

Ch.3. GAs : Why Do They Work?

Schema

- **Definition**

a similarity template describing a subset of strings with similarities at certain string of positions

don't care symbol: *

- **Example**

schema (*0000) matches 2 strings {(00000), (10000)}

schema (*111*) matches 4 strings {(01110), (01111), (11110), (11111)}

schema (10100) matches 1 string {(10100)}

- **Properties**

1) Every schema matches 2^r strings (r : no. of don't care symbols)

2) Each string of the length m is matched by 2^m schema

(ex) string (101) , $m=3$

schema: (101),
 (*01), (1*1), (10*),
 (**1), (*0*), (1**),
 (***)

3) strings of length $m \Rightarrow 2^m$ different strings, 3^m possible schema

(ex) $m=3$

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strings: $2(0/1) \times 2 \times 2 = 8$

schema: $3(0/1/*) \times 3 \times 3 = 27$

4) Number of possible schema in a population of size $n \Rightarrow 2^m \sim n 2^m$

(ex) $m=3$

1	0	1
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 $\rightarrow 2^3$ schema

.....

0	0	1
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 $\rightarrow 2^3$ schema

Schema Properties

(1) Schema order : $o(S)$

- number of fixed positions
- 0 / 1 의 개수

(ex)

$$S1 = (0\ 1\ 1\ * \ 1\ * \ *) : o(S1) = 4$$

$$S2 = (0\ * \ * \ * \ * \ * \ *) : o(S1) = 1$$

- mutation 에 대한 schema 의 survival probability 계산에 사용

(2) Defining length : $\delta(S)$

- distance between the first and the last fixed positions

(ex)

$$\delta(S1) = 5-1 = 4$$

$$\delta(S2) = 1-1 = 0$$

- crossover 에 대한 schema 의 survival probability 계산에 사용

Reproductive Schema Growth Equation

(1) Reproduction 고려

- $\xi = \xi(S, t)$

: number of strings in a population at time t , matched by schema S

(ex) population at time t

$$v_1 = (0100) \rightarrow eval(v_1) = 15$$

$$v_2 = (1110) \rightarrow eval(v_2) = 20$$

$$v_3 = (1011) \rightarrow eval(v_3) = 5$$

$$v_4 = (1111) \rightarrow eval(v_4) = 10$$

$$S_0 = (*11*)$$

$$\Rightarrow \xi(S_0, t) = 2$$

- $eval(S, t)$, **schema fitness**

: average fitness of all strings in a population at time t , matched by schema S

$$(ex) eval(S_0, t) = \frac{eval(v_2) + eval(v_4)}{2} = 15$$

- **Schema growth equation**

$$\xi(S, t+1) = \xi(S, t) \cdot \text{popsize} \cdot \frac{\text{eval}(S, t)}{F(t)}$$

↯, $F(t)$: total fitness of the population at time t

let $\bar{F}(t) = \frac{F(t)}{\text{popsize}}$: average fitness,

$$\xi(S, t+1) = \xi(S, t) \cdot \frac{\text{eval}(S, t)}{\bar{F}(t)}$$

=> reproductive schema growth eq.

let $\xi(S, t) = \bar{F}(t) + \epsilon \bar{F}(t)$

$$\xi(S, t) = \xi(S, 0)(1 + \epsilon)^t$$

$\epsilon > 0$: above average, exponentially increasing

$\epsilon < 0$: below average, exponentially decreasing

(ex) continued

$$\begin{aligned} \xi(S_0, t+1) &= \xi(S_0, t) \frac{\text{eval}(S_0, t)}{\bar{F}(t)} \\ &= 2 \times \frac{15}{12.5} = 2.4 \end{aligned}$$

$$\epsilon = \frac{15 - 12.5}{12.5} = 0.2$$

$$\xi(S, t) = 2 \times 1.2^t$$

(ex)

Population at time t

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v1 = (100110100000001111111010011011111)
v2 = (111000100100110111001010100011010)
v3 = (000010000011001000001010111011101)
v4 = (10001100010110100111100000110010)
v5 = (00011101100101001101011111000101)
v6 = (00010100001001010100101011111011)
v7 = (00100010000011010111101101111011)
v8 = (10000110000111010001011010110011)
v9 = (0100000010110001011000001111100)
v10 = (000001111000110000011010000111011)
v11 = (011001111110110101100001101111000)
v12 = (11010001011110110100010101000000)
v13 = (111011111010001000110000001000110)
v14 = (010010011000001010100111100101001)
v15 = (111011101101110000100011111011110)
v16 = (110011110000011111100001101001011)
v17 = (01101011111100111101000110111101)
v18 = (011101000000001110100111110101101)
v19 = (000101010011111111110000110001100)
v20 = (101110010110011110011000101111110)

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- pop_size = 20, length of string $m=33$

Schema S_0

$S_0 = (*****111*****)$

- $\xi(S_0, t) = 3$ (v_{13}, v_{15}, v_{16})

- $eval(S_0, t) = (27.316702 + 30.060205 + 23.867227)/3 = 27.081378$

$\bar{F}(t) = \sum_{i=1}^{20} eval(v_i) / popsize = 387.776822/20 = 19.388841$

$eval(S_0, t) / \bar{F}(t) = 1.396751$

☞ $\xi(S_0, t+1) = 3 \times 1.396751 = 4.19, \xi(S_0, t+2) = 3 \times 1.396751^2 = 5.85, \dots$

Population at time t+1

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v'1 = (011001111110110101100001101111000) (v11)
v'2 = (10001100010110100111100000110010) (v4)
v'3 = (00100010000011010111101101111011) (v7)
v'4 = (011001111110110101100001101111000) (v11)
v'5 = (00010101001111111110000110001100) (v19)
v'6 = (10001100010110100111100000110010) (v4)
v'7 = (111011101101110000100011111011110) (v15)
v'8 = (000111011001010011010111111000101) (v5)
v'9 = (011001111110110101100001101111000) (v11)
v'10 = (000010000011001000001010111011101) (v3)
v'11 = (111011101101110000100011111011110) (v15)
v'12 = (010000000101100010110000001111100) (v9)
v'13 = (00010100001001010100101011111011) (v6)
v'14 = (10000110000111010001011010110011) (v8)
v'15 = (101110010110011110011000101111110) (v20)
v'16 = (111001100110000101000100010100001) (v1)
v'17 = (111001100110000100000101010111011) (v10)
v'18 = (111011111010001000110000001000110) (v13)
v'19 = (111011101101110000100011111011110) (v15)
v'20 = (110011110000011111100001101001011) (v16)

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5 strings ($v'_{7}, v'_{11}, v'_{18}, v'_{19}, v'_{20}$) are matched with the schemata S_0

(2) Crossover 고려

- δ (defining length) 가 작은 schema 가 crossover 시 survival probability 가 높다.

$$\text{(ex)} \quad S_1 = (*1****0) : \delta(S_1) = 7 - 2 = 5$$

$$S_2 = (***10**) : \delta(S_2) = 5 - 4 = 1$$

crossover position = 3 경우

=> S_1 : destroy, S_2 : survive

- **Destruction probability of a schema**

$$p_d(S) = \frac{\delta(S)}{m-1}$$

$$\text{(ex)} \quad p_d(S_1) = \frac{5}{6}, \quad p_d(S_2) = \frac{1}{6}$$

- **Survival probability of a schema**

$$p_s(S) = 1 - p_d(S) = 1 - \frac{\delta(S)}{m-1}$$

$$\text{(ex)} \quad p_s(S_1) = \frac{1}{6}, \quad p_s(S_2) = \frac{5}{6}$$

- **Crossover 선택 확률 p_c 고려**

$$p_d(S) = p_c \cdot \frac{\delta(S)}{m-1}$$

$$p_s(S) = 1 - p_c \cdot \frac{\delta(S)}{m-1}$$

- **Mate (pair) 고려**

$$p_s(S) \geq 1 - p_c \cdot \frac{\delta(S)}{m-1}$$

$$\text{(ex)} \quad S_1 = (111* \updownarrow **00)$$

$$S_2 = (100* \updownarrow **00)$$

- Reproduction 과 Crossover를 고려한 schema growth equation

$$\begin{aligned}\xi(S, t+1) &= \xi(S, t) \cdot \frac{eval(S, t)}{\bar{F}(t)} \cdot p_s(S) \\ &\geq \xi(S, t) \cdot \frac{eval(S, t)}{\bar{F}(t)} \cdot \left(1 - p_c \frac{\delta(S)}{m-1}\right)\end{aligned}$$

(ex) continued

$$\delta(S_0) = 1, p_c = 0.25, m = 5,$$

$$\Rightarrow \frac{eval(S_0, t)}{\bar{F}(t)} \left(1 - p_c \frac{\delta(S_0)}{m-1}\right) = \frac{15}{12.5} \left(1 - 0.25 \frac{1}{4}\right) = 1.205$$

$$\xi(S_0, 0) = 2$$

$$\xi(S_0, 1) \geq 2 \times 1.205 \approx 2.4$$

$$\xi(S_0, 2) \geq 2 \times 1.205^2 \approx 2.9$$

$$\xi(S_0, 3) \geq 2 \times 1.205^3 \approx 3.5$$

(3) Mutation 고려

- o (order) 가 작은 schema 가 crossover 시 survival probability 가 높다.

(ex) $S_1 = (** * 10 **)$: $o(S_1) = 2$

$S_2 = (1011 ** 1)$: $o(S_2) = 5$

- Survival probability of a schema

$$p_s(S) = (1 - p_m)^{o(S)}$$

p_m : mutation probability

*	*	1	*	0	*	*
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$$1 - p_m \quad 1 - p_m$$

if $p_m \ll 1$, $p_s(S) \approx 1 - o(S) \cdot p_m$

- Reproduction, Crossover, Mutation 을 고려한 schema growth equation

$$\xi(S, t+1) \geq \xi(S, t) \cdot \frac{eval(S, t)}{\bar{F}(t)} \cdot (1 - p_c \frac{\delta(S)}{m-1} - o(S)p_m)$$

(ex) continued

$$o(S_0) = 2, \quad p_m = 0.01$$

$$\Rightarrow \epsilon = \frac{15}{12.5} (1 - 0.25 \times \frac{1}{4} - 2 \times 0.01) = 1.101$$

$$\xi(t+k) = 2 \times 1.101^k$$

$$\xi(t+5) \geq 3.2$$

■ Schema Theorem

Short, low-order, above-average schemata receive *exponentially* increasing trials in subsequent generation of a genetic algorithm.

■ Building Block Hypothesis

A genetic algorithm seeks *near-optimal* performance through the juxtaposition of short, low-order, above-average schemata, called the building blocks.