4	4	4	4	4	4	
8	7	t (sec)	105.2	258.3	236.3	178.1	194.9	215.4	
		Iteration	10	47	42	28	32	53	
		γ_{ir}	5	5	5	5	5	5	
9	8	t (sec)	121.8	389.5	311.6	202.8	243.1	298.3	
		Iteration	17	69	50	35	41	59	
		γ_{ir}	5	5	5	5	5	5	
10	9	t (sec)	156.4	478.1	382.2	274.5	299.3	385.9	
-		Iteration	31	78	36	29	56	45	
		γ_{ir}	6	6	6	6	6	6	
11	10	t (sec)	194.2	535.7	447.8	351.4	367.9	457.4	
11	10	Iteration	22	93	48	56	75	63	
			6	6	6	6	6	6	
12	11	γ_{ir} $t \text{ (sec)}$	229.1	589.2	513.1	415.9	406.5	529.3	
12	11	Iteration	27	71	39	64	47	82	
			7	7	7	7	7	7	
12	10	\gamma_{ir}	283.4	711.9	599.5	501.3	484.2	591.7	
13	12	t (sec)							
		Iteration	35	105	31	82	63	59	
1.4	1.2	γ_{ir}	7	7	7	7	7	7	
14	13	t (sec)	372.9	794.4	678.4	604.8	567.9	673.5	
		Iteration	45	93	58	69	108	74	
		γ_{ir}	8	8	8	8	8	8	
15	14	t (sec)	307.2	881.7	794.5	647.3	685.2	757.9	
		Iteration	41	62	84	95	81	103	
		γ_{ir}	8	8	8	8	8	8	
16	15	t (sec)	356.6	1013.5	937.4	705.9	779.1	869.5	
		Iteration	29	118	95	49	54	73	
		γ_{ir}	9	9	9	9	9	9	
17	16	t (sec)	328.5	1183.9	1021.8	802.4	895.3	957.3	
	- 0	Iteration	16	78	110	56	64	81	
			9	9	9	9	9	9	
10	17	γ_{ir} $t \text{ (sec)}$							
18	17		411.8	1357.2	1109.3	896.7	961.5	1034.6	
		Iteration	34	103	62	78	89	88	
		γ_{ir}	10	10	10	10	10	10	
19	18	t (sec)	443.3	1562.4	1239.7	975.3	1198.4	1156.1	
		Iteration	25	99	76	62	70	91	

Table 4. Comparison between BRSOA, BWOA, BPSO, BGSA, BGWO and BMFO for computing connected domination resolving number for cycle graph C_n , $4 \le n \le 15$, BRSOA has reached an optimal solution

N	М	-	BRSOA nas	BWOA	BPSO	BGSA	BGWO	BMFO
10	IVI	γ_{ir}	2	2	2	2	2	2
4	4	t (sec)	25.4	61.8		32.9	37.2	43.5
4	4	Iteration	1	3	42.7	1	1	2
			2	2	2 2	2	2	2
5	5	γ_{ir}		_				
)	3	t (sec) Iteration	34.8 1	79.3 7	56.2 3	48.2 2	54.9	63.6
			3	3	3	3	3	3
6	6	γ_{ir} $t \text{ (sec)}$	3 61.7	3 104.8	3 73.8	59.7	68.3	81.4
0	O	Iteration	3			39.7 6	9	
				19	8	-		13
7	7	γ_{ir}	3	3	3	3	3	3
/	/	t (sec) Iteration	80.9	135.6	97.5	91.3	87.1	108.7
			4	42	17	15	23	34
0	0	γ_{ir}	-		4	4		
8	8	t (sec) Iteration	104.2 14	176.9	91.7	118.2 24	128.6	137.9
				56	31		38	48
9	9	γ_{ir} $t \text{ (sec)}$	4	4	4	4	4	4
9	9	t (sec) Iteration	129.1 11	211.2 73	116.4 45	173.9 36	159.4 24	172.1 53
			5	5	5	5	5	5
10	10	γ_{ir} $t \text{ (sec)}$	146.3	267.9	174.8	198.4	182.1	207.2
10	10	Iteration	26	109	82	61	49	74
		γ _{ir}	5	5	5	5	5	5
11	11	t (sec)	188.9	341.6	229.4	224.3	216.1	283.9
11	11	Iteration	37	81	89	75	56	103
		γ _{ir}	6	6	6	6	6	6
12	12	t (sec)	207.5	419.2	367.8	268.1	289.4	342.6
12	12	Iteration	24	134	52	69	48	82
			6	6	6	6	6	6
13	13	γ_{ir} $t \text{ (sec)}$	173.9	485.7	442.9	309.7	351.9	407.2
13	13	Iteration	32	159	78	83	72	113
				7				7
1.4	1.4	γ_{ir}	7	,	7	7	7	
14	14	t (sec) Iteration	228.6 21	562.1 183	499.5 93	375.2	412.5	488.1 66
						56	54	
1.5	1.5	γ_{ir}	7	7	7	7	7	7
15	15	t (sec)	294.1	739.8	617.3	476.8	532.5	761.4
		Iteration	17	201	72	54	69	75

Table 5. Comparison between BRSOA, BWOA, BPSO, BGSA, BGWO and BMFO for computing connected domination resolving number for friendship graph F_{r_n} , $3 \le n \le 25$, BRSOA has reached an optimal solution

	T_n , $S \subseteq R \subseteq 2S$, Bits of this reached an optimal solution									
N	Μ		BRSOA	BWOA	BPSO	BGSA	BGWO	BMFO		
3	3	γ_{ir}	1	1	1	1	1	1		
		t (sec)	11.9	29.6	18.2	16.3	22.4	32.8		
		Iteration	1	2	1	1	1	2		
5	6	γ_{ir}	2	2	2	2	2	2		
		t (sec)	16.8	47.9	35.8	28.4	31.9	46.7		
		Iteration	1	7	4	3	2	5		
7	9	γ_{ir}	3	3	3	3	3	3		
		t (sec)	34.5	83.1	72.7	46.2	57.8	89.5		
		Iteration	6	25	16	7	13	11		

9	12	γ_{ir}	4	4	4	4	4	4
		t (sec)	56.7	139.8	96.8	71.5	81.9	127.4
		Iteration	9	44	35	16	21	32
11	15	γ_{ir}	5	5	5	5	5	5
		t (sec)	95.4	215.3	83.9	107.9	111.3	185.9
		Iteration	16	59	28	25	38	44
13	18	γ_{ir}	6	6	6	6	6	6
		t (sec)	143.6	294.1	154.7	185.6	125.4	248.7
		Iteration	28	39	49	42	30	73
15	21	γ_{ir}	7	7	7	7	7	7
		t (sec)	197.1	376.9	236.3	248.4	221.8	367.3
		Iteration	43	64	73	32	49	57
17	24	γ_{ir}	8	8	8	8	8	8
		t (sec)	262.7	456.3	298.2	293.7	306.7	433.9
		Iteration	36	75	109	58	74	86
19	27	γ_{ir}	9	9	9	9	9	9
		t (sec)	311.3	563.9	374.8	389.5	412.5	524.8
		Iteration	21	117	64	70	85	104
21	30	γ_{ir}	10	10	10	10	10	10
		t (sec)	352.9	657.4	458.9	479.3	464.9	561.2
		Iteration	28	159	86	55	78	123
23	33	γ_{ir}	11	11	11	11	11	11
		t (sec)	388.7	689.5	534.2	515.7	527.2	598.5
		Iteration	19	208	78	67	61	82
25	36	γ_{ir}	12	12	12	12	12	12
		t (sec)	296.2	774.8	592.8	588.4	596.3	634.3
		Iteration	14	186	91	45	82	97

Table 6. Comparison between BRSOA, BWOA, BPSO, BGSA, BGWO and BMFO for computing connected domination resolving number for triangular snake graph T_n , $3 \le n \le 31$, BRSOA has reached an optimal solution.

N	М		BRSOA	BWOA	BPSO	BGSA	BGWO	BMFO
		γ_{ir}	1	1	1	1	1	1
3	3	t (sec)	10.5	26.3	20.8	19.6	27.3	25.2
		Iteration	1	4	2	1	1	3
		γ_{ir}	2	2	2	2	2	2
5	6	t (sec)	18.2	63.7	46.2	28.4	31.9	46.7
		Iteration	1	9	3	2	3	6
		γ_{ir}	3	3	3	3	3	3
7	9	t (sec)	41.4	99.3	87.2	58.9	74.1	95.4
		Iteration	5	31	15	10	8	19
		γ_{ir}	4	4	4	4	4	4
9	12	t (sec)	59.8	161.7	103.5	41.6	91.2	117.5
		Iteration	12	38	25	18	21	32
		γ_{ir}	5	5	5	5	5	5
11	15	t (sec)	76.3	208.1	174.2	89.3	124.9	154.6
		Iteration	16	45	41	27	38	50
		γ_{ir}	6	6	6	6	6	6
13	18	t (sec)	105.9	246.3	211.9	137.6	192.8	204.5
		Iteration	11	71	62	44	56	78
		γ_{ir}	7	7	7	7	7	7
15	21	t (sec)	189.3	297.5	287.1	201.8	274.9	297.3
		Iteration	37	93	51	49	74	86
		γ_{ir}	8	8	8	8	8	8
17	24	t (sec)	256.4	406.2	358.3	287.9	326.2	382.9
		Iteration	24	59	45	76	60	147

		γ_{ir}	9	9	9	9	9	9
19	27	t (sec)	317.8	485.9	402.4	346.3	398.7	459.3
		Iteration	40	133	49	83	67	95
		γ_{ir}	10	10	10	10	10	10
21	30	t (sec)	383.6	517.4	467.2	398.2	456.8	562.4
		Iteration	19	106	208	53	104	203
		γ_{ir}	11	11	11	11	11	11
23	33	t (sec)	437.9	593.5	538.9	443.9	523.5	623.8
		Iteration	13	181	169	62	77	122
		γ_{ir}	12	12	12	12	12	12
25	36	t (sec)	393.2	649.2	592.6	562.8	587.2	692.3
		Iteration	15	93	103	116	159	181
		γ_{ir}	13	13	13	13	13	13
27	39	t (sec)	431.8	737.5	678.2	642.7	675.9	728.9
		Iteration	29	84	178	96	113	79
		γ_{ir}	14	14	14	14	14	14
29	42	t (sec)	490.3	821.9	731.7	703.5	765.8	784.1
		Iteration	21	138	217	71	85	193
		γ_{ir}	15	15	15	15	15	15
31	45	t (sec)	356.9	879.3	812.6	786.9	807.6	843.7
		Iteration	14	186	204	82	99	161

6. Comparison

To further demonstrate the excellence of proposed BRSOA, we choose BWOA, BPSO, BGSA, BGWO and BMFO algorithms to conduct experiments under the same conditions and compared the results. The results on graphs are shown in Tables 3-6, which indicate that proposed BRSOA algorithm, outperforms other algorithms on graphs, reaching 443.3 sec in BRSOA, 1562.4 sec in BWOA, 1239.7 sec in BPSO, 975.3 sec in BGSA, 1198.4 sec in BGWO, 1156.1 sec in BMFO for path graph and 294.3 sec in BRSOA, 739.8 sec in BWOA, 617.3 sec in BPSO, 476.8 sec in BGSA, 532.5 sec in BGWO and 761.4 sec in BMFO for cycle graph and 296.2 sec in BRSOA, 774.8 sec in BWOA, 592.8 sec in BPSO, 588.4 sec in BGSA, 596.3 sec in BGWO, and 634.3 sec in BMFO for friendship graph and 356.9 sec in BRSOA, 879.3 sec in BWOA, 812.6 sec in BPSO, 786.9 sec in BGSA, 807.6 sec in BGWO and 843.7 sec in BMFO for triangular snake graph. It proves the correctness and superiority of proposed BRSOA. Figs. 2, 3, 4 and 5 show the superiority of the proposed BRSOA on the BWOA, BPSO, BGSA, BGWO and BMFO according to the independent domination resolving number.

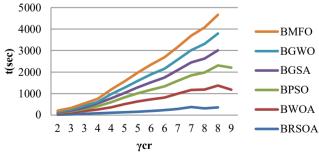


Fig. 2. The superiority of BRSOA on the BWOA, BPSO, BGSA, BGWO and BMFO

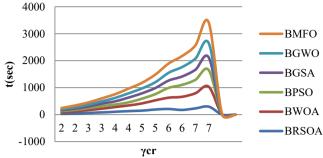


Fig. 3. The superiority of BRSOA on the BWOA, BPSO, BGSA, BGWO and BMFO

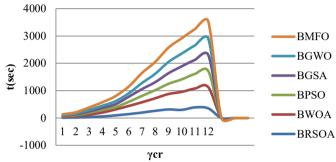


Fig. 4. The superiority of BRSOA on the BWOA, BPSO, BGSA, BGWO and BMFO

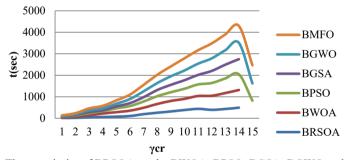


Fig. 5. The superiority of BRSOA on the BWOA, BPSO, BGSA, BGWO and BMFO

7. Conclusions

In this paper, a binary variant of the basic Rat Swarm Optimizer Algorithm (BRSOA) is adapted for determining the minimum independent domination resolving set of graphs and compared to BWOA, BPSO, BGSA, BGWO and BMFO. Comparisons were performed on the graphs: path graph, cycle graph, friendship graph and triangular snake graph. Experimental results and their analysis confirmed the superiority of the proposed BRSOA for solving the independent domination metric dimension problem. For further work in the future, we plan to compute other variants of metric dimension by other metaheuristic algorithms and compare them with competitive algorithms.

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Data availability

The datasets generated during and/or analyzed during the current study are available from the corresponding author on reasonable request.

Author contributions

Iqbal M. Batiha: conceptualization, methodology, software, validation, formal analysis, investigation data curation. Basma Mohamed: conceptualization, methodology, software, validation, formal analysis, investigation, data curation.

Conflict of interest

The authors declare that they have no conflict of interest.

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