## Advanced Algorithms in Bioinformatics (P4) Sequence and Structure Analysis

Freie Universität Berlin, Institut für Informatik David Weese, Sandro Andreotti Sommersemester 2011

1. Review, 2011-05-11

Name, Vorname Matrikelnummer

Zur Bearbeitung des Reviews stehen Ihnen 50 Minuten zur Verfügung. Jeder Punkt entspricht in etwa einer Minute.

Geben Sie auf diesem Titelblatt und auf allen eventuell zusätzlich abgegebenen Blättern ihren Namen und ihre Immatrikulationsnummer an.

Schreiben Sie ihre Lösungen direkt auf die entsprechenden Aufgabenbögen. Sollte dort der Platz nicht ausreichen und Sie weitere Blätter benötigen, vermerken Sie dies bitte, damit wir auch den Rest ihrer Antwort finden und bei der Bewertung berücksichtigen können. Am Ende des Reviews sind sämtliche Aufgabenblätter wieder abzugeben.

## **Ergebnis:**

Aufgabe	maximal	erreicht
1	12	
2	15	
3	10	
4	13	×
Σ	50	

6 + 2 + 4 = 12 PtsExercise 1.

- 1. For the pattern AGATA and text AGATACGATATATAC apply the Horspool algorithm and explain the single steps.
- 2. What is the worst case runtime (number of comparisons) when searching a pattern of length m in a text of length n?
- 3. Give an example of a text T of length  $\geq$  20 and pattern P of length 5 where the number of character comparisons equals the worst case and P does not occur in T.

1) Ilropiocessis

A GET A

shipt table:

13 12 F5 V

II - Searching

POS = 0

A GATA C GATATATA C

5=0 => patternoccois at position 1

AGATA ATA

& [A] = 2 = ) shift 2 6 \$ A, & [6] = 3 C = A , L [A] = 2 T \$ 6, & [A] = 2

T = 6, & [A] = 2

V 0/6 pos is not & n-m => algoriths ands

output: pottern acconsat position 1

O (m.n) V212

## Exercise 2. 6 + 6 + 3 = 15 Pts

The Myers Bitvector algorithm uses binary encoding of the dynamic programming matrix.

1. Use the bitvectors to fill out the dynamic programming matrix

$VN_1$	=	000000
$VP_1$	=	111110
$D0_2$	=	111110
$HN_3$	=	111100
$HP_3$	=	000010

		$t_1$	$t_2$	$t_3$
	0	0	0	0
$p_1$	1	0	1	1
$p_2$	2	1	0	1
$p_3$	3	5	į	0
$p_4$	4	3	2	į
$p_5$	5	٧(	3	2
$p_6$	6	2	4	3

- 2. Below you find the pseudocode of the Myers Bitvector algorithm where the variables are renamed to  $\alpha$ ,  $\beta$ ,  $\gamma$ ,  $\delta$ ,  $\theta$ . Map these identifiers back to the original names D0, HN, HP, VN, VP.
- 3. How can you modify the algorithm to compute edit distance (global alignment) instead of the semi-global alignment?

```
// Preprocessing for c \in \Sigma do B[c] = 0^m od for j \in 1 \dots m do B[p_j] = B[p_j] \mid 0^{m-j}10^{j-1} od \delta = 1^m; \ \gamma = 0^m; score = m;
```

```
// Searching
 for pos \in 1 \dots n do
     X = B[t_{pos}] \mid \gamma;
      \alpha = ((\delta + (X \& \delta)) \land \delta) \mid X;
      \theta = \delta \& \alpha;
     \beta = \gamma \mid \sim (\delta \mid \alpha);
X = \beta \ll 1;
     \gamma = X \& \alpha;
      \delta = (\theta \ll 1) \mid \sim (X \mid \alpha);
      // Scoring and output
     if \beta \& 10^{m-1} \neq 0^m
        then score += 1;
         else if \theta \& 10^{m-1} \neq 0^m
                  then score = 1;
               fi
     fi
      if score \leq k report occurrence at pos fi;
 od
```

2) 
$$x = D0$$
 V

 $\beta = HPV$ 
 $y = VNV$ 
 $\delta = MMVPI$ 
 $\theta = HNV$ 

3) modicalding [  $0^{m-1}$  [ to

the second definition of  $X$ 
 $\Rightarrow X = HPZZI | 0^{m-1}$  1

V 3/3

## Exercise 3. 6 + 4 = 10 Pts

- 1. State the q-gram Lemma for contiguous shapes and prove it.
- 2. Is the threshold we compute by that Lemma tight (if  $\geq$  0)? Justify your answer.

THE THE

1) of grown: a short subsequer of length of let pard S be two strings of length w with at most K differences

then the number of possible of grams arounds to
with - of and the number of of-grams that
with - of and the number of or sometiment to kg
ten be at a destroyed at most arounds to kg

threshold:

t = W+1-9-9h = W+1-9(K+1)

Pard S share at least + common of grams

2) Yes. degrade and stopen that the

A DOWN

why?

1/4

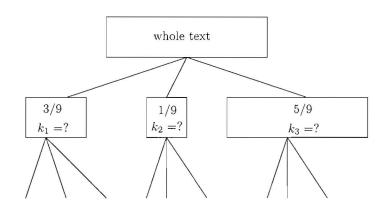
Exercise 4. 6 + 4 + 3 = 13 Pts

1. Prove the pigeonhole Lemma:

**Lemma 1.** Let Occ match P with k errors,  $P = p^1, \ldots, p^j$  be a concatenation of subpatterns, and  $a_1, \ldots, a_j$  be nonnegative integers such that  $A = \sum_{i=1}^j a_i$ . Then, for some  $i \in 1, \ldots, j$ , Occ includes a substring that matches  $p^i$  with  $|a_ik/A|$  errors.

2. Anne, Paul und Peter want to use hierarchical filtering. They are asked for error bounds  $(k_1, k_2, k_3)$  for hierarchical verification. The pattern shall be searched with 5 errors and is split in three parts.

Anne suggests (1,0,2), Paul (1,0,1) and Peter (1,1,2).



- (a) Are all three error bounds valid? Justify your answer.
- (b) Who suggested the best bounds? Justify your answer.

1) Let hi be the number of errors in pi  
then 
$$N = \frac{2}{5}$$
 hi

according to the Lemma Ji. [ail ] > ki

Than applying chaining rules:

now it is easy to device the controdiction

=> K7K 5