CSE422 ASSIGNMENT !

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SECTION: 06



Question 1

(1)

$$A B C D E F G H$$
 $CI \rightarrow ABH$ 
 $CI \rightarrow ABCDH$ 
 $CS \rightarrow EFH$ 
 $CS \rightarrow DEGH$ 
 $CI \rightarrow DEGH$ 

(2) We = 
$$\sum_{i=1}^{8} w(0i) * B(0i)$$
  
Re =  $\sum_{i=1}^{8} Re(0i) * B(0i)$ 

fitness, f(n) = of Pe, we < man - weight

1110

1110 > man weight

ч	we 5	Me > wan-najant	we & man-welant Town False	4(n) 37 -4
ርጉ	16	Tow	faloe	-1
C3	13	Tow	faise	-10
CA	$\mathcal{P}$	Trow		

CA has the least fitness function, so is excluded. and as have the best fitness function

(3) 
$$C1 \times C3 \rightarrow A \downarrow B \qquad H \rightarrow AFH$$

$$OY \longrightarrow FQH$$

Randomly, a gene is selected from both offsprings and that gene is charged due to mutation.

$$02 \rightarrow EBH \longrightarrow EAH$$

$$2nd agne is$$
mutated

$$mut - OI \longrightarrow ADH$$

$$mut - O 2 \longrightarrow EAH$$

On checking the fitness function of both the mutated offspoing, mut-OI have the higgest fitness function. During coossover, oI have necessed the best genes forom parents and during mutation, the change in gene caused an advantage to be the best in this scenario.

Question 2

A\* Gnaph seanch

Visited: — A\* | E6 | H6 | C9 | F8 | G9 | X8

X8 Bio Dis Go Bio Dis Xx X8 Co Bio Co Lo Dis Xo Co Hr. Bio Co Lo Dis Xo Er Bio Co Lo Ho Er Bio Co Lo Ho

 $path: A \longrightarrow C \longrightarrow G \longrightarrow X$ 

S: tea

Coneady Best Finst Search Visited: - AT | EA | H2 | X0

4

Hy B2 GP Ed

Ey B. G. ty Hr

Hy Be Co to Do Xo

Xo By Cp Ly Da

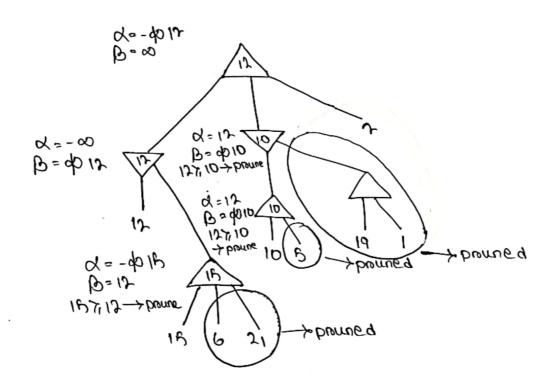
bath: A -> E -> H -> X

6: teas

Question 3

 $d = -\infty$  [update with man value at  $\triangle$ ]  $B = \infty$  [update with min value at  $\nabla$ ]  $d = \infty$  [update with min value at  $\nabla$ ]

ary wet



Question 4

(1)

(2) bin -to-dec = int(0);  $0 \rightarrow chnomosome$ N = bin - to-dec

$$fitness = \begin{cases} x, f(x) = 0 \end{cases}$$

CI has the lowest fitness function, so is excluded.

$$01 \longrightarrow 0010$$

(3) 
$$01 \rightarrow 0011 \xrightarrow{\text{Ath agne is}}$$
 $01 \rightarrow 0010 \xrightarrow{\text{Ath agne is}}$ 
 $01 \rightarrow 0010 \xrightarrow{\text{3rd agne is}}$ 

mutated

Randomly, a gene is selected from both offsprings and that gene is changed due to mutation.

$$mut - 01 \longrightarrow 0000$$

$$mut - 02 \longrightarrow 0000$$

$$mut - 01 \qquad 0 \qquad 1$$

$$mut - 02 \qquad 6 \qquad 0$$

On checking the fitness function of both the mutated offsproing, mut\_01 have the highest function.

During crossover, mut\_01 have neceived the best genes from parents and during mutation, the change in gene caused as an advantage to be the best in this function.

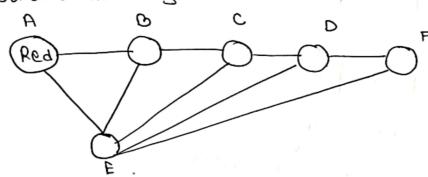
## Question 5.

Ponward checking can reduce the domain of unassigned variable using assigned variable.

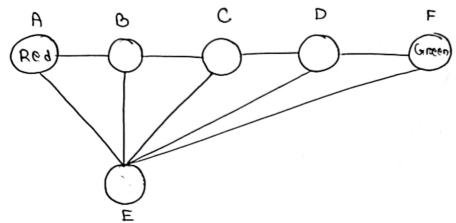
we can bostically product the outcome of the next unassigned variable by using a known variable. beforehard.

Node consistency checking is the forward checker where a random node is chosen and color is assigned to it and its following nodes are reduced with respect to the color assigned, and constraint,

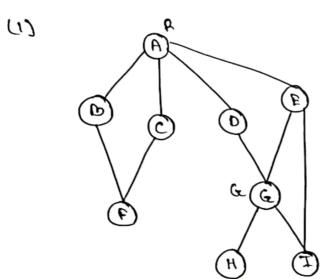
Constraint: Two adjacent nodes cannot be of same colors



Anc consistency Node is stronger than Node consistency checking where domain is checked and remaining value can be used to reduce the domain.



ango stops and anothing is tent



Deance neunistic is the most constraining variable. It is the vaniable which can neduce the highest vaniables. can neduce 4 vaniables assigned node (1) (1) (0) (1) (2) (4) (0)

(1) (1) (0) (1) (2) (4) (0) Red GD GD D E Red Green RD RE

(A) (O) (O) (O) (I) (I) (2) node (A) (O) (O)

(A) (O) (O) (O) (I) (I) (2) node (A) (O) (O)

Red GD GD D E Red Green RD RB

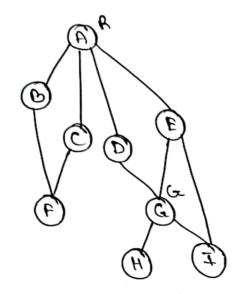
Red GD GD D Blue Red Green RD RB

Red GD GD D Blue Red Green RD RB

Red GD GD GD D Blue Red Green RD RB

Red GD GD GD D Blue Red Green RD RB no de

(2)



The colon of node "B" can either be oneen on blue.

It node "B" is assigned with either some on blue, then B can neduce only node "F". So, either blue on some can be assigned to node B.

Hene, node b is assigned with