# **Session: Bacterial Foraging Algorithm**

Course Title: Computational Intelligence
Course Code: 19MIE501A

#### Course Leader: Dr. Vaishali R. Kulkarni

Assistant Professor, Department of Computer Science and Engineering Faculty of Engineering and Technology
Ramaiah University of Applied Sciences, Bengaluru
Email: vaishali.cs.et@msruas.ac.in

Tel: +91-804-906-5555 Ext:2222

Website: http://www.msruas.ac.in/staff/fetcseVaishali



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- 2. The foraging theory and foraging strategies

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- 3. Social and intelligent foraging
- 4. Chemotaxis, elimination and dispersal and reproduction in E. Coli bacteria
- 5. The bacterial foraging optimization; and algorithm and all its important parameters



At the end of this session, the student will be able to:

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- 3. Appreciate E.Coli life-cycle as an optimization exercise
- 4. Implement Bacterial Foraging Optimization (BFO) Algorithm to solve a continuous optimization problem
- 5. Choose proper values of algorithm parameters based on the nature of the problem at hand

#### **Recommended Resources for this Session**

- 1. K. M. Passino. (2002). "Biomimicry of bacterial foraging for distributed optimization and control". IEEE Control Systems Journal. volume 22, number 3, pp. 52–67.
- Kulkarni R. V. and Venayagamoorthy, G. K. (2010). "Bioinspired algorithms for autonomous deployment and localization of sensor nodes." *IEEE Transactions on Systems, Man, and Cybernetics, Part C: Applications and Reviews*, issue 6, volume 40, pp. 663–675.

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- This evolutionary principle has inspired scientists in "foraging theory" to model foraging as an optimization process
- An animal attempts to maximize the energy obtained per unit time spent foraging, under the constraints of its own physiology (e.g., sensing and cognitive capabilities) and environment (e.g., density of prey, risks from predators, physical characteristics of the search area)



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- Patches are encountered sequentially, and great effort and risk are needed to travel from one patch to another



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  to deplete its resources, so there should be an optimal time to
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  readily available, but it also does not want to waste time in the
  face of diminishing energy returns
- Optimal foraging theory formulates the foraging as an optimization problem and via computational or analytical methods can provide an optimal foraging policy that specifies how foraging decisions are made





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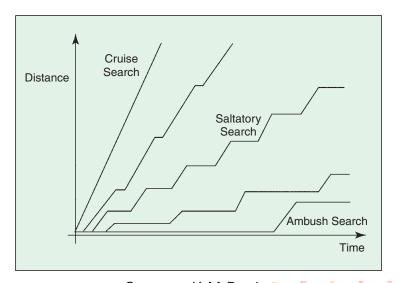
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- Researchers have shown that foraging decision heuristics are used very effectively by animals to approximate optimal policies



# **Search Strategies for Foraging Animals**





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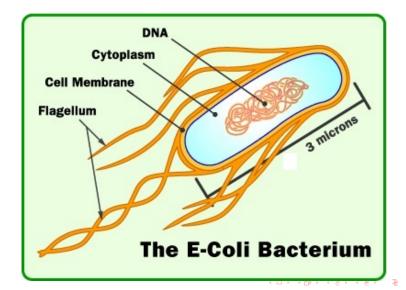
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- Even extremely simple organisms like Bacteria can still work together for the benefit of the group

# Escherichia Coli (E. coli) Bacterium





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- It swims away from alkaline and acidic environments and toward more neutral ones



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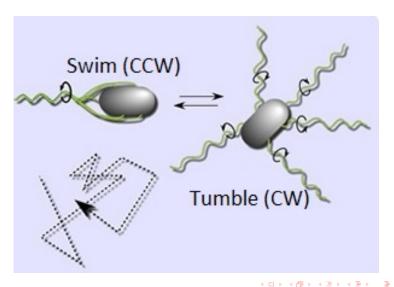
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- After a tumble, the cell is pointed in a random direction







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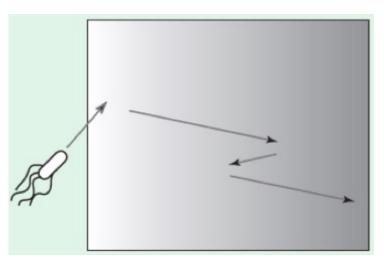
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- When it reaches constant nutrient concentration, it returns to baseline behavior. It is never satisfied with the amount of surrounding food; it always seeks higher concentrations



## E. coli Chemotaxis





Courtesy: K M Passino

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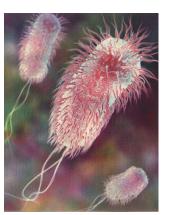
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- Elimination and dispersal are parts of the population-level longdistance motile behavior





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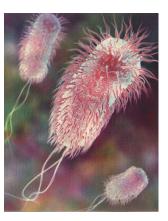


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- The goal is to achieve better fitness in a way similar to natural bacteria that thrive to move to higher concentration of nutrients



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- Let  $J(P_i)$  represent an objective function. Let  $J(P_i) < 0$ ,  $J(P_i) = 0$  and  $J(P_i) > 0$  represent the bacterium at location  $P_i$  in nutrient rich, neutral and noxious environments respectively



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- Chemotaxis is a foraging behavior that captures the process of optimization where bacteria try to achieve positions having lower values of  $J(P_i)$ , and avoid being at positions  $P_i$  where  $J(P_i) \ge 0$



# Chemotaxis and Reproduction in BFOA

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- This swimming is continued until a minimum fitness value is reached, but only for a maximum number of steps  $N_s$
- After  $N_c$  chemotactic steps, a reproduction step is taken, in which the population is sorted in ascending order of the objective function value J and the least healthy bacteria are replaced by the copies of the healthier bacteria



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- After  $N_{re}$  reproduction steps, an elimination-dispersal step is taken
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- ullet The optimization stops after  $N_{ed}$  elimination-dispersal rounds

## **Swarming in BFOA**

• Bacteria create swarms by means of cell-to-cell signalling via an attractant and a repellent. Cell-to-cell attraction for bacterium i is represented with  $J_{cc}(\mathcal{P}, P_i), i = 1, 2, \cdots, S$ 

$$J_{cc}(\mathscr{P}, P_i) = \sum_{t=1}^{S} \left[ -d_a exp \left( -w_a \sum_{m=1}^{p} (P_{i,m} - P_{t,m})^2 \right) \right]$$
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• Here,  $J_{cc}(P_i, \mathscr{P})$  denotes the combined cell-to-cell attraction and repulsion effects for bacterium i at position  $P_i = [P_{i,1}, P_{i,2}, \cdots, P_{i,p}]^T$  and the whole swarm of bacteria  $\mathscr{P} = \{P_1, P_2, \cdots, P_S\}$ 



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- For BFOA, the maximum number of objective function evaluations is  $N_{ed} \cdot N_{re} \cdot N_c \cdot S \cdot N_s$
- A general biologically inspired rule of thumb for choosing the parameters of BFOA is:  $N_c > N_{re} > N_{ed}$

#### **BFOA** Pseudocode

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- Important parameters: Numbers of elimination and dispersion rounds, reproduction rounds, chemotaxis rounds, swimming rounds



# **Any Questions?**





# Thank You

