Department of Engineering/Informatics, King's College London Nature-Inspired Learning Algorithms (7CCSMBIM) Assignment 1

This coursework is assessed. A type-written report needs to be submitted online through KEATS by the deadline specified on the module's KEATS webpage.

Q1 Binary Genetic Algorithm by Hand

- a) Write down the first 7 digits of your student ID as $s_1s_2s_3s_4s_5s_6s_7$. Form an 8-digit number as $s_2s_3s_4s_5s_6s_7b_1b_2$ where the two digits b_1b_2 denote your day of birth. (5 Marks)
- b) Write down the cost function with two variables: $f(x,y) = -(s_2 + s_3)x^2 + (s_4 + s_5)xy (s_6 + s_7)y^2 (b_1 + b_2)$. In the following, we are going to find the optimal solution for the optimisation problem: $\min_{x,y} f(x,y)$ using binary genetic algorithm. (5 Marks)
- c) Population Initialisation Greate a population with 8 chromosomes and list then in a table with the healings of n, Chromosome, Decoded x, y, and 'Cost' where n denotes the chromosome number. It is assumed that both variables x and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and y are in the range of $-(s_1 + s_2)$ and $s + s_6 + s_7$ where s denotes the larges Sigit and s are in the range of s and s are in the second distance of s and s are in the range of s and
- d) Natural Selection: Show the ranked population after the process of Natural Selection with $N_{keep} = 4$ in a table with the heading of 'n', 'Chromosome', 'Decoded x, y' and 'Cost'. (10 Marks)
- e) **Selection:** Using the cost weighting technique, show the probability table with the heading of 'n', 'Chromosome', 'Decoded x, y', 'Cost', ' $C_n = c_n c_{N_{keep+1}}$ ', ' P_n ' and ' $\sum_{i=1}^n P_n$ '. (10 Marks)
- f) Crossover: Show the population after the process of Single-Point Crossover. The chromosomes are selected for crossover according to the probability table in the Selection process. The first pair of parents are chosen according to the random numbers $\{\frac{s_2}{s_2+s_3+s_4+s_5+s_6}, 1-\frac{s_2}{s_2+s_3+s_4+s_5+s_6}\}$. The next pair of parents are the rest two chromosomes in the pool of N_{keep} . If the remaining number of chromosomes is more than two. Make a remark and randomly pick two as the second pair of parents. The crossover point for the first pair of parents is right after the $round(\frac{s_2+s_3}{2})^{th}$ bit and the crossover point for the second pair of parents is right after the $round(\frac{s_4+s_5}{2})^{th}$ bit, where the operator $round(\cdot)$

rounds off the value to the nearest integer. Different colours should be used to show clearly where each part of the chromosome goes to in the crossover process. *Note: The left most bit of the chromosome is the first bit.* (10 Marks)

- g) **Mutation:** Show the population in a table with the heading of 'n', 'Chromosome', 'Chromosome after mutation', 'Decoded x, y' and 'Cost'. The mutation rate is $\mu = 0.25$. Calculate the number of bits to be mutated (#mutation) and randomly choose #mutation bits of your choice. Highlight the mutated bits.

 (10 Marks)
- Q2 Denote R_1 as the remainder of $\frac{s_2+s_3+s_4}{9}$. Write a Matlab script to find the minimum of the function R_1+1 in Appendix using binary genetic algorithm. All variables are in the range of -20 and 20. The precision of the solution should be up to 4 decimal places.
 - a) Show your calculation getting R_1 . (5 Marks)
 - b) Determine the number of bits for each variable for binary decoding. Explain Assignment Project Exam Help (5 Marks)
 - c) Run the procedure of binary genetic algorithm for 10 times with the combinations of population size $(p \text{ or } N_{rep})$ 10, 50 100 and μ : 0.1, 0.5, 0.9. 9 combinations of the total. Postant the best the worst costs of the 10 runs for each combination; the mean and the standard deviation of the best the worst costs for the 10 runs for each combination. Specify other control parameters of binary seneral abovirum usellar this quarter of iterations, population size, N_{keep} , methods of crossover and mutation, etc. List the results in Table 1 and Table 2. (20 Marks)
 - d) Comment briefly on the results obtained (with support/evidence) in terms of reliability and sensitivity of the binary genetic algorithm. (10 Marks)

"Runs" Number	$p = 10$ $\mu = 0.1$		$p = 10$ $\mu = 0.9$		$p = 50$ $\mu = 0.5$			$p = 100$ $\mu = 0.5$	$p = 100$ $\mu = 0.9$
1									
2									
3									
4									
5									
6									
7									
8									
9									
10									
Mean									
Standard Deviation									
Best Cost									
Worst Cost	\	~40.40	2424	Dag 2	2 24]			-1	
F	Assignment Project Exam Help Table 1: Statistics of 10 runs.								

https://powcoder.com							
Mean of "Mean"							
Standard deviation of "mean"							
Mean of "Standard Deviation" Standard deviation of "Standard Deviation"	owcoder						
Standard deviation of "Standard Deviation"	JW COUCI						
Mean of "Best Cost"							
Standard deviation of "Best Cost"							
Mean of "Worst Cost"							
Standard deviation of "Worst Cost"							

Table 2: Overall Statistic.

Appendix 1: Test Functions

$$N = 2 + s_2 + s_3$$
F1:
$$\sum_{i=1}^{N} (|x_i| + \cos(x_i))$$
F2:
$$\sum_{i=1}^{N} (|x_i| + \sin(x_i))$$
F3:
$$\sum_{i=1}^{N} x_i^2$$
F4:
$$\sum_{i=1}^{N-1} (100(x_{i+1} - x_i^2)^2 + (1 + x_i)^2)$$

F5:
$$10N + \sum_{i=1}^{N} (x_i^2 - 10\cos(2\pi x_i))$$

F6: $1 + \sum_{i=1}^{N} \frac{x_i^2}{4000} - \prod_{i=1}^{N} \cos(x_i)$
F7: $0.5 + \frac{\sin^2(\sqrt{x_1^2 + x_2^2}) - 0.5}{1 + 0.1(x_1^2 + x_2^2)}$
F8: $(x_1^2 + x_2^2)^{0.25} \sin(30((x_1 + 0.5)^2 + x_2^2)^{0.1}) + |x_1| + |x_2|$
F9: $-x_1 \sin(\sqrt{|x_1 - (x_2 + 9)|}) - (x_2 + 9) \sin(\sqrt{|0.5x_1 + x_2 + 9|})$

In the above test functions, all x_i , $i = 1, 2, \dots, N$, are decision variables.

Marking: The learning outcomes of this assignment are that students will understand the fundamental principle of binary genetic algorithm (BGA) and appreciate the BGA performance through the evaluation of some benchmark functions. The assessment will look into the knowledge and understanding on the topics. When answering the questions, show/explain/describe clearly the steps with reference to the equations/theory/algorithms (stated in the lecture slides). When making comments, provide statements with support from the results obtained.

from the results obtained Project Exam Help

Purposes of Assignment: This assignment is designed to reinforce the understanding of Genetic Algorithm by going through every single step and to get an idea how it performs when applying the Country of the country color of Genetic Algorithm and thus know what to note when applying it.

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