

Genetic and Evolutionary Algorithms: Schema Theorem

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Theoretical Basis of Trust in the Potential of Genetic Algorithms!

Schema **T**heorem





Schema Theorem (1)

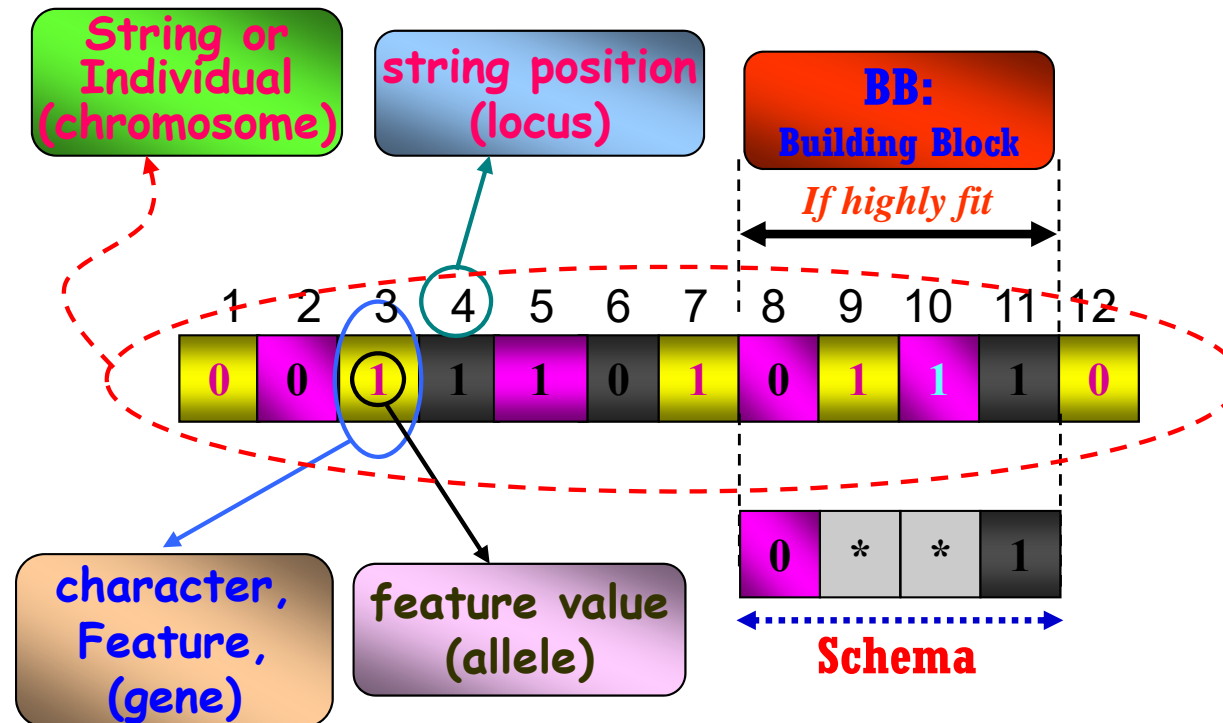


❖ Schema Theorem

- It guarantees the **success of GAs** by just applying genetic operations
 - ✓ **Concept:** the **good schemata** increase their **market share** in the population

Cf) What is **Schema**?

- ✓ A **subset** of the space in which **all** the strings **share** a **particular defined set**





Schema Theorem (2)



- **Order** of a schema H : $o(H)$

- Number of fixed positions present in the template

011*1**

$$o(011*0**) = 4$$

0*****

$$o(0******) = 1$$

- **Length** of a schema H : $\delta(H)$

- The distance between the first and last specific string position

011*1**

$$\delta(H_1) = 5 - 1 = 4$$

0*****

$$\delta(H_2) = 1 - 1 = 0$$

- How many **expected number** of schema will be inherited to the next generation?

- The **survival rate under simple reproduction**:

$$\frac{f(H)}{\bar{f}}$$

Proportional selection

where $f(H)$: average fitness of the strings representing schema H

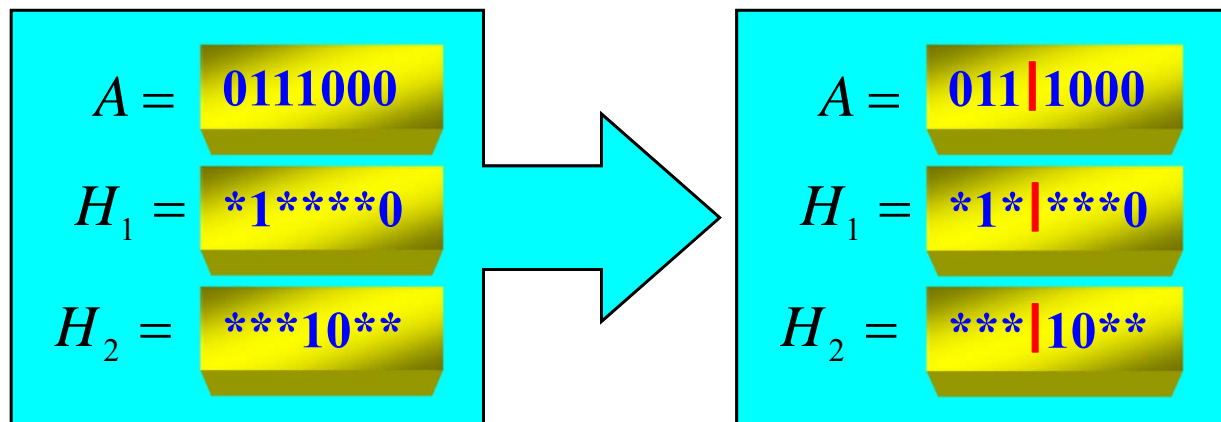
\bar{f} : average fitness of the entire population ($= \sum f_i / n$)



Schema Theorem (3)



- Consider the following situation on Crossover:



The schema H_1 will be **destroyed** by the **one-point crossover**!

- If the **crossover site** is **selected uniformly among** $l - 1 = 7 - 1 = 6$, then the **prob.** that schema H_1 is **destroyed** is given as

$$p_d = \delta(H_1) / (l - 1) = 5 / 6$$

: Just a conditional prob.

- The **survival prob. under simple crossover** is given by

$$p_s \geq 1 - p_c \cdot \frac{\delta(H)}{l - 1}$$

where p_c is a crossover prob.



Schema Theorem (4)



- **Mutation** is the *random alteration* of a **single position** with the prob. p_m
- Single allele **survives** with the prob. $(1 - p_m)$
- The **schema survival prob.** is obtained by

$$(1 - p_m)^{o(H)} \approx 1 - o(H) \cdot p_m$$

- Particular **schema H** receives **an expected number of copies** in the **next generation**:

$$\begin{aligned} m(H, t+1) &= m(H, t) \cdot P[H \text{ is survival under reproduction} \cap \text{crossover} \cap \text{mutation}] \\ &= m(H, t) \cdot P[H \text{ is survival under reproduction}] \cdot P[H \text{ is survival under crossover}] \\ &\quad \cdot P[H \text{ is survival under mutation}] \\ &\geq m(H, t) \cdot \frac{f(H)}{f} \cdot \left[1 - p_c \cdot \frac{\delta(H)}{l-1} \right] \cdot [1 - o(H) \cdot p_m] \geq m(H, t) \cdot \frac{f(H)}{f} \cdot \left[1 - p_c \cdot \frac{\delta(H)}{l-1} - o(H) \cdot p_m \right] \end{aligned}$$



Short, Low-order schemata receive **exponentially increasing trials** in subsequent generation!