

Slime Mould Algorithm: A New Method for Stochastic Optimization

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Abstract

Slime mould algorithm (SMA) is proposed based on 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 and to form the optimal path for connecting food with excellent exploratory ability and exploitation propensity. Source codes of SMA are publicly available at <http://www.alimirjalili.com/SMA.htm> and <https://tinyurl.com/Slime-mould-algorithm>.

Keywords: Slime mould algorithm, Adaptive weight, Engineering design problems, Constrained optimization

1. An Overview of slime mould algorithm

For complete details, please refer to main paper¹ established by [1].

1.1. Approach food

To model the approaching behavior of slime mould as a mathematical equation, the following rule is proposed to imitate the contraction mode:

$$X_{t+1} = \begin{cases} X_b(t) + v_b \cdot (W \cdot X_A(t) - X_B(t)) & r < p \\ v_c \cdot X_t & r \geq p \end{cases} \quad (1)$$

where v_b is a parameter with a range of $[-a, a]$, v_c 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 the swarm, W represents the weight of slime mould. The formula of p is as follows:

$$p = \tanh |S(i) - DF| \quad (2)$$

¹The paper is available at <https://doi.org/10.1016/j.future.2020.03.055> and <https://tinyurl.com/Slime-mould-algorithm>.

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

The formula of v_b is as follows:

$$v_b = [-a, a] \quad (3)$$

$$a = \operatorname{arctanh} \left(-\left(\frac{t}{max_t} \right) + 1 \right) \quad (4)$$

The formula of W is listed as follows:

$$W(SmellIndex(i)) = \begin{cases} 1 + r \log((b_F - S(i))/(b_F - w_F) + 1) & \text{condition} \\ 1 - r \log((b_F - S(i))/(b_F - w_F) + 1) & \text{others} \end{cases} \quad (5)$$

$$SmellIndex = sort(S) \quad (6)$$

where *condition* indicates that $S(i)$ ranks first half of the population, r denotes the random value in the interval of $[0, 1]$, b_F denotes the optimal fitness obtained in the current iterative process, w_F 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).

1.2. Wrap food

The mathematical formula for updating the location of slime mould is as follows:

$$X^* = \begin{cases} rand(UB - LB) + LB & rand < z \\ X_b(t) + v_b(WX_A(t) - X_B(t)) & r < p \\ v_c X(t) & r \geq p \end{cases} \quad (7)$$

where LB and UB denote the lower and upper boundaries of the search range, $rand$ and r denote the random value in $[0, 1]$.

1.3. Oscillation

The value of v_b oscillates randomly between $[-a, a]$ and gradually approaches zero as the iterations increase. The value of v_c oscillates between $[-1, 1]$ and tends to zero eventually.

The logic of SMA is shown in Fig. 1 and its Pseudo-code represented in Algorithm 1.

Algorithm 1 Pseudo-code of SMA algorithm

Inputs: The population size N and maximum number of iterations max_t

Outputs: The best solution

Initialize the the positions of slime mould $X_i (i = 1, 2, \dots, n)$

while (Stopping condition is not met) **do**

 Calculate the fitness of all slime mould

 Calculate the W by Eq. (5)

for (each search portion (X_i)) **do**

 Update p, v_b, v_c ;

 Update positions by Eq. (7)

Return bestFitness and X_b

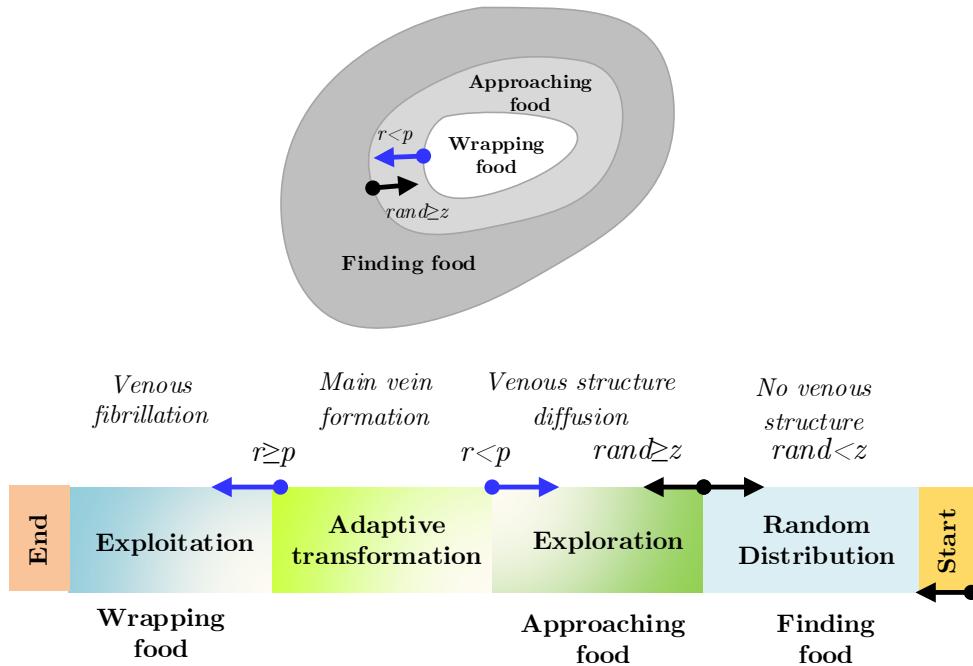


Figure 1: The overall steps of SMA

References

- [1] S. Li, H. Chen, M. Wang, A. A. Heidari, S. Mirjalili, Slime mould algorithm: A new method for stochastic optimization, Future Generation Computer Systems (2020). doi:<https://doi.org/10.1016/j.future.2020.03.055>.