# 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 paper<sup>1</sup> 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:

$$X_{t+1} = \begin{cases} X_b(t) + v_b \cdot (W \cdot X_A(t) - X_B(t)) & r (1)$$

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:

$$p = \tanh |S(i) - DF| \tag{2}$$

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

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

The formula of  $v_b$  is as follows:

$$v_b = [-a, a] \tag{3}$$

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

The formula of W is listed as follows:

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

$$SmellIndex = sort(S)$$
 (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:

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

$$(7)$$

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 positions of slime mould  $X_i (i = 1, 2, ..., n)$ 

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  $X_b$ 

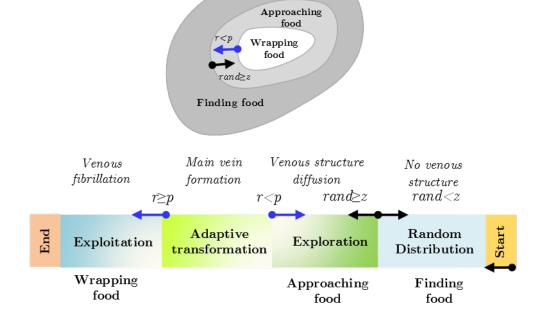


Figure 1: The overall steps of SMA  $\,$ 

# 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.