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Slime Mould Algorithm: A New Method for Stochastic Optimization

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Abstract

In this paper, a new stochastic optimizer, which is called slime mould algorithm (SMA), is proposed based upon the oscillation mode of slime mould in nature. The proposed SMA has several new features with a unique mathematical model that uses adaptive weights to simulate the process of producing positive and negative feedback of the propagation wave of slime mould based on bio-oscillator to form the optimal path for connecting food with excellent exploratory ability and exploitation propensity. The proposed SMA is compared with up-to-date metaheuristics in an extensive set of benchmarks to verify the efficiency. Moreover, four classical engineering structure problems are utilized to estimate the efficacy of the algorithm in optimizing engineering problems. The results demonstrate that the proposed SMA algorithm benefits from competitive, often outstanding performance on different search landscapes. Source codes of SMA are publicly available at<http://www.alimirjalili.com/SMA.html>

Keywords

Slime mould optimization algorithm; Adaptive weight; Engineering design problems; Constrained optimization

1 Introduction

Metaheuristic algorithms (MAs) have become prevalent in many applied disciplines in recent decades because of higher performance and lower required computing capacity and time than deterministic algorithms in various optimization problems [1]. Simple concepts are required to achieve favorable results, and it is facile to transplant to different disciplines. Also, the lack of randomness in the later stage of some deterministic algorithm makes it inclined to sink into local optimum, and random factors in MAs can make the algorithm search for all optimal solutions in search space, thus effectively avoiding local optimum. In linear problems, some gradient descent algorithms such as [2] are more efficient than stochastic algorithms for the utilization of gradient information. The convergence speed of MAs will be less than gradient descent algorithms and can be considered as a drawback. In non-linear problems, however, MAs typically commence the optimization process with randomly generated solutions and do not demand gradient information, which makes the algorithm eminently suitable for practical problems when the derivative information is unknown. In real-world scenarios, the solution space of many problems is often indeterminate or infinite. It may be infeasible to find optimal solutions by traversing the solution space under current circumstances. MAs detect the proximate optimal solution of the problem by sampling the enormous solution space randomly in a certain way, to find or generate better solutions for the optimization problem under limited circumstances or computational capacity.

MAs are typically inspired by real-world phenomena find better heuristic solutions for optimization problems by simulating physical rules or biological phenomena. MAs can be divided into two main categories: swam-based methods and evolutionary techniques. The first kind mainly simulate physical phenomena, apply mathematical rules or methodologies including: Multi-Verse Optimizer (MVO) [3], Gravitational Local Search Algorithm (GLSA) [4], Charged System Search (CSS) [5], Gravitational Search Algorithm (GSA) [6], Sine Cosine Algorithm (SCA) [7], Simulated Annealing (SA) [8], Teaching-Learning-Based Optimization (TLBO) [9], Central Force Optimization (CFO) [10] and Tabu Search (TS) [11]. Nature-inspired methods mainly include two types: evolutionary methods and intelligent swarm techniques. The inspiration of the evolutionary algorithm (EA) originates from the process of biological evolution in nature. Compared with the traditional optimization algorithm, it is a global optimization method with better robustness and applicability.

Some of the widespread algorithms in the class of EA are Genetic Algorithm (GA) [12], Genetic Programming (GP) [13], Evolution Strategy (ES) [14], Evolutionary Programming (EP) [15] and Differential Evolution (DE) [16]. The application of ES and EP in scientific research and practical problems is also becoming more and more extensive. Swarm Intelligence (SI) [17] includes a collective or social intelligence that artificially simulates the decentralization of biological clusters in nature or the collective behavior of self-organizing systems. In this class of algorithms, the inspiration usually comes from biological groups in nature that have collective behavior and intelligence to achieve a certain purpose. In general, SI algorithms are more advantageous than evolutionary algorithms because SI algorithms are accessible to appliance than evolutionary algorithms with less operators that need to be controlled. Moreover, the SI algorithm has a stronger capability to record and utilize historical information than EA. Established and recent algorithms in this class are: Particle Swarm Optimization (PSO) [18], Wasp Swarm Optimization (WSO) [19], Bat-inspired Algorithm (BA) [20] , Grey Wolf Optimization (GWO) [21], Fruit Fly Optimization

(FOA) [22] , Moth Flame Optimization (MFO) [23], Ant Colony Optimization (ACO) [24], Harris Hawk Optimizer (HHO) [25], and Artificial Bee Colony (ABC) [26]. A schematic design for the classification of evolutionary and SI methods are shown in **Figure 1**.

Figure 1 classification of evolutionary and SI methods

Although different MAs have some distinctness, they all have two identical stages in the search gradation: exploration and exploitation [27, 28]. Exploration phase refers to the process of searching solution space as widely, randomly, and globally as possible. Exploitation phase refers to the competence of the algorithm to search more accurately in the area acquired by the exploration phase, and its randomness decreases while its precision increases. When the exploration ability of the algorithm is dominant, it can search the solution space more randomly and produce more differentiated solution sets to converge fleetly. When the exploitative ability of the algorithm is dominant, it searches more locally to enhance the quality and precision of the solution sets. However, when the exploration facility is improved, it will lead to reductions in the exploitation capability, and vice versa. Another challenge is that the balance of these two abilities is not necessarily identical to different problems. Therefore, it is relatively challenging to attain an appropriate balance between the two phases that are efficient for all optimization problems.

Despite the success of conventional and recent MAs, none of them can guarantee finding the global optimum for all optimization problems. This has been proven logically the No-Free-Lunch (NFL) theory [29]. This theorem motivated numerous researchers to design a new algorithm and solve new classes of problems more efficiently. With the aspiration of proposing a more versatile and efficient algorithm, this paper introduces a new meta-heuristic algorithm: slime mould algorithm (SMA). This method is aroused by the diffusion and foraging conduct of slime mould. An overall set of 33 benchmarks and four famous manufacturing design problems has rigorously verified the effectiveness and robustness of SMA.

The remainder of the paper is structured as below. Section 2 illustrated the concept and elicitation source of slime mould algorithm, and the mathematical model was established. Section 3 firstly gave a qualitative analysis of the algorithm and made a comprehensive comparison of 33 benchmark functions, then tested it on four engineering design problems. Section 4 summarized the whole work and put forward some inspirations for future work.

2 Slime mould algorithm

In this section, the basic concept and conduct of slime mould will be introduced. Then a mathematical model inspired by its behavior pattern will be established.

2.1 Originality

Before this article, some scholars have proposed similar naming algorithms, but the way of designing the algorithm and usage scenarios are quite different from the algorithms proposed in this paper. Monismith and Mayfield [30] solves the single-objective optimization problem by simulating the five life cycles of amoeda Dictyostelium discoideum: a state of vegetative, aggregatice, mound, slug, or dispersal while using ε-ANN to construct an initial position-based mesh. Li et al. [31] proposed a method to construct wireless sensor networks by using two forms of slime mould tubular networks to correspond to two different regional routing protocols. Qian. et al. [32] combined the Physarum network with the ant colony system to improve the algorithm's competence to avoid local optimal values to handle the Traveling Salesman Problem better. Inspired by the diffusion of slime mould, Schmickland Crailsheim [33] proposed a bio-inspired navigation principle designed for swarm robotics. Becker [34] generated inexpensive and fault-tolerant graphs by simulating the foraging process of the slime mould Physarum polycephalum. As can be seen from the above discussion, most of the modeled slime mould algorithms were used in graph theory and generation networks. The algorithm used to optimize the problem [30] simulates the five life cycles of amoeda Dictyostelium discoideum, but the experiments and proofs in the article are slightly less.

The SMA proposed in this paper mainly simulates the behavior and morphological changes of slime mould Physarum polycephalum in foraging and does not model its complete life cycle. At the same time, the use of weights in SMA is to simulate the positive and negative feedback generated by slime mould during foraging, thus forming three different morphotype, is a brand new idea. This paper also conducted a full experiment on the characteristics of the algorithm. The results in the next sections demonstrate the superiority of the SMA algorithm.

2.2 Concept and elicitation

The slime mould mentioned in this article generally refers to Physarum polycephalum. Because it was first classified as a fungus, thus it was named "slime mould" whose life cycle was originally specified by Howard [35] in a paper published in 1931. Slime mould is a eukaryote that inhabits cool and humid places. The main nutritional stage is Plasmodium, the active and dynamic stage of slime mould, and also the main research stage of this paper. In this stage, the organic matter in slime mould seeks food, surrounds it, and secretes enzymes to digest it. During the migration process, the front end extends into a fan-shaped, followed by an interconnected venous network that allows cytoplasm to flow inside [36], as shown in **Figure 2**. Because of their unique pattern and characteristic, they can use multiple food sources at the same time to form a venous network connecting them. If there is enough food in the environment, slime mould can even grow to more than 900 square centimeters [36].

Owing to the feature of slime mould can be easily cultured on agar and oatmeal [37], they were

widely used as model organisms. Kamiya and his colleagues [38] were the first team to study the detailed process of the cytoplasmic flow of slime mould. Their work is of great help to our subsequent understanding of the way slime mould move and connects food in the environment. We now cognize that when a vein approaches a food source, the bio-oscillator produces a propagating wave [39] that increases the cytoplasmic flow through the vein, and the faster the cytoplasm flows, the thicker the vein. Through this combination of positive-negative feedback, the slime can establish the optimal path to connect food in a relatively superior way. Therefore, slime mould was also mathematically modeled and applied in graph theory and path networks [40-42].

Figure 2 Foraging morphology of slime mould

The venous structure of slime mould develops along with the phase difference of the contraction mode [39], so there are three correlations between the morphological changes of the venous structure and the contraction mode of slime mould.

1) Thick veins form roughly along the radius when the contraction frequencies vary from outside

to inside.

2)When the contraction mode is unstable, anisotropy begins to appear.

3)When the contraction pattern of slime mould is no longer ordered with time and space, the venous structure is no longer present.

Therefore, the relationship between venous structure and contraction pattern of slime mould is consistent with the shape of naturally formed cells. The thickness of each vein is determined by the flow feedback of the cytoplasm in the Physarum solver [43]. The raise in the flow of cytoplasm leads to an increase in the diameter of veins. As the flow decreases, the veins contract because of the decrease of the diameter. Slime mould can build a stronger route where food concentration is higher, thus ensuring that they get the maximum concentration of nutrients. Recent studies have also revealed that slime mould have the competence of making foraging arrangements based on optimization theory [44]. When the quality of various food sources is different, slime mould can choose the food source with the highest concentration. However, slime mould also needs to weigh speed and risk in foraging. For instance, slime mould needs to make faster decisions in order to avoid environmental damage to them. Experiments have shown that the quicker the decision-making speed is, the possibilities of slime mould to find the prime food source is smaller [45]. Therefore, when deciding the source of food, slime mould obviously needs to weigh the speed and accuracy.

Slime mould need to decide when to leave this area and search in another area when foraging. When lacking complete information, the best way for a slime mould to estimate when to leave the current position is to adopt heuristic or empirical rules based on the insufficient information currently available. Experience has shown that when slime mould encounter high-quality food, the probability of leaving the area is reduced [46]. However, due to its unique biological characteristics, slime mould can utilize a variety of food sources at the same time. Therefore, even if the slime mould has found a better source of food, it can still divide a component of the biomass to exploit both resources simultaneously when higher quality food is found [43].

Slime mould can also dynamically adjust their search patterns according to the quality of foodstuff provenience. When the quality of food sources is high, the slime mould will use the region-limited search method [47], thus focusing the search on the food sources that have been found. If the denseness of the food provenience initially found is low, the slime mould will leave the food source to explore other alternative food sources in the region [48]. This adaptive search strategy can be more reflected when different quality food blocks are dispersed in a region. Some of the mechanisms and characteristics of the slime mould mentioned above will be mathematically modeled in the subsequent sections.

2.3 Mathematical model

In this part, the mathematical model and method proposed will be described in details.

2.3.1 Approach food

Slime mould can approach food according to the odor in the air. To express its approaching behavior in mathematical formulas, the following formulas are proposed to imitate the contraction mode:

$$
\overrightarrow{X(t+1)} = \begin{cases} \overrightarrow{X_b(t)} + \overrightarrow{vb} \cdot (\overrightarrow{W} \cdot \overrightarrow{X_A(t)} - \overrightarrow{X_B(t)}), r < p \\ \overrightarrow{vc} \cdot \overrightarrow{X(t)}, r \ge p \end{cases} \tag{2.1}
$$

where \overrightarrow{vb} is a parameter with a range of $[-a, a]$, \overrightarrow{vc} decreases linearly from one to zero. t represents the current iteration, X_b represents the individual location with the highest odor concentration currently found, X represents the location of slime mould, X_A and X_B represent two individuals randomly selected from slime mould, \vec{W} represents the weight of slime mould. The formula of p is as follows:

$$
p = \tanh|S(i) - DF|
$$
\n(2.2)

where $i \in 1, 2, ..., n$, $S(i)$ represents the fitness of \vec{X} , DF represents the best fitness obtained in all iterations.

The formula of \overrightarrow{vb} is as follows:

$$
\overrightarrow{vb} = [-a, a] \tag{2.3}
$$

$$
a = \operatorname{arctanh}(-\left(\frac{t}{\max_{-t}}\right) + 1) \tag{2.4}
$$

The formula of \vec{W} is listed as follows:

$$
\overrightarrow{W(SmellIndex(i))} = \begin{cases} 1 + r \cdot \log\left(\frac{bF - S(i)}{bF - wF} + 1\right), condition\\ 1 - r \cdot \log\left(\frac{bF - S(i)}{bF - wF} + 1\right), \quad others \end{cases}
$$
(2.5)

$$
SmellIndex = sort(S) \tag{2.6}
$$

where *condition* indicates that $S(i)$ ranks first half of the population, r denotes the random value

in the interval of $[0,1]$, bF denotes the optimal fitness obtained in the current iterative process, wF denotes the worst fitness value obtained in the iterative process currently, *SmellIndex* denotes the sequence of fitness values sorted(ascends in the minimum value problem).

Figure 3 visualizes the effects of Eq. (2.1). The location of searching individual \vec{X} can be updated according to the best location X_b currently obtained, and the fine-tuning of parameters vb , \vec{v} and \vec{W} can change the location of the individual. **Figure 3** is also used to illustrate the position change of the searching individual in three-dimensional space. $rand$ in the formula can make individuals form search vectors at any angle, that is, search solution space in any direction, so that the algorithm has the possibility of finding the optimum solution. Therefore, **Eq. (2.1)** enables the searching individual to search in all possible directions near the optimal solution, thus simulating the circular sector structure of slime mould when approaching food. It is also applicable to extend this concept to Hyper-dimensional space.

Figure 3 Possible locations in 2-dimention and 3-dimention

Figure 4 Assessment of fitness

2.3.2 Wrap food

This part simulates the contraction mode of venous tissue structure of slime mould mathematically when searching. The higher the concentration of food contacted by the vein, the stronger the wave generated by the bio-oscillator, the faster the cytoplasm flows, and the thicker the vein. **Eq. (2.5)** mathematically simulated the positive and negative feedback between the vein width of the slime mould and the food concentration that was explored. The component r in Eq. (2.5) simulates the uncertainty of venous contraction mode. \log is used to alleviate the change rate of numerical value so that the value of contraction frequency does not change too much. *condition* simulates the slime mould to adjust their search patterns according to the quality of food. When the food concentration is content, the bigger the weight near the region is; when the food concentration is low, the weight of the region will be reduced, thus turning to explore other regions. **Figure 4** shows the process of evaluating fitness values for slime mould.

Based on the above principle, the mathematical formula for updating the location of slime mould is as follows:

$$
\overrightarrow{X^*} = \begin{cases}\n\overrightarrow{rand} \cdot (UB - LB) + LB, rand < z \\
\overrightarrow{X_b(t)} + \overrightarrow{vb} \cdot \left(W \cdot \overrightarrow{X_A(t)} - \overrightarrow{X_B(t)}\right), r < p \\
\overrightarrow{vc} \cdot \overrightarrow{X(t)}, r \ge p\n\end{cases} \tag{2.7}
$$

where LB and UB denote the lower and upper boundaries of search range, rand and r denote the random value in $[0,1]$. The value of z will be discussed in the parameter setting experiment.

2.3.3 Grabble food

Slime mould mainly depends on the propagation wave produced by the biological oscillator to change the cytoplasmic flow in veins, so that they tend to be in a better position of food concentration. On the purpose of simulating the variations of venous width of slime mould, we used W , vb and \overrightarrow{vc} to realize the variations.

 \vec{W} mathematically simulates the oscillation frequency of slime mould near one at different food

concentration, so that slime mould can approach food more quickly when they find high-quality food, while approach food more slowly when the food concentration is lower in individual position, thus improving the efficiency of slime mould in choosing the optimal food source.

The value of \overrightarrow{vb} oscillates randomly between $[-a, a]$ and gradually approaches zero as the increasement of iterations. The value of $\vec{v}\vec{c}$ oscillates between [-1,1] and tends to zero eventually. The trend of the two values is shown as **Figure 5**. Synergistic interaction between $v\dot{b}$ and $\vec{v}\vec{c}$ mimics the selective behavior of slime mould. In order to find a better source of food, even if slime mould has found a better source of food, it will still separate some organic matter for exploring other areas in an attempt to find a higher quality source of food, rather than investing all of it in one source.

Moreover, the oscillation process of \overrightarrow{vb} simulates the state of slime mould deciding whether to approach the food source or find other food sources. Meanwhile, the process of probing food is not smooth. During this period, there may be various obstacles, such as light and dry environment, which restrict the spread of slime mould. However, it also improves the possibility of slime mould to find higher quality food and evades the trapping of local optimum.

The pseudo code of the SMA is shown in **Algorithm 1**. The intuitive and detailed process of SMA is shown in **Figure 6**.

There are still many mechanisms that can be added to the algorithm, or more comprehensive simulation of the life cycle of slime mould. However, to enhance the extensibility of the algorithm, we simplify the process and operators of the algorithm, leaving only the simplest algorithm as possible.

2.3.4 Computational complexity analysis

SAM mainly consists of the subsequent components: initialization, fitness evaluation, and sorting, weight update, and location update. Among them, *N* denotes the number of cells of slime mould, *D* denotes the dimension of functions, and *T* denotes the maximum number of iterations. The computation complexity of initialization is $O(N)$, the computation complexity of fitness evaluation and sorting is $O(N + N \log N)$, the computational complexity of weight update is $O(N \times D)$, the complexity of location update is $O(N \times D)$. Therefore, the total complexity of SMA is $O(N * (1 +$ $T * N * (1 + log N + 2 * D))).$

Figure 6 Flowchart of SMA

3 Experimental results and analyses

In this sector, we compared the SMA with some competitive MAs in an all-inclusive set of 33 benchmarks. The experimentations were ran on the operating system of Windows Server 2012 R2 Datacenter with 128 GB RAM and CPU of Intel (R) Xeon (R) E5-2650 v4 (2.20 GHz). The algorithms for comparison were coded by MATLAB R2018b.

3.1 Qualitative analysis

The qualitative analysis results of SMA in handling unimodal functions and multimodal functions are presented in **Figure 7** to intuitively analyze the position and fitness changes of slime mould during foraging. The figure is comprised of four concernment indicators: search history, the trajectory of the slime mould in the 1st dimension, the average fitness of slime mould, and convergence curve. Search history represents the location and distribution of slime mould in the iteration process. The trajectory of slime mould reveals the behavior of the position change of slime mould in the first part of the first dimension. Average fitness indicates the variation trend of the average fitness of the slime mould colony changes with the iteration process. Convergence curve shows the optimal fitness value in the slime mould during the iteration process.

From the search history subplot, the slime mould in different benchmark functions put up a similar cross-type search trajectory clustered near the optimal value, thus accurately searching in reliable search areas and reflecting fast convergence. Meanwhile, the distribution of slime mould is mainly concentrated in multiple regions with local optimum, which shows the tradeoff of slime mould between multiple local optimums.

The trajectory of the first slime mould in the first dimension can be used as a representative of other parts of slime mould, revealing the primary exploratory behavior of slime mould. The fast oscillation in the prophase and the slight oscillation in the anaphase can ensure the fast convergence of slime mould and the accurate search near the optimal solution [49]. As can be perceived from the figure, the position curve of slime mould has very large amplitude in the early iteration process, even up to 50% of the exploration space. In the later iteration period, if the function is smooth, the amplitude of the position of slime mould begins to decrease; if the amplitude of the function changes significantly, the position amplitude also changes greatly. This reflects the high adaptability and robustness of slime mould in different functions.

By observing the average fitness curve, the variation tendency of the fitness of slime mould during the iterative procedure can be visually observed. Although the average fitness curve of slime mould is oscillating, the average fitness value tends to decrease, and the oscillation frequency decreases inversely proportional to iterations, thus ensuring the rapid convergence of slime mould in the prophase and the precise search in the anaphase.

Convergence curve reveals the average fitness of the optimal fitness value searched by slime mould varies with iterations. By observing the downtrend of the curve, we can observe the convergence rate of slime mould and the time when it switches between the exploration and exploration gradation.

Figure 7 Qualitative analysis

3.2 Benchmark function validation

In this section, SMA was assessed on a comprehensive set of functions from 23 benchmarks and CEC 2014. These functions cover unimodal, multimodal, hybrid, and composite functions, as shown in **Tables 1-3**. Some composite functions of CEC 2014 are shown in **Figure 8**. Dim denotes the dimension of function; Range denotes the definition domain of the function, and f_{min} denotes the optimal value of the function. The MAs used for comparison include well-regarded and recent ones: WOA [50], GWO [21], MFO [23], BA [20], SCA [7], FA[51], PSO[18], SSA [52], MVO [3], ALO

[53], PBIL [54], DE [55] and advanced MAs: AGA[56], BLPSO [57], CLPSO [58], CBA [59], RCBA [60], CDLOBA [61], m_SCA [62], IWOA [63], LWOA [64], and CSSA [65]. The parameter setup of traditional MAs is detailed in **Table 4**. The parameter selection was based on the parameters used by the original author in the article or the parameters widely used by various researchers.

Figure 8 Illustration of CEC 2014 composite functions

Table 1

Unimodal and multimodal test functions of 23 standard benchmarks

Table 2

Unimodal and simple multimodal functions of CEC2014

Functions	Dim	Range	f_{min}
$f_{14}(x)$ = Rotated High Conditioned Elliptic Function	n	$[-100, 100]$	100
$f_{15}(x)$ = Rotated Bent Cigar Function	\boldsymbol{n}	$[-100, 100]$	200
$f_{16}(x)$ = Shifited and Rotated(SR) Ackley's Function	\boldsymbol{n}	$[-100, 100]$	500
$f_{17}(x)$ = SR Weierstrass Function	n	$[-100, 100]$	600
$f_{18}(x) = SR$ HappyCat Function	\boldsymbol{n}	$[-100, 100]$	1300
$f_{10}(x)$ = SR HGBat Function	\boldsymbol{n}	$[-100, 100]$	1400
$f_{20}(x)$ = SR Expanded Griewank's plus Rosenbrock's Function	n	$[-100, 100]$	1500

Table 3

Hybrid and Composition functions of CEC 2014

Functions	Dim	Range	f_{min}
$f_{22}(x)$ = Hybrid Function 1	\boldsymbol{n}	$[-100, 100]$	1700
$f_{23}(x)$ = Hybrid Function 2	\boldsymbol{n}	$[-100, 100]$	1800
$f_{24}(x)$ = Hybrid Function 3	\boldsymbol{n}	$[-100, 100]$	1900
$f_{25}(x)$ = Hybrid Function 4	\boldsymbol{n}	$[-100, 100]$	2000
$f_{26}(x)$ = Hybrid Function 5	\boldsymbol{n}	$[-100, 100]$	2100
$f_{27}(x)$ = Hybrid Function 6	\boldsymbol{n}	$[-100, 100]$	2200
$f_{28}(x)$ = Composite function 1	\boldsymbol{n}	$[-100, 100]$	2300
$f_{29}(x)$ = Composite function 2	\boldsymbol{n}	$[-100, 100]$	2400
$f_{30}(x)$ = Composite function 3	\boldsymbol{n}	$[-100, 100]$	2500
$f_{31}(x)$ = Composite function 4	\boldsymbol{n}	$[-100, 100]$	2600
$f_{32}(x)$ = Composite function 5	\boldsymbol{n}	$[-100, 100]$	2700
$f_{33}(x)$ = Composite function 6	\boldsymbol{n}	$[-100, 100]$	2800

Table 4

Parameter settings of counterparts

Algorithm	Parameter settings
WOA	$a_1 = [2,0]$; $a_2 = [-2,-1]$; $b = 1$
GWO	$a = [2,0]$
MFO	$b = 1$; $t = [-1,1]$; $a \in [-1,-2]$
BA	$A = 0.5: r = 0.5$
SCA	$A=2$
FA	$\alpha = 0.5; \ \beta = 0.2; \ \nu = 1$
PSO	$c_1 = 2$: $c_2 = 2$: $vMax = 6$
SSA.	$c_1 \in [0 \; 1]; c_2 \in [0 \; 1];$
MVO	existence probability $\in [0.2 1]$; travelling distance rate $\in [0.6 1]$
ALO	$k = 500$
PBIL	learning rate = 0.05 ; elitism parameter = 1;
	probability vector mutation rate $= 0$
DE	scaling $factor = 0.5$; crossover probability = 0.5

All algorithms were performed under the same conditions to achieve fairness in comparative experiments. Among them, the population was set to 30, the dimension and the iteration time was set to 30 and 1000 respectively. To reduce the impacts of random factors in the algorithm on the results, all the compared algorithms were run individually 30 times in each function and averaged as the final running result. On the purpose of measuring experiment results, Standard deviation (STD), Average results (AVG) and Median (MED) were employed to evaluate the results. Note that best results will be bolded (take one in the case of juxtaposition).

3.2.1 Exploitation competence analysis

The data in **Table 5** demonstrates that SMA ranked first or tied first on average when solving F1-5, F7, and F14. The convergence curves of F2 and F5 in **Figure 9** can be visually observed that SMA has the fastest convergence trend among all the comparative functions. The data in **Table 6** demonstrates that SMA can still exhibit significant advantages even when compared to a modified Ma, such as ranking first among other unimodal functions other than F5 and F14. These functions are unimodal functions in the benchmarks, reflecting SMA's efficient exploration capability. Moreover, in order to more fairly evaluate the local search efficiency of the algorithm, an evaluation version of the experiment has been added. The data in **Table 7** demonstrate the experimental results obtained by 300,000 evaluations of the SMA with 10 other participants on the unimodal functions. In the experimental results, the values obtained by SMA were still better than those of other algorithms on F1-5 and F7. At the same time, the median values of the solutions were also consistent with the ranking of the optimal values, indicating the stability of the SMA.

Table 6

Comparison results on the unimodal functions with advanced algorithms

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Comparison results on unimodal functions during 3E5 evaluations

3.2.2 Exploration competence analysis

The data in **Table 8** represents that SMA is still competitive in multimodal functions. In F8-F11

and F20-21, the AVG of SMA was the smallest or the smallest in parallel compared with other algorithms. From the convergence curves of F8 and F21 in **Figure 9**, it can be observed that SMA can search for the highest accuracy fitness value in these two multimodal functions, while some algorithms fail to obtain a superior solution after a certain amount of iterations. This is due to local optima stagnation, which illustrates that SMA can still show better exploration ability in case of preferable exploration. From the data in **Table 9**, it can be seen that the results of SMA in F9-F11, F17, and F20-21 are optimal, and only slightly lower than other algorithms in F8, F18, and F19, which indicates that SMA can still maintain its advantages over advanced algorithms and reflect SMA's capability to avoid local optimum solutions. **Figure 10** also shows that SMA can find a superior solution at a relatively fast convergence tendency in multimodal functions such as F9-11, F17, and F21. **Table 10** illustrates the experimental results of SMA with 10 other comparators on the multimodal function. Among them, SMA obtained the best average and median results on F8-F11 compared with other algorithms, and AGA obtained the best average and median on F16-21. Compared with AGA, SMA has a greater advantage in unimodal functions, while AGA has a preferable performance in multimodal functions.

Table 9

Comparison results on the multimodal functions with advanced algorithms

Table 10

Comparison results on multimodal functions during 3E5 evaluations

3.2.3 Analysis of avoiding locally optimal solutions

All functions in **Tables 11-12**, as fix-dimension multimodal functions, have multiple local optima, which are challenging for MAs, thus can discriminate the overall efficacy of algorithms in exploration and exploration. According to the data in **Tables 11-12**, SMA ranked first in AVG on F28, F29, F30, F32, and F33, which show a very potential comprehensive ability. It can also be seen from the optimum curve of F28-33 in **Figures 9** that SMA achieves superior solutions faster than other counterparts, thus well coordinating the ability of exploration and exploration. The statistics of **Tables 13-14** illustrate that SMA can also maintain certain advantages in composition functions compared with the advanced algorithm, which further reflects that SMA can avert falling into local optimum with fast convergence. F25, F32, and F33 in **Figure 10** also intuitively incarnate the

performance preponderance of SMA in composition functions.

fable

Comparison results on the Hybrid functions of CEC 2014 with traditional algorithms

Table 12

Comparison results on composite functions of CEC2014 with traditional algorithms

Table 13

Comparison results on the Hybrid functions of CEC 2014 with advanced algorithms

Table 14

Comparison results on composite functions of CEC2014 with advanced algorithms

Figure 9 Comparisons between SMA and traditional MAs

Figure 10 Comparisons between SMA and advanced MAs

3.2.4 Significance of superiority analysis

Wilcoxon sign-rank test method [66] was exerted to verify whether SMA has obvious advantages over pairwise comparison. If the p-value produced by the comparison is below the significant level of 0.05 in this case, it means that the achievements of the algorithm in pairwise comparison have obvious superiority in the statistical sense. Otherwise, it is considered that the discrepancies between the two contestants are inconspicuous in a statistical sense. In order to draw further comprehensive conclusions and control the family-wise error rate (FWER), the true statistical significance (#TSS) of

the combined pairwise comparison is shown in Eq 3.1 [67]:

$$
p = 1 - \prod_{i=1}^{k-1} 1 - p_{H_1}
$$
 (3.1)

The p value achieved from this expression is shown in **Table 15**, where the TSS in F1-8, F10, F12, F15, F28-30, and F32-33 were all less than 0.05 when compared with traditional algorithms. Therefore, SMA has significant differences on these functions compared to the traditional algorithms. TSS in **Table 15** when compared with advanced algorithms indicates that SMA outperforms other algorithms in F1-8, F10, F17, F19-22, F25, F28-30, F32-33.

Although pairwise comparisons can be used for comparisons between algorithms, the FWER generated during the experiment cannot be corrected in advance, and the choice of algorithms in multiple comparisons can greatly affect the results of the analysis. In order to reduce the effect of algorithm selection in each result set, multiple comparison processes are used to modify FWER. In multiple comparisons, first check whether the results obtained by the algorithm are unequal. When inequality exists, then perform post-hoc analysis to know which algorithms have significant differences. Therefore, non-parametric Friedman's test [68] was utilized. **Table 16** illustrates the average ranking of the results of the algorithms on the benchmarks compared in the three sets of experiments. In a non-hypothesis, there is equality between all algorithms, so if the hypothesis is reversed, it means that there are differences between the algorithms being compared. Then we chose Holm 's test [69] as the method of post hoc analysis, which is a multiple comparison method that can be used for control algorithms. Using the z-value obtained in **Table 17** to find the corresponding p-value from the normal distribution table and compare it with the corrected *α* value. Take SMA as a control algorithm and compared it with other algorithms. The p-values have been sorted according to their significance. If the p-value is lower than the corresponding significant level α , the corresponding hypothesis is reversed, that is, the algorithm is significantly different. This paper selected two significant level $\alpha = 0.10$ and $\alpha = 0.05$, which indicate that there are marginal and significant differences between the two methods. As can be seen from **Table 17**, compared with the traditional algorithms other than DE, the z-value is smaller than the corrected value with $\alpha = 0.05$ as the significant level, that is, there are significant differences in benchmarks. Compared with advanced algorithms other than LWOA, there are significant differences among the benchmark functions, and slightly different from LWOA. In the experiments with other algorithms in the evaluation version, SMA is slightly different compared to GWO and WOA, not significantly different from AGA and DE, while significantly different from the remaining algorithms.

	True p-value obtained from comparison on unity-unce benemiativis											
		F1	F ₂	F3	F ₄	F5	F6	F7	F8	F9	F10	F11
	#TSS	2.08E-05	2.08E-05	2.08E-05	2.08E-05	2.08E-05	4.13E-03	2.68E-05	1.34E-04	$1.00E + 00$	2.31E-05	$1.00E + 00$
		F12	F13	F14	F15	F ₁₆	F17	F ₁₈	F19	F20	F21	F22
Traditional MAS	#TSS	3.71E-05	5.23E-02	5.53E-02	6.34E-03	7.95E-01	1.32E-01	3.26E-01	2.24E-01	1.98E-01	6.88E-02	7.01E-01
		F23	F24	F25	F26	F27	F28	F ₂₉	F30	F31	F32	F33
	#TSS	4.53E-01	7.59E-01	9.96E-01	5.34E-01	4.96E-01	2.29E-05	2.08E-05	$2.63E-04$	4.36E-01	2.08E-05	2.16E-05
Adv		F ₁	F2	F3	F4	F5	F6	F7	F8	F9	F10	F11
نم	#TSS	1.56E-05	1.56E-05	1.56E-05	1.56E-05	1.85E-02	1.40E-03	2.60E-03	1.60E-05	1.00E+00	2.74E-05	$1.00E + 00$

Table 15 True p-value obtained from comparison on thirty-three benchmarks

nced

Table 166

Results of Friedman test of iterative version and function evaluation version

Table 177

Holms' test (take SMA as the control algorithm)

3.3 Wall-clock time analysis

In this section of the experimentations, SMA was compared with other 11 participants in the calculation of time-consuming experiments in the 33 benchmarks mentioned above. The time-consuming calculation approach is that all participants independently run 10 times on each function and recorded the results in **Table 18**. As can be observed from the data in the table, the computation of SMA took relatively longer time, because the calculation of the oscillation factor requires more computing power. However, SMA can still outperform some algorithms while taking less time, such as GOA, DA, and ALO. In general, even if it is relatively time-consuming, SMA still possess tremendous effectiveness advantages over other algorithms, so the time-consuming is worth it.

Table 18 Wall-Clock Time costs of SMA and other candidates on 33 benchmarks

	SMA	SCA	SSA	GWO	MFO	WOA	GOA	DA	ALO	MVO	PSO	DE
F1	14.040	1.310	0.811	1.825	1.513	0.562	119.46	90.262	218.35	3.822	0.967	5.054
F2	13.291	1.139	0.796	1.622	1.342	0.577	118.10	110.88	216.74	3.806	0.920	4.446
F ₃	13.478	2.480	2.215	2.839	2.636	2.278	110.01	116.26	207.60	4.165	1.919	5.288
F4	12.776	1.123	0.796	1.560	1.123	0.546	117.18	86.659	207.51	2.761	0.640	3.838
F ₅	12.995	1.404	1.030	1.950	1.466	0.780	123.64	115.76	233.11	3.401	0.874	4.976
F ₆	15.241	1.513	1.045	2.090	1.544	0.686	146.37	122.92	260.16	3.760	0.889	4.914
F7	16.037	2.356	1.919	2.855	2.465	1.638	141.66	118.85	257.99	4.883	1.794	5.678
F8	15.709	1.763	1.295	2.434	1.778	0.983	141.11	172.03	255.46	2.824	1.264	5.491
F9	16.115	1.576	1.123	2.028	1.669	0.764	142.13	136.65	259.30	3.994	1.139	4.742
F10	14.726	1.794	1.279	2.090	1.997	0.936	143.20	111.15	251.09	4.025	1.092	5.444
F11	16.115	2.215	1.607	2.309	1.981	0.998	144.87	121.33	264.06	4.040	1.217	5.600
F12	19.032	4.602	4.134	5.023	4.836	3.900	149.91	124.59	265.56	6.880	4.134	8.596
F13	18.939	4.852	4.274	5.101	4.243	3.604	149.15	126.20	266.68	6.833	4.087	8.518
F14	16.411	2.090	1.732	2.855	2.340	1.326	140.80	145.54	256.34	4.399	1.544	5.647
F15	15.725	2.106	1.498	2.465	1.950	1.108	144.89	136.08	261.58	4.524	1.373	5.990
F ₁₆	15.803	2.106	1.638	2.652	2.044	1.295	146.76	152.35	261.16	4.243	1.342	5.351
F17	31.715	16.677	15.943	17.254	16.224	15.678	163.83	185.08	281.14	19.438	16.021	20.686
F18	16.177	2.090	1.544	2.621	2.246	1.279	146.37	170.30	263.08	4.508	1.342	5.288

3.4 Parameter sensitivity analysis

In this section, parameter sensitivity test was utilized to evaluate the impacts of population size, iterations and parameter z on the algorithm. The range of parameter z is $[0,0.1]$, and there are 11 values at intervals of 0.01. The population size was set to 5,10,30,50,100 and 200. The number of iterations was set to 50,100,200,500,1000 and 2000. Under other conditions remained, different values of parameter z were tested on F1-13 and the results are shown in **Table 19**. SMA0 indicates that z takes a value of 0, SMA1 indicates that z takes a value of 0.01, and so on. The values in the table are ranking. From the results in **Table 20**, it can be recognized that the result of the algorithm is superior when z was taken as 0.03, because the probability maintains the balance between explosion and exploitation. Experimenter can also take different values for z according to specific problems.

Table 19

Ranking of results with varied values of parameter z

	$\tilde{}$			л.							
Function	SMA0	SMA1	SMA ₂	SMA3	SMA4	SMA5	SMA6	SMA7	SMA8	SMA9	SMA10
						6	$\overline{7}$	8	9	10	11
$\overline{2}$			3	$\overline{4}$	5	6	$\overline{7}$	9	8	10	11
3					5	6	7	8	9	10	11
4			3	$\overline{4}$	5	6	7	8	9	10	11
5	11	10	9	8	7	3	$\overline{4}$	2	6	5	
6	11		$\overline{2}$	3	$\overline{4}$	5	6	9	$\overline{7}$	8	10
7		2	$\overline{4}$	3	6	τ	10	8	9	5	11
8	11		$\overline{4}$	5	6	8	9	$\overline{2}$	$\overline{7}$	10	3
9		-1					1	ı			
10											
11											
12	11	10	9	$\overline{7}$	6	8	5	$\overline{4}$	$\overline{2}$	3	

To explore the influence of populations and iterations on the algorithm, we chose F13 to test the synergistic effect of the two parameters on the algorithm. As can be seen visually from **Figure 11**, as the population size and iterations increased, the average became better. The reason is that the increase in the number of populations improves the search efficiency, and the increase in iterations leads to an incensement in the times of searches and the accuracy of subsequent searches. However, the results were not increased proportionally when the population size and iterations continue to grow due to the global approximate optimal solution has been roughly discovered. Researchers can select the appropriate populations and iterations based on specific questions.

Figure 11 The influence of populations and iterations

3.5 Experiments on engineering design problems

Most problems have constraints in the real production environment. The process of considering constraints of equality and inequality during optimization is called constraints processing. The candidate solutions of the heuristic algorithm can be divided into feasible and infeasible according to the constraints. There are currently several types of constraint methods: death penalty, annealing, static, dynamic, co-evolutionary, and adaptive. Although useful information may be lost in the process of abandonment, we still adopted a relatively simple method of the death penalty with low computational cost to deal with search individuals who violated constraints and then re-assigned them a relatively large target value.

In the following sections, SMA is tested four engineering-constrained design problems: a welded beam problem, a pressure vessel problem, a cantilever, and I-beam.

3.5.1 Welded beam structure problem

The main purpose of the problem is to constrain side constraints, end deflection of the beam (δ) , buckling load on the bar(P_c), bending stress in the beam(θ), moreover, shear stress(τ) with the least economic cost of welded beams.

There are four variables, for instance, the thickness of the weld(h), length of the attached bar(l), the height of the bar(t), the thickness of the bar(b). The design diagram for this problem is shown in **Figure 12**. The formulations were list below:

Consider:

$$
X = [x_1, x_2, x_3, x_4] = [h l t b]
$$

Minimize:

$$
F(X) = 1.10471x_1^2x_2 + 0.04811x_3x_4(14.0 + x_2)
$$

Subject to:

$$
g_1(X) = \tau(X) - \tau_{max} \le 0
$$

\n
$$
g_2(X) = \sigma(X) - \sigma_{max} \le 0
$$

\n
$$
g_3(X) = \delta(X) - \delta_{max} \le 0
$$

\n
$$
g_4(X) = x_1 - x_4 \le 0
$$

\n
$$
g_5(X) = P - P_C(X) \le 0
$$

\n
$$
g_6(X) = 0.125 - x_1 \le 0
$$

\n
$$
g_7(X) = 1.10471x_1^2 + 0.04811x_3x_4(14.0 + x_2) - 5.0 \le 0
$$

where

$$
\tau(\vec{x}) = \sqrt{(\tau')^2 + 2\tau'\tau'' \frac{x_2}{2R} + (\tau'')^2}
$$

$$
\tau' = \frac{P}{\sqrt{2}x_1x_2}, \tau'' = \frac{MR}{J}, M = P(L + \frac{x_2}{2})
$$

$$
R = \sqrt{\frac{x_2^2}{4} + \frac{x_1 + x_3}{2}}?
$$

$$
J = 2\left\{\sqrt{2}x_1x_2\left[\frac{x_2^2}{4} + \frac{x_1 + x_3}{2}\right]^2\right\}
$$

$$
\sigma(\vec{x}) = \frac{6PL}{x_4x_3^2}, \delta(\vec{x}) = \frac{6PL^3}{Ex_3^2x_4}
$$

$$
P_C(\vec{x}) = \frac{4.013E\sqrt{\frac{x_3^2x_4^6}{36}}}{L^2} (1 - \frac{x_3}{2L}\sqrt{\frac{E}{4G}})
$$

 $P = 60001b, L = 14in.$ $\delta_{max} = 0.25$ in $E = 30 \times 1^6 \text{psi}, G = 12 \times 10^6 \text{psi}$ $\tau_{max} = 13600\pi\eta$, $\sigma_{max} = 30000\pi\eta\eta$

In this problem, SMA was compared with MFO[23], SSA[52], Random[70], Siddall[71], Ragsdell[70], Coello and Montes[72], GWO[21], WOA[50], GSA, Simplex[70] and David[70]. **Table 20** illustrates that SMA can obtain the optimal value.

Figure 12 Structure of welded beam design

Table 20

Results of welded beam structure problem compared with other competitors

3.5.2 Pressure vessel structure problem

The intention of the problem is to find the parameters of cylindrical pressure vessels which can minimize the total cost of production and meet the pressure requirements. The parameters including the thickness of the shell (T_s) , inner radius (T_h) , the thickness of the head (T_h) and the length of the cylindrical portion. Both ends of the container are covered with a hemispherical shell at one end. **Figure 13** illustrates the design of the object and its corresponding parameters.

The formulations of four constraints are listed as follow:

Consider:

$$
X = [x_1 \ x_2 \ x_3 \ x_4] = [T_s \ T_h \ R \ L]
$$

Objective:

 $f(X)_{min} = 0.6224x_1x_3x_4 + 1.7781x_3x_1^2 + 3.1661x_4x_1^2 + 19.84x_3x_1^2$

Subject to:

$$
g_1(X) = -x_1 + 0.0193x_3 \le 0
$$

$$
g_2(X) = -x_3 + 0.00954x_3 \le 0
$$

$$
g_3(X) = -\pi x_4 x_3^2 - \frac{4}{3}\pi x_3^3 + 1296000 \le 0
$$

$$
g_4(X) = x_4 - 240 \le 0
$$

Variable ranges:

Table 21

$$
0 \le x_1 \le 99, 0 \le x_2 \le 99, 10 \le x_3 \le 200, 10 \le x_4 \le 200
$$

From the data of **Table 21**, it is obvious that SMA can obtain fairly superior optimal values compared with MFO[23], BA[73], HPSO[74], CSS[5], CPSO[75], ACO[76], GWO[21], WOA[50], MDDE[77], Lagrangian multiplier[78] and Branch-bound[79].

Figure 13 Structure of pressure vessel

3.5.3 Cantilever structure problem

The cantilever beam is made up of five hollow square cross-sections, as exhibited in **Figure 14**. Since the thickness is fixed, only six parameters identified in the figure need to be considered. The intention of the problem is to dwindle the total mass of the cantilever beam when the bearing

capacity is satisfied. The formulas of this optimization problem are listed as follow: Consider:

$$
X = [x_1 \ x_2 \ x_3 \ x_4 \ x_5]
$$

Minimize:

$$
F(X) = 0.6224(x_1 + x_2 + x_3 + x_4 + x_5)
$$

Subject to:

$$
G(X) = \frac{61}{x_1^3} + \frac{37}{x_2^3} + \frac{19}{x_3^3} + \frac{7}{x_4^3} + \frac{1}{x_5^3} \le 1
$$

Variable ranges:

 $0.01 \leq x_1, x_2, x_3, x_4, x_5 \leq 100$

Figure 14 Structure of cantilever beam

Compared to MFO[23], SOS[80], CS[81], MMA[82] and GCA[82], SMA can achieve better results in the cantilever design problem, as shown in **Table 22**.

Table 22

Results of cantilever beam structural problem in comparison with other competitors

3.5.4 I-beam structure problem

The intention of this engineering problem is to minimize the vertical deviation of I-beam by adjusting four parameters as shown in **Figure 15**.

Figure 15 Structure of I-beam

Table 23 lists the optimization results for SMA compared to ARSM[83], SOS[80], CS[81] and IARSM[83]. The data reveals that SMA can obtain excellent optimal values in this engineering problem, reflecting the applicability of SMA to engineering problems. **Table 23**

		Optimum values for variables							
Algorithm	\boldsymbol{b}	h		$T_{\scriptscriptstyle\mathsf{F}}$	cost				
SMA	49.998845	79.994327	1.764747	4.999742	0.006627				
ARSM[83]	37.0500	80.0000	1.7100	2.3100	0.0157				
SOS[80]	50.0000	80,0000	0.9000	2.3218	0.0131				
CS[81]	50.0000	80,0000	0.9000	2.3217	0.0131				
IARSM[83]	48.4200	79.9900	0.9000	2.4000	0.1310				

Results of I-beam structural problem in comparison with other methods

4 Conclusions and future perspectives

This paper proposed a brand-new metaheuristic enlightened by slime behavior to tackle the optimization problem. The algorithm mainly uses the weights to simulate the positive and negative feedback of the bio-oscillator during the foraging to the food source to form a different thickness of the feeding vein network. The morphology of the slime mould also changes with three different contraction patterns.

To qualitatively analyze the algorithm, four metrics (search history, the trajectory of the first dimension, average fitness, and convergence curve) were applied. Then, the algorithm was evaluated in 33 benchmark functions consisting of unimodal, multimodal, fix-dimension multimodal, and composite functions. Most of the functions tested are composite functions. Wilcoxon sign-rank test and Freidman test were applied to estimate the efficacy of the algorithm more scientifically. The experimental results illustrate that SMA can guarantee the performance of explorations while achieving superior exploitations, thus maintaining an outstanding balance between exploitations and explorations, which reflects the superior performance of the algorithm in a statistical sense compared with other algorithms.

Meanwhile, SMA was used in four classical engineering structural problems, including welded beam, pressure vessel, cantilever, and I-beam design problems. The results demonstrate that SMA is also applicable to engineering optimization problems in real life with satisfactory optimization results.

The accounts for the satisfactory performance of SMA in maintaining the balance of exploitation and exploration can be theoretically attributed to the following points:

- The adaptive weight *W* enables the SMA to maintain a certain disturbance rate while guaranteeing fast convergence, thus avoiding optimal local trapping during fast convergence.
- Vibration parameter \overrightarrow{vb} allows the individual position of slime mould to contract in a specific way, thus ensuring the efficiency of the early exploration and the accuracy of the later exploitation.
- The adequate utilization of individual fitness values allows SMA to make better decisions based on historical information.
- The location updating decision parameter *p* and three different location updating methods ensure better adaptability of the SMA in different search phases.

On the purpose of improving the extensibility of the algorithm, the development of the algorithm is established on the principle of being as simple as possible. In future work, various mutation mechanisms or acceleration mechanisms can be employed to enhance the efficacy of the algorithm. The binary version of the algorithm can also be developed for feature selection. Moreover, SMA can also be used to optimize parameters of classifiers such as kernel extreme learning machine or support vector machine.

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