Schema Theory

Sabah Sayed

Department of Computer Science
Faculty of Computers and Artificial Intelligence
Cairo University
Egypt

Why do Genetic Algorithms work?

In this section we take an in-depth look at the working of the standard genetic algorithm, explaining why GA constitutes an effective search procedure

For simplicity we discuss binary string representation of individuals

Schema Theory

- Schema theory seeks to give a theoretical justification for the efficacy of the field of genetic algorithms.
- What is a Schema:
 - a template
 - the new gene alphabet \Rightarrow {0,1,*} where * is a don't care

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* * 1 1 0 * * 1 1 * * *
```

- Schema is favorable traits in a solution, where a favorable schema is called an above average schema
- allows exploration of similarities among chromosomes
- represents all matching strings

Schema Theory

Example:

the schema (* 1 1 1 1 0 0 1 0 0) matches two strings: (0 1 1 1 1 0 0 1 0 0) and (1 1 1 1 1 0 0 1 0 0)

Q. which strings does this schema match?

$$(011*1011**)$$
 $\gamma^{2} = 8$

A schema matches 2^r strings

r: # of (*) in the schema

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011 0 1611 0 0
```

Schema Properties

Order of schema S: o(S)

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- number of fixed (non-*) positions

S1 = **11**101**

12 \quad 345

o(S1) = 5
```

 The order of a schema is useful to calculate survival probability of the schema for mutations

Defining length of schema S: d(S)

- distance between first and last **fixed** string positions
$$S1 = {**11**101**}$$
 $S2 = {**101***}$ $S2 = {**101***}$ above d(S1) = 9-3= 6 d(S2)=?

 The defining length of a schema is useful to calculate survival probability of the schema for crossovers

- Assume total number of chromosomes is PopSize.
- The number of above average individuals abiding by the above average schema S is P.
- f_i: fitness of chromosome <u>i</u> in the population of all chromosomes.
- Average fitness of population = $\frac{\sum_{i=1}^{PopSize} f_i}{PopSize} = \overline{F}$

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 The average fitness of above average individuals abiding with favorable schema S=

$$F_{av}(S) = \frac{\sum_{j=1}^{P} f_j}{P}$$

- where P is the number of individuals abiding by S
- Let:
 - m(S, t) denote the expected no. of individuals matched by schema S at time(t)
 - $-\mathbf{F}$ = total fitness of population, $F = \sum_{i=1}^{PopSize} f_i$
 - m(S, t+1) denote the expected no. of individuals matched by schema S at time (t+1)(next iteration).

•
$$m(S, t+1) = m(S, t) * PopSize * \frac{f_{av}(S)}{F}$$

- But F = total fitness where $\bar{F} = \frac{F}{PopSize}$, so F= \bar{F} * PopSize
- Substitute in (1): $m(S, t + 1) = m(S, t) * PopSize * \frac{f_{av}(S)}{\overline{F} * PopSize}$ $m(S, t + 1) = m(S, t) * \frac{f_{av}(S)}{\overline{F}}$

$$m(S,t+1) = m(S,t) * \frac{f_{av}(S)}{\overline{F}}$$
Note that $\frac{f_{av}(S)}{\overline{F}} > 1$

$$f_{av}(s) > \overline{F}$$

$$f_{av}(s) = \overline{F} + \in \overline{F}$$
Substitute in (2):
$$m(S,t+1) = m(S,t) * \frac{(1+\epsilon)\overline{F}}{\overline{F}}$$

$$m(S,t+1) = m(S,t) * \frac{(1+\epsilon)\overline{F}}{\overline{F}}$$

$$m(S,t) * (1+\epsilon) = m(S,t) * \frac{(1+\epsilon)\overline{F}}{\overline{F}}$$

$$m(S,t) * \frac{(1$$

Effect of Operators on Schema 2-Crossover

Remember: Effect of selection on schema:

$$m(S, t + 1) = m(S, t) * \frac{f_{av}(S)}{\overline{F}}$$

Effect of Crossover:

What is the probability of <u>destruction</u> of a schema by crossover?

Output

Description:

Output

Description

where *d(S)* is the defining length of the schema, and *I* is length of chromosome.

Effect of Operators on Schema 2-Crossover

What is the probability of <u>survival</u> of a schema after crossover?

-->It should be (1-probability of destruction), but... destruction only happens if crossover will occur, and crossover occurs with a probability of **P**_c. So:

$$P_s = 1 - P_c \left(\frac{d(S)}{l-1} \right)$$

Substitute in equation (1):

$$m(S, t + 1) = m(S, t) * \frac{f_{av}(S)}{\overline{F}} * (1 - P_c \frac{d(S)}{l-1})$$

Equation (2) represents the combined effects of selection and crossover.

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Effect of Operators on Schema 3-Mutation

/hat is the procent nutation?
i.e. all bits survive mutation without any constant \rightarrow probability of destruction of 1 bit: $P_d = P_m$ Thability of survival of 1 bit: $-1 - P_d = 1 - P_m$

$$P_s = 1 - P_d = 1 - P_m$$

→ probability of survival of schema: [given order of schema o(S)]

$$P_s(S) = (1 - P_m) * (1 - P_m) * \cdots [for o(S) times]$$

$$P_s(S) = (1 - P_m)^{o(S)}$$
 -- but Pm is a small number $P_s(S) \simeq 1 - o(S)P_m$

Effect of Operators on Schema 3-Mutation

So probability of **survival** of a schema after crossover and mutation is:

$$P_{s} = 1 - P_{c} \frac{d(S)}{l-1} - o(S)P_{m}$$

 \rightarrow Substitute in (2):

$$\underline{m(S, t + 1)} = \underline{m(S, t)} * \frac{f_{av}(S)}{\overline{F}} * (1 - P_c \frac{d(S)}{l-1} - o(S)P_m)$$

which is called the **Reproductive Schema Growth** Equation **RSG**

Conclusion:

'Short', 'low-order' & 'above-average' schema are <u>liable</u> to grow, increase and flourish in population

Exercise

Given the following schema 'S':



A- What is the probability of schema S survival after crossover knowing that Pc = 0.8? $P_S = I - P_s = \frac{d(s)}{I-I} = \frac{d(s)}{A}$