

PART-2

PROBLEM-3

Given initial data table D is:

Goal is to transform the data table

D to TD representing the population

with the fitness function $F = 0$

for all chromosomes.

income	student	rating
high	no	fair
high	yes	fail
low	yes	fail
medium	yes	excellent
low	no	fail

PART-1

We are asked to write the set of Binary Encoding chromosomes representing the data table D.

We can call this initial population as P.

The possible values for each attribute are

Income = high, low, medium.

student = yes, no.

rating = fair, excellent.

So each chromosome would be of the format

$C_1 C_2 C_3 C_4 C_5 C_6 C_7$ as there are total of 7 unique values.

And we are also asked to calculate the fitness value of each chromosome using Definition 2.

Chromosome	$In=h$	$In=l$	$In=m$	$St=Y$	$St=N$	R_{refair}	R_{GxL}	F
Ch 1	1	0	0	0	1	1	0	3
Ch 2	1	0	0	1	0	1	0	3
Ch 3	0	1	0	1	0	1	0	3
Ch 4	0	0	1	1	0	0	1	3
Ch 5	0	1	0	0	1	1	0	3

Note: The number of chromosomes don't change from the initial data.

The fitness function is also evaluated for all the chromosomes & we got 3 for all of them. Importantly fitness value is same for all the chromosomes.

We have also kept all these values in a single table.

PART-2

- (1) We need to create ONE generations P_1 from our initial population P using our GA operators.
We are also asked to use Single Point Crossover & Single Point Mutation.

SELECTION:

If we see, the fitness values are same for all chromosomes, so we can select randomly any two chromosomes from P .

Note: We keep the population size to be constant.
so after the entire process 5 chromosomes should be present.

REPRODUCTION:

We will follow single point crossover as mentioned in Definition 1 which is after C_p . Since the fitness values are same, we can randomly pick 2 for crossover.

We'll pick ch1 & ch2 i.e. 1000110, 1001010 & perform the cross over after C4.

ch1	1000 110
ch2	1001 010
Off1	1000 010
Off2	1001 110

Similarly next we'll pick ch3 & ch4

ch3	0101 010
ch4	0011 001
Off3	0101001
Off4	0011010

RECOMBINATION:

Since the population size needs to be constant, we'll pick ch5 as our off5. (fitness values are same)

After this step the chromosomes obtained are

1000010, 1001110, 0101001, 0011010 & 0100110

MUTATION:

Now single point mutation has to be applied on each of the obtained off5 sparrings.

Since it's Single point Mutation, we have randomly selected 1 bit from each offspring chromosome to obtain the final set of offspring i.e. Since mutation pro. is not mentioned.

1000010, 100110, 0101001, 0011010 & 0100110

Inverted C₁ C₅ C₂ C₄ C₅

0000010, 1001010, 0001001, 0010010 & 0100010

TERMINATE :

Since we are asked for only one generation, the DNA generation population P, looks like this using the process we followed.

Chromosome	I _{n=h}	I _{n=l}	I _{n=m}	C _{t=Y}	C _{t=N}	R _{refar}	R _{exec}	F
Ch1	0	0	0	0	0	1	0	1
Ch2	1	0	0	1	0	1	0	3
Ch3	0	0	0	1	0	0	1	2
Ch4	0	0	1	0	0	1	0	2
Ch5	0	1	0	0	0	1	0	2

(2) The best chromosome after this one generation is $\text{chi } 0000010$ as its fitness value is 1
 \Rightarrow
 which is closer to $F=0$ (our goal) the optimum ideal solution.

(3) Now we need to write Data Table D_1 using the population P_1 .

$D_1 =$

Income	Student	Rating
		Fair
high	yes	Fair
	yes	Excellent
medium		Fair
low		Fair

If we observe there are few missing values in the data table D_1 .

The reasons behind them are explained below.

Ch1: This chromosome doesn't have values for
Income & student attributes.

Ch3: No value for Income attribute

Ch4 & Ch5: No defined values for student attribute

So finally datatable D_i is generated of the same
format as of D . (Initial datatable).