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Section ? BCS-6B

Assignment 3

Question 1:

eg (A,B,C,D,E,F)

· Chromosome: A sequence of Genes representing order of cities visited.

Initial population A, C, B, D, E, F] . . . [A,B,D,C,F,E][A, C, F, D, C, E]

If Fitness = if valid sequencel): return total Distance

elsė;

yeturn & o

A valid sequence which teasible and visits all cities once.

1) [A,C,B,D,E,F] = 4+3+5+1+4=17 = 0.0588 @ [A,B,D,C,F,E] = 2+5+3+11+4= 1/25=0.04

[A,C,F,D,C,E] = 3+11+6+3+6=1/29=0.034





1,0) selected

This is not the best way to cross over since it results in invalid sequences and howe zero fitness. The sequence have cities repeated and miss at least one of the cities.

This isnot the best way to perform mutation. It can cause both repeated cities, missing cities as well as invalid sequences.

Question 2
$$f(x) = x^3 - 60x^2 + 900x + 100$$

Chromosome	Binary String	×	Fitness
P2	001111	15	3475
Pa	000100	4	2804





b)	00)111	
	000/100	
	001000 -> 001000	
	000011	
i i		•
	Chromosome Binary String oc	Fitness
12.	01 001000 8	3972
	02 000011 3	2287
c)	PI+ P2 = 3475+2804 = 6549	
	$D_1 + D_2 = 3972 + 2287 = 6250$	7
	No the overall fitness has n	reduced.
d)	001000	**





Quest	ion 3 1	
E	x yz lo	de la
Pr	0000	$\vec{N} = \begin{bmatrix} -2 & 2 & 32 \end{bmatrix}$
P_	10000	$\alpha = 1$
P3	0010	
P4	0101	ì
B	1011	
P6	1101	
P7	01111	
Pg	1 1 1 1	
iteration	1	2.232]
P,(0,0,0)	d=0	
(0,70)	y = 0	nochange
Pa (1,0,0		
1	7 = 21	mochange [-2 2 32] +[-1-1 00]
P3(0,0,		-[3, 3,2]
		no change
	y = 0	
P4(010) 0=1	
1000)	no change
0-6.01) d=1	
P5 (101	for any and the same	no change
0 (110)	J=1	
P6 (110)	_	nochanje
1	9=1	
P7(011)	d=1	no change.
	7=1	





iteration 2 [-3132]

$$P_{1}(0,0,0)$$
 $d=0$ $y=0$ no change $P_{2}(1,0,0)$ $d=0$ $y=0$ no change $P_{3}(0,0,1)$ $d=0$ $y=0$ no change $P_{4}(0,1,0)$ $d=1$ $y=1$ no change $P_{5}(1,0,1)$ $d=1$ $y=1$ no change $P_{6}(110)$ $d=1$ $y=1$ no change $P_{7}(011)$ $d=1$ $y=1$ no change $P_{8}(111)$ $d=1$ $y=1$ no change

No change in weights