

WEST BENGAL STATE UNIVERSITY

B.Sc. Honours 6th Semester Examination, 2022

BOTADSE05T-BOTANY (DSE3/4)

BIOINFORMATICS

Time Allotted: 2 Hours

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Full Marks: 40

 $1 \times 16 = 16$

The figures in the margin indicate full marks. Candidates should answer in their own words and adhere to the word limit as practicable. All symbols are of usual significance.

- 1. Answer *all* the following questions briefly:
 - (a) What is a cladogram?
 - (b) Define orthologs.
 - (c) What is dynamic programming?
 - (d) What is CLUSTAL W?
 - (e) Name a software used to create phylogenetic tree.
 - (f) Define FASTA.
 - (g) Name the NCBI database for transcriptomic data.
 - (h) Name two tools used in phylogenetic analysis.
 - (i) What is consensus sequence?
 - (j) Define molecular docking.
 - (k) What is MSA?
 - (l) In the diagram which node represents the most recent common ancestor for organism B and C?



- (m) What is e-value of alignment score?
- (n) Which kind of mutation is more likely to be encountered: Transition or transversion?
- (o) Expand DDBJ.
- (p) Define topology.

CBCS/B.Sc./Hons./6th Sem./BOTADSE05T/2022

2.		Answer any <i>eight</i> questions from the following:	$3 \times 8 = 24$
	(a)	What is PIR? Describe the various resources of PIR.	1+2
	(b)	What do you understand by sequence alignment? Differentiate between global and local alignment.	1+2
	(c)	Discuss the importance of publically available biological databases in Bioinformatics.	3
	(d)	State principle of parsimony. What are the basic premises of concept of biological parsimony?	1+2
	(e)	What is Bootstrap test? What are its application and limitations?	1+2
	(f)	What is bioinformatics? Mention its role in crop improvement.	1+2
	(g)	Give one example each of (i) Nucleotide database, (ii) Protein database, (iii) Gene expression database.	
	(h)	What is a BLAST tool used for? What is the format used for submitting a sequence in a search base? Name the type of blast program used for proteins and nucleotide sequences.	1+1+1
	(i)	Briefly explain how PAM is derived.	1+2
	(j)	What is genomics? How does bioinformatics support genomic research?	1+2
	(k)	Explain monophyletic group, paraphyletic group and polyphyletic group with the help of a diagram.	1+1+1
	(l)	Define the following terms:	1 + 1 + 1
		(i) Lead compound	

- (ii) CADD
- (iii) Virtual screening.
- **N.B.**: Students have to complete submission of their Answer Scripts through E-mail / Whatsapp to their own respective colleges on the same day / date of examination within 1 hour after end of exam. University / College authorities will not be held responsible for wrong submission (at in proper address). Students are strongly advised not to submit multiple copies of the same answer script.

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