Reg. No. :	••••	
Name :		
Improve (2 Core Course in Life So	Degree (CBCSS – OBE – Regular/Supplement ement) Examination, April 2024 (020 Admission Onwards) ciences (Zoology) and Computational Biolog (NTALS OF COMPUTATIONAL BIOLOGY AND BIOINFORMATICS	у
Time : 3 Hours	Max. Marks	s : 40
	PART – A	
Write about <b>each</b> of the follo <b>1</b> mark.	owing in <b>2</b> or <b>3</b> sentences. <b>Each</b> question carries (6×	<1=6)
1. Proteomics.		
2. MEGA.		
3. SWISS-PROT.		
4. Cladogram.		
5. Maximum parsimony.		
6. Data mining.	VUR UNIVERS	
	PART – B	
Explain about any 6 of the fo	ollowing. Each question carries 2 marks. (6×2	2=12)
7. PDB.		
8. BLOSUM.		
9. BLAST.		

10. Genomics.

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- 11. Ultra metric trees.
- 12. Multiple sequence alignment.
- 13. PHYLIP.
- 14. Global sequence alignment.

## PART - C

Answer any 4 questions. Each question carries 3 marks.

 $(4 \times 3 = 12)$ 

- 15. Discuss the emerging areas of bioinformatics.
- 16. Briefly explain any two database search tools used in bioinformatics.
- 17. Explain PAM.
- 18. Comment on FASTA.
- 19. What are the goals of HGP?
- 20. Comment on KEGG.

## PART - D

Answer any 2 questions. Each question carries 5 marks.

 $(2 \times 5 = 10)$ 

- 21. Discuss various applications of bioinformatics.
- 22. Briefly explain the different types of databases used in bioinformatics.
- 23. Elaborate on the dynamic programming methodologies for sequence alignment.
- 24. Briefly explain any two distance based methods of phylogenetic tree construction.

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