Bioinformatics

(a) Parameters of the positional independence of a transcription factor binding site were estimated by the experimental positional nucleotide frequencies shown in the following table:

/	1	2	3	4	5	6 γ	
T	0.16	0.05	0.01	0.03	0.12	0.14	
C	0.08	0.04	0.01	0.03	0.05	0.11	
A	0.68	0.11	0.02	0.90	0.16	0.51	
$\backslash G$	0.08	0.80	0.96	0.04	0.67	$\begin{pmatrix} 0.14 \\ 0.11 \\ 0.51 \\ 0.24 \end{pmatrix}$	

Explain what a *logo* is and determine the parameters of the logo graph. Compute the information content of one column. [8 marks]

- (b) Discuss the complexity of the algorithm for finding a global alignment between two DNA sequences that have a high degree of similarity. Present an example and analyse it using the following scoring parameters: +1 for match, -1 for mismatch, and d = -1 for a linear gap penalty. [7 marks]
- (c) In modelling a metabolic process, describe the advantages and disadvantages of using a stochastic approach (for example agents) as opposed to using a set of deterministic differential equations.