School of Information Technology Indian Institute of Technology Kharagpur

IT 60108 :: Soft Computing Applications

Mid-Spring Semester Test Session 2013-2014

Full Marks: 40 Time: 02 hours

Answer ALL questions

(Numbers in the right side indicate the marks for questions)

- 1. Draw a flowchart of the steady state GA (SSGA). 3 (a) (b) Answer the following questions with reference to SSGA. How SSGA is different from simple GA (SGA)? 2 ii. To solve what type of optimization problem the SGA can be thought 1 iii. Comment on the performance of SGA with respect to the following. (You should justify your answers.) Convergence rate 1 2 Population diversity 2 Selection pressure
- 2. A binary search tree is a binary tree such that the key value of each node is greater than the key value of any node in its left sub-tree and less than the key value of any node in its right sub-tree. All key values in a binary search tree are assumed to be unique.

There is a need to construct an optimum binary search tree with the following specification.

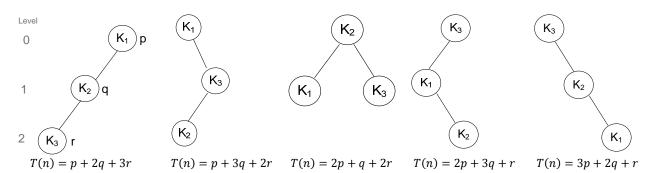
- Given a set $K = \{K_1, K_2, \dots, K_n\}$ with n distinct key values such that $K_i < K_{i+1}$, $i=1,2,\dots,n$. We have a probability p_i such that a search will be for key K_i where $\sum_{i=1}^n p_i = 1$
- Assuming that the root node is at level 0. The average search time T(n) for a given binary search tree is defined as

$$T(n) = \sum_{i=1}^{n} p_i \ (level (K_i) + 1)$$

where $level(K_i)$ denotes as level of K_i .

The problem is to find an optimum binary search tree with minimumT(n).

For an example, with tree key values K_1 , K_2 and K_3 (such that $K_1 < K_2 < K_3$) with probabilities p,q and r, (such that p+q+r=1), there are many binary search trees with different T(n) values few of which are shown below.



With reference to the above problem answer the following questions.

(a) Whether solving the above problem comes under hard computing, soft computing or hybrid computing?

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(b) It is proposed to solve the problem using genetic algorithm (GA). In that case state your approach to the following.

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Encoding scheme (clearly explain genotype and phenotype for an instance).

4

- ii. Crossover technique (explain your crossover technique with a small but clear example).
- 3. Selection strategy according to *Roulette Wheel* scheme is stated as follows.

Input: Given a population of size N with fitness values of individuals are $f_1, f_2, ..., f_N$ Output: Selection of a sub-population of size N_p for some N_p .

- 1. Calculate $p_i = \frac{f_i}{\sum_{i=1}^N f_i}$ for all $i=1,2,\ldots,N$
- 2. Calculate cumulative probability for each individual $P_i = \sum_{j=1}^i p_j$ for all $i=1,2,\ldots,N$
- 3. Generate a random number r (between 0 and 1, both inclusive)
- 4. Select the *j*-th individual such that $P_{j-1} < r \le P_j$.
- 5. Repeat Step 3-4 until N_p number of individuals are selected.

With reference to the above algorithm, answer the following questions.

(a) How many counts the i –th individual is expected to be selected?

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(b) Whether there is(are) case(s) that some individual(s) will be selected more than once. Argue your answer.

1

- (c) It is proposed to modify the above replacing Step 3—5 as follows.
 - 3. Choose i –th individual such that $P_{i-1} < r \le P_{ii-1}$ as a candidate
 - 4. Compute $e_i = p_i \times N$ with the p_i value of the candidate
 - 5. If $[e_i] \neq 0$, then select the i –th individual
 - 6. $r = e_i \lfloor e_i \rfloor$
 - 7. Repeat Step 3-6 until $N_p < N$ number of individuals are selected.

Assume that the procedure starts with an initial value of r=0.5. Here $\lfloor e_i \rfloor$ denotes the integer value not exceeding e_i .

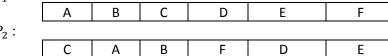
Calculate the selection of population, if we run the modified version of *Roulette Wheel* for 8 times, with the following population with their fitness values.

Individual	1	2	3	4	5	6	7	8
Fitness value	1.0	2.1	3.1	4.0	4.6	1.9	1.9	4.5

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4.	For the following mating pairs and their crossover techniques, obtain the offspring
	chromosomes.

(a) P_1 :



with precedence Preservative Crossover (PPX) with a vector

1	2	1	1	2	2

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(b) P_1 :

1	1	0	1	0	0	0	1

 P_2 :

0	1	1	0	1	0	1	1

 P_3 :

0	1	1	0	1	1	0	0

with three Three Parent Crossover.

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(c) For the following offspring in Real-coded GA, obtain three mutated offspring chromosomes using the *Random mutation* scheme.

$$P_1 = 10 \text{ and } P_2 = 30$$

Assume any other value(s) which might involve in the calculation.

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