

# CSCI3220 2018-19 First Term Assignment 4

I declare that the assignment here submitted is original except for source material explicitly acknowledged, and that the same or closely related material has not been previously submitted for another course. I also acknowledge that I am aware of University policy and regulations on honesty in academic work, and of the disciplinary guidelines and procedures applicable to breaches of such policy and regulations, as contained in the following websites.

University Guideline on Academic Honesty: <http://www.cuhk.edu.hk/policy/academichonesty/>

Student Name: ZHANG Chongzhi  
Student ID : 1155077072

1. (a). Not possible.

We have  $d(s_2, s_3) + d(s_2, s_4) = 2 + 6 = 8 < d(s_3, s_4) = 10$ .

Thus the distance Matrix doesn't meet the properties (iv), the rule of Triangle.

Therefore it can not construct an additive tree.

(b).

	$s_1$	$s_2$	$s_3$	$s_4$	$s_5$
$s_1$	0	8	6	8	2
$s_2$	8	0	2	6	8
$s_3$	6	2	0	10	6
$s_4$	8	6	10	0	6
$s_5$	2	8	6	6	0

Merge  
 $\{s_3\}, \{s_5\} \Rightarrow$

	$s_1, s_5$	$s_2$	$s_3$	$s_4$
$s_1, s_5$	0	8	6	7
$s_2$	8	0	2	6
$s_3$	6	2	0	10
$s_4$	7	6	10	0

Merge  
 $\{s_3\}, \{s_5\} \Rightarrow$

	$s_1, s_3$	$s_2, s_4$	$s_5$
$s_1, s_3$	0	7	8
$s_2, s_4$	7	0	7
$s_5$	8	7	0

$s_1 \quad s_2 \quad s_3 \quad s_4 \quad s_5$

$s_1 \quad s_3 \quad s_4 \quad s_5 \quad s_1$

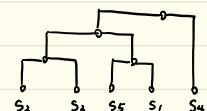
$s_2 \quad s_3 \quad s_4 \quad s_5 \quad s_1$

Merge  
 $\{s_1, s_5\}, \{s_2, s_3\} \Rightarrow$

	$s_1, s_5, s_5$	$s_4$
$s_1, s_5, s_5$	0	7.5
$s_4$	7.5	0

Merge  
 $\{s_1, s_2, s_3, s_5\}, \{s_4\} \Rightarrow$

	$s_1, s_2, s_3, s_4, s_5$
	0



Distance Matrix (DM)

(C).

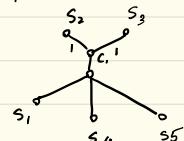
<b>d</b>	<b>S<sub>1</sub></b>	<b>S<sub>2</sub></b>	<b>S<sub>3</sub></b>	<b>S<sub>4</sub></b>	<b>S<sub>5</sub></b>
<b>S<sub>1</sub></b>	0	8	6	8	2
<b>S<sub>2</sub></b>	8	0	2	6	8
<b>S<sub>3</sub></b>	6	2	0	10	6
<b>S<sub>4</sub></b>	8	6	10	0	6
<b>S<sub>5</sub></b>	2	8	6	6	0

<b>U</b>	
<b>S<sub>1</sub></b>	24
<b>S<sub>2</sub></b>	24
<b>S<sub>3</sub></b>	24
<b>S<sub>4</sub></b>	30
<b>S<sub>5</sub></b>	22

Q matrix (Q.M.)

<b>Q</b>	<b>S<sub>1</sub></b>	<b>S<sub>2</sub></b>	<b>S<sub>3</sub></b>	<b>S<sub>4</sub></b>	<b>S<sub>5</sub></b>
<b>S<sub>1</sub></b>	0	-24	-30	-30	-40
<b>S<sub>2</sub></b>	-24	0	-42	-36	-22
<b>S<sub>3</sub></b>	-30	-42	0	-24	-28
<b>S<sub>4</sub></b>	-30	-36	-24	0	-34
<b>S<sub>5</sub></b>	-40	-22	-28	-34	0

Merge  
 $\{S_3\}, \{S_3\} \Rightarrow$



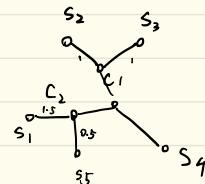
$$d(S_2, C_1) = \frac{1}{2} \times 2 + \frac{0}{6} = 1, \quad d(S_3, C_1) = 2 - 1 = 1.$$

<b>d</b>	<b>S<sub>1</sub>, S<sub>3</sub></b>	<b>S<sub>1</sub></b>	<b>S<sub>4</sub></b>	<b>S<sub>5</sub></b>
<b>S<sub>1</sub>, S<sub>3</sub></b>	0	6	7	6
<b>S<sub>1</sub></b>	6	0	8	2
<b>S<sub>4</sub></b>	7	4	0	6
<b>S<sub>5</sub></b>	6	2	6	0

<b>U</b>	
<b>S<sub>1</sub>, S<sub>3</sub></b>	19
<b>S<sub>1</sub></b>	16
<b>S<sub>4</sub></b>	21
<b>S<sub>5</sub></b>	14

<b>Q</b>	<b>S<sub>1</sub>, S<sub>3</sub></b>	<b>S<sub>1</sub></b>	<b>S<sub>4</sub></b>	<b>S<sub>5</sub></b>
<b>S<sub>1</sub>, S<sub>3</sub></b>	0	-23	-20	-21
<b>S<sub>1</sub></b>	-23	0	-21	-26
<b>S<sub>4</sub></b>	-26	-21	0	-23
<b>S<sub>5</sub></b>	-21	-26	-23	0

Merge  
 $\{S_1, S_3\}, \{S_1, S_3\} \Rightarrow$



$$d(S_1, C_2) = \frac{1}{2} \times 2 + \frac{16-14}{4} = 1 + \frac{1}{2} = 1.5$$

$$d(S_5, C_2) = 2 - 1.5 = 0.5$$

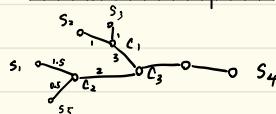
<b>d</b>	<b>S<sub>1</sub>, S<sub>5</sub></b>	<b>S<sub>2</sub>, S<sub>3</sub></b>	<b>S<sub>4</sub></b>
<b>S<sub>1</sub>, S<sub>5</sub></b>	0	5	6
<b>S<sub>2</sub>, S<sub>3</sub></b>	5	0	7
<b>S<sub>4</sub></b>	6	7	0

<b>U</b>	
<b>S<sub>1</sub>, S<sub>5</sub></b>	11
<b>S<sub>2</sub>, S<sub>3</sub></b>	12
<b>S<sub>4</sub></b>	13

<b>Q</b>	<b>S<sub>1</sub>, S<sub>5</sub></b>	<b>S<sub>2</sub>, S<sub>3</sub></b>	<b>S<sub>4</sub></b>
<b>S<sub>1</sub>, S<sub>5</sub></b>	0	-18	-18
<b>S<sub>2</sub>, S<sub>3</sub></b>	-18	0	-18
<b>S<sub>4</sub></b>	-18	-18	0

$$d(S_1, C_3) = \frac{1}{2} \times 5 + \frac{11-12}{2} = 2$$

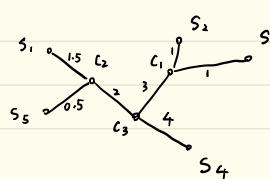
$$d(S_2, C_3) = 5 - 2 = 3.$$



<b>d</b>	<b>S<sub>1</sub>, S<sub>5</sub></b>	<b>S<sub>4</sub></b>
<b>S<sub>1</sub>, S<sub>5</sub></b>	0	4
<b>S<sub>4</sub></b>	4	0

<b>U</b>	
<b>S<sub>1</sub>, S<sub>5</sub></b>	4
<b>S<sub>4</sub></b>	4

Merge the  
 $\{S_1, S_5\}$  and  $\{S_4\}$ . We get the final graph

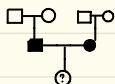


3. (a). Yes.

Assumed the disease is **recessive** which determined by a single locus, and both of the parents have a genotype **Aa**, then it is possible for their children to both have & have not disease.

(b). Yes, it is possible because we may determine the genotype of parents by the grandparents.

For instance, gives a pedigree chart as follow:



From the status of grandparents, we can know the disease is recessive and both the parents of child has a genotype aa.

Therefore, their children is bound to have a disease.

(c). In some situations, yes.

Situations	Location	Trait Dominance	Daughter genotype	Infer	Father genotype	Father's Phenotype
1.	X-linked	Recessive	X <sup>A</sup> X <sup>A</sup>	→	X <sup>A</sup> Y	Unaffected
2.	X-linked	Recessive	X <sup>a</sup> X <sup>a</sup>	→	X <sup>a</sup> Y	Affected
3.	X-linked	Dominant	X <sup>A</sup> X <sup>A</sup>	→	X <sup>A</sup> Y	Affected
4.	X-linked	Dominant	X <sup>a</sup> X <sup>a</sup>	→	X <sup>a</sup> Y	Unaffected

(d). Assumed that

P: Both parents are heterozygous.

F<sub>1</sub>: First child is heterozygous.

F<sub>2</sub>: Second child is heterozygous.

$$\Pr(F_1 \wedge F_2 | P) = \frac{1}{2} \times \frac{1}{2} = \frac{1}{4}.$$

$$\Pr(F_1 \vee F_2 | P) = 1 - \Pr(\neg F_1 \wedge \neg F_2 | P) = 1 - \frac{1}{4} = \frac{3}{4}.$$

$$\Pr((F_1 \wedge F_2) \wedge (F_1 \vee F_2) | P) = \Pr(F_1 \wedge F_2 | P) = \frac{1}{4}.$$

$$\Pr(F_1 \wedge F_2 | (F_1 \vee F_2) \wedge P) = \frac{\Pr((F_1 \wedge F_2) \wedge (F_1 \vee F_2) | P)}{\Pr(F_1 \vee F_2 | P)} = \frac{\frac{1}{4}}{\frac{3}{4}} = \frac{1}{3}.$$

(e). Explanation 1:

The situation just happened by chance. In fact, all the descendants have same probability to have a disease, but all the females were lucky while all the males were unlucky.

Explanation 2:

The disease is X-Linked, and it is recessive. The parents have a genotype  $X^A Y$  and  $X^A X^a$ . Thus all the female descendants have a genotype  $X^A X^a$ , which is not affected. But all the male descendants have a genotype  $X^a Y$ , which is affected.

Explanation 3:

All the descendants actually have the disease genotype  $aa$ , but there is another gene (Marked B.b) which is X-linked and dominant, which have a effect that can cover the disease phenomenon.

Cous father  $aa X^B Y$ , mother  $aa X^b X^b$ .

All the females have the genotype  $aa X^B X^b$ , which is not affected.

All the males have the genotype  $aa X^b Y$ , which is affected.