

IT

bioinformatics Exam
2014-2015

Question 4

(a)

Lis o ~

$$(b) F = (x + \bar{z})(\bar{y} + z)(y + \bar{x})$$

000	001
010	100
110	101
011	111

$x=1$

$+_{=0}$

100
110
101
111

N_1

000	001
010	011

N_1'

N_2

000
010

$+ (N_1, \bar{z})$

$$N_3 = \text{merge}(N_1, N_2) =$$

100	110
101	111
000	010

N_4

100	101
000	.

$+ (N_3, \bar{y})$

110	111
010	.

$y=1$

N_4'

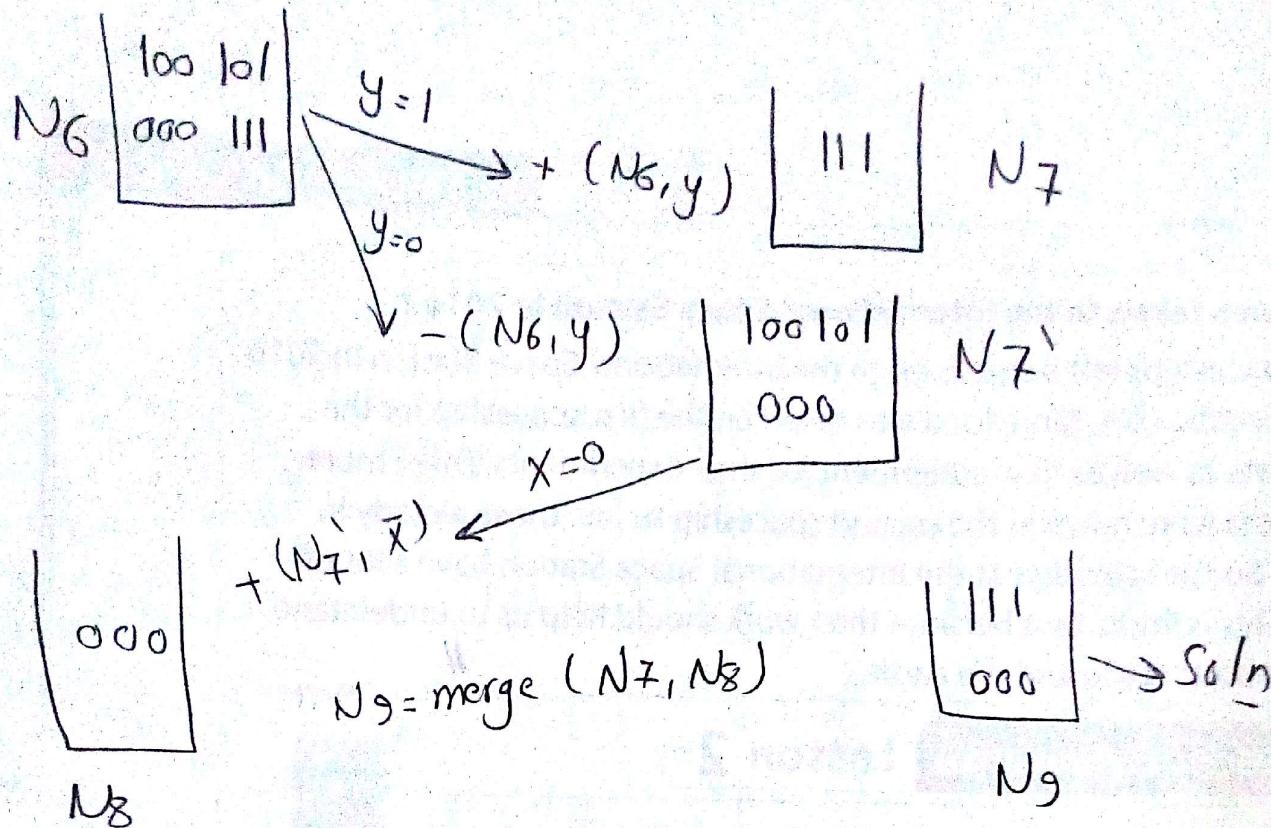
N_5

111
.

$$N_6 = \text{merge}(N_4, N_5) =$$

100	101
000	111

2



Lipton's Algorithm in TTP_L

- | | |
|------------------------------------|-------------------------------------|
| (1) input (N_0) | (10) $N_7 = +(N_6, y)$ |
| (2) $N_1 = +(N_0, x)$ | (11) $N_7' = -(N_6, y)$ |
| (3) $N_1' = -(N_0, x)$ | (12) $N_8 = +(N_7', x)$ |
| (4) $N_2 = +(N_1, z)$ | (13) $N_9 = \text{merge}(N_7, N_8)$ |
| (5) $N_3 = \text{merge}(N_1, N_2)$ | (14) detect (N_9) |
| (6) $N_4 = +(N_3, \bar{y})$ | |
| (7) $N_4' = -(N_3, \bar{y})$ | # |
| (8) $N_5 = +(N_4, z)$ | |
| (9) $N_6 = \text{merge}(N_4, N_5)$ | |

3

Question 2

1 → 2
5

$$(a) \pi = (1 \ 6 \ 2 \ 3 \ 4 \ 5)$$

$$\begin{array}{l} i=1 \\ \pi_j=i \\ j=1 \end{array} \quad \begin{array}{l} i=2 \\ \pi_j=i \\ j=3 \end{array} \quad \pi = (1 \ 2 \ 6 \ 3 \ 4 \ 5)$$

$$\boxed{\rho(2,3)} \quad \pi = (1 \ 2 \ 3 \ 6 \ 4 \ 5)$$

$$\begin{array}{l} i=3 \\ \pi_j=i \\ j=4 \end{array} \quad \begin{array}{l} i=4 \\ \pi_j=i \\ j=5 \end{array} \quad \pi = (1 \ 2 \ 3 \ 4 \ 6 \ 5)$$

$$\boxed{\rho(3,4)} \quad \boxed{\rho(4,5)} \quad \pi = (1 \ 2 \ 3 \ 4 \ 5 \ 6)$$

number of reversal = 4

$$\begin{array}{l} i=5 \\ \pi_j=i \\ j=6 \end{array} \quad \boxed{\rho(5,6)}$$

$$\pi = (1 \ 6 \ 2 \ 3 \ 4 \ 5)$$

$$\rho(2,6)$$

$$\pi = (1 \ 5 \ 4 \ 3 \ 2 \ 6)$$

$$\rho(2,5)$$

$$\pi = (1 \ 2 \ 3 \ 4 \ 5 \ 6)$$

min number of Reversal = 2

#

[4]

(b)

$$L = \{2, 2, 3, 3, 4, 5, 6, 7, 8, 10\}$$

$$X = \{0\} \quad \text{max element in } L = 10$$

$$X = \{0, 10\} \quad L = \{2, 2, 3, 3, 4, 5, 6, 7, 8\}$$

Remove 8

$$y = 2$$

$$\Delta(y, X) = \{2, 8\}$$

$$y = 8$$

$$\Delta(y, X) = \{8, 2\} \rightarrow \text{back tracking}$$

$$X = \{0, 2, 10\}$$

$$L = \{2, 3, 3, 4, 5, 6, 7\}$$

Remove 7

~~$$y = 3$$~~

~~$$\Delta(y, X) = \{3, 7\}$$~~

$$y = 7$$

$$\Delta(y, X) = \{7, 5, 3\}$$

$$X = \{0, 2, 7, 10\}$$

$$L = \{2, 3, 4, 6\}$$

Remove 6

$$y = 4$$

$$\Delta(y, X) = \{4, 2, 3, 6\}$$

$$X = \{0, 2, 4, 7, 10\}$$

~~$$y = 6$$~~

~~$$\Delta(y, X) = \{6, 4, 1, 4\}$$~~

$$L = \{3\}$$

+

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Question 3

(a) Problem \rightarrow this algorithm may work forever

There is no guarantee that eliminating some break points will not introduce others, leading to an endless cycle.

Improved Breakpoint Reversal Sort Algorithm

Improved Breakpoint Reversal Sort (Π)

while $b(\Pi) > 0$

if Π has a decreasing strip

Among all reversals, choose reversal ρ minimizing $b(\Pi \cdot \rho)$

else

Choose a reversal ρ that flips an increasing strip in Π

$\Pi \leftarrow \Pi \cdot \rho$

output Π

return

#

(b) list a

(6)

Question 4

(a) Alignment

A	G	G	T	A	C	T	T
C	C	A	T	A	C	G	T
A	C	G	T	T	A	G	T
A	C	G	T	C	C	A	T
C	C	G	T	A	C	G	G

Profile matrix

A	3	0	1	0	3	1	1	0
C	2	4	0	0	1	4	0	0
G	0	1	4	0	0	0	3	1
T	0	0	0	5	1	0	1	4

Consensus

AC GT ACGT

Score $3+4+4+5+3+4+3+4$

Total Distance $2+1+1+0+2+1+2+1$

Sum 5 5 5 5 5 5 5 5

The motif finding Problem is a maximization problem while
The median String Problem is a minimization Problem

motif finding problem need to maximizing Score

median string problem need to minimizing total distance

#

17)

(b)

Motif finding problem

- motifs encode for a message in the "Genetic language"
- analyze the frequencies of pattern in DNA message
- establish regulatory Motif

hints

- hidden seq. is of length 8
- the pattern is not exactly the same in each array because random points mutation may occur in the seq.

Gold bug Problem

- symbols encode for a message in the "English language"
- analyze the frequencies of pattern in Gold bug message
- word in the English dictionary

hints

- English message
- Each symbol correspond to one letter
- No punctuation

but the motif finding Problem is harder than the Gold bug Problem

because

- we don't have the complete dictionary of Motifs
- Genetic language doesn't have standard Grammer

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