

Suggested reading:

D. E. Goldberg, *Genetic Algorithm in Search, Optimization, and Machine Learning*, Addison Wesley Publishing Company, January 1989

# **Schema Theorem**

- Schema theorem serves as the analysis tool for the GA process
- Explain why GAs work by showing the expectation of schema survival
- Applicable to a canonical GA
   binary representation
   fixed length individuals
   fitness proportional selection
   single point crossover
   gene-wise mutation

## Schema

- A schema is a set of binary strings that match the template for schema H
- A template is made up of 1s, 0s, and \*s where
   \* is the 'don't care' symbol that matches either
   0 or 1

## **Schema Examples**

The schema H = 10\*1\* represents the set of binary strings

#### 10010, 10011, 10110, 10111

The string '10' of length l = 2 belongs to  $2^{l} = 2^{2}$  different schemas

## Schema: Order o(H)

- The order of a schema is the number of its fixed bits, i.e. the number of bits that are not '\*' in the schema H
- Example: if H = 10\*1\* then o(H) = 3

## **Schema: Defining Length** $\delta(H)$

- The defining length is the distance between its first and the last fixed bits
- Example: if H = \*1\*01 then  $\delta(H) = 5 2 = 3$
- Example: if  $H = 0^{****}$  then  $\delta(H) = 1 1 = 0$

## Schema: Count

- Suppose x is an individual that belongs to the schema H, then we say that x is an instance of  $H(x \in H)$
- *m*(*H*, *k*) denotes the number of instances of *H* in the *k* th generation

#### **Schema: Fitness**

■ f(x) denotes fitness value of x

■ *f*(*H*,*k*) denotes average fitness of *H* in the *k*-th generation

$$f(H,k) = \frac{\sum_{x \in H} f(x)}{m(H,k)}$$

## Effect of GA On A Schema

- Effect of Selection
- Effect of Crossover
- Effect of Mutation
- $\blacksquare$  = Schema Theorem

#### **Effect of Selection on Schema**

Assumption: fitness proportional selection

Selection probability for the individual x

$$p_s(x) = \frac{f(x)}{\sum_{i=1}^N f(x_i)}$$

#### where the N is the total number of individuals

#### **Net Effect of Selection**

The expected number of instances of *H* in the mating pool *M*(*H*,*k*) is

$$M(H,k) = \frac{\sum_{x \in H} f(x)}{\overline{f}} = m(H,k) \frac{f(H,k)}{\overline{f}}$$

Schemas with fitness greater than the population average are likely to appear more in the next generation

## **Effect of Crossover on Schema**

Assumption: single-point crossover

Schema H survives crossover operation if
 one of the parents is an instance of the schema H AND

 $\Box$  one of the offspring is an instance of the schema H

#### **Crossover Survival Examples**

Consider H = \*10\*\*

P<sub>1</sub> = 1 1 0 1 0 ∈ H P<sub>2</sub> = 1 0 1 1 1 ∉ H  $S_1 = 1 1 0 1 1 ∈ H$  Schema H S<sub>2</sub> = 1 0 1 1 0 ∉ H survived

P<sub>1</sub> = 1 1 0 1 0 ∈ H P<sub>2</sub> = 1 0 1 1 1 ∉ H  $S_1 = 1 1 1 1 1 ∉ H$  Schema H S<sub>2</sub> = 1 0 0 1 0 ∉ H destroyed

## **Crossover Operation**

- Suppose a parent is an instance of a schema *H*. When the crossover is occurred within the bits of the defining length, it is destroyed unless the other parent repairs the destroyed portion
- Given a string with length l and a schema H with the defining length  $\delta(H)$ , the probability that the crossover occurs within the bits of the defining length is  $\delta(H)/(l-1)$

## **Crossover Probability Example**

Suppose H = \*1\*\*0
We gave
l = 5
δ(H) = 5 - 2 = 3

□Thus, the probability that the crossover occurs within the defining length is 3/4

## **Crossover Operation**

The upper bound of the probability that the schema *H* being destroyed is

$$D_c(H) \le p_c \frac{\delta(H)}{l-1}$$

where  $p_{\rm c}$  is the crossover probability

#### **Net Effect of Crossover**

The lower bound on the probability S<sub>c</sub>(H) that H survives is

$$S_{c}(H) = 1 - D_{c}(H) \ge 1 - p_{c} \frac{\delta(H)}{l - 1}$$

Schemas with low order are more likely to survive

#### **Mutation Operation**

Assumption: mutation is applied gene by gene

For a schema H to survive, all fixed bits must remain unchanged

Probability of a gene not being changed is

$$(1-p_m)$$

where  $p_m$  is the mutation probability of a gene

## **Net Effect of Mutation**

The probability a schema H survives under mutation

$$S_m(H) = (1 - p_m)^{o(H)}$$

Schemas with low order are more likely to survive

#### **Schema Theorem**

Exp. # of Schema H in Next Generation > Exp. # in Mating Pool ( $M(H,k) = m(H,k) \frac{f(H,k)}{\overline{f}}$ ) Prob. of Surviving Crossover ( $S_c(H) \ge 1 - p_c \frac{\delta(H)}{l-1}$ ) Prob. of Surviving Mutation ( $S_m(H) = (1 - p_m)^{o(H)}$ )

#### **Schema Theorem**

Mathematically

$$E[m(H,k+1)] \ge m(H,k) \frac{f(H,k)}{\bar{f}} \left(1 - p_c \frac{\delta(H)}{l-1}\right) (1 - p_m)^{o(H)}$$

The schema theorem states that the schema with *above average fitness*, *short defining length* and *lower order* is more likely to survive