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2022/TDC/ODD/SEM/ BOTDSE-501T(A/B)/145

TDC (CBCS) Odd Semester Exam., 2022

BOTANY

(5th Semester)

Course No.: BOTDSE-501T

Full Marks: 50
Pass Marks: 20

Time: 3 hours

The figures in the margin indicate full marks for the questions

Candidates have to answer either from Option—A

or Option—B

OPTION—A

Course No.: BOTDSE-501T (A)

(Analytical Techniques in Plant Science)

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- **1.** Answer any *three* of the following: $1 \times 3 = 3$
 - (a) What is confocal microscopy?
 - (b) What is the magnification of electron microscopy?
 - (c) What is freeze etching?
 - (d) What is freeze fracture?

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2.	Ans	swer any one of the following: 2	
	(a)	Differentiate between light microscopy and electron microscopy.	
	(b)	Write about cryofixation.	
3.	Ans	swer any one of the following:	
	(a)	Give a detailed account of transmission electron microscopy (TEM).	
	(b)	Describe briefly fluorescence microscopy and its applications.	
		Unit—II	
4.	Ans	wer any three of the following: 1×3=3	
	(a)	Define centrifugation.	
	(b)	What is autoradiography?	
	(c)	What is analytical centrifugation?	
	(d)	Define density gradient centrifugation.	
5.	Ans	wer any one of the following:	
	(a)	Write the principle of pulse chase experiment.	
	(b)	Write about marker enzymes.	
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6.	Ansv	wer any one of the following:						
	(a) Write a note on differential centrifugation.							
	(2)	Give a detailed account of ultracentrifugation.						
		UNIT—III						
7.	Ans	wer any three of the following: 1×3=3						
	(a)	What is R_f ?						
	(b)	What is stationary phase?						
	(c)	What is mobile phase?						
	(d)	What do you mean by GLC?						
8.	Ans	wer any one of the following: 2						
	(a)	Write a note on ion-exchange chromatography.						
	(b)	Write about TLC.						
9.	Ans	wer any one of the following:						
	(a)	Give an account of general aspects of chromatography.						
	(b)	Give an illustrated account of HPLC.						
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(3)

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UNIT-IV

		OMI—IV	
10.	Ans	swer any three of the following:	1×3=3
	(a)	Define mass spectrometry.	
	(b)	Name a technique for protein identification.	n
	(c)	Mention two uses of mass spectrometry	7.
	(d)	Define nucleotides.	
11.	Ans	swer any one of the following:	2
	(a)	Write a note on characterisation proteins.	of
	(b)	Give a brief account of AGE.	
12.	Ans	swer any one of the following:	5
ō	(a)	Describe briefly X-ray diffraction and it uses.	s
	(b)	Give a detailed account of PAGE.	
		UNIT—V	
13.	Ans	swer any three of the following:	×3=3
	(a)	Define data.	1
	(b)	What is sample?	
	(c)	What is statistics?	
	(d)	Define parameter.	
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14.	Answer any one of the following: 2
ri.	(a) Write a note on arithmetic mean.
	(b) Give an account of range.
15.	Answer any one of the following: 5
	(a) Write notes on: 2½+2½=5
	(i) Median
	(ii) Mode
	(b) Give a detailed account of chi-squared test for goodness of fit.
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OPTION—B

Course No.: BOTDSE-501T (B)

(Bioinformatics)

UNIT-I

1.	An	Answer any <i>three</i> of the following: 1×3=3						
in the	(a)	What is bioinformatics?						
	(b)	Give the full form of NCBI.						
	(c)	Name a nucleotide database.						
	(d)	Name the different branches of						
	lar.	bioinformatics.						
2.		swer any one of the following:	2					
	(a)	Classify the biological database with examples.						
	(b)	Write a note on protein data bank (PDB).						
3.	Ans	wer any one of the following:	5					
	(a)	Write a note on biological database retrieval system.						
	(b)	What are the different aspects of						

bioinformatics including

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4.	Answer any three of the following:				
		all Treath Information of Removes			
	(a)	Give the full form of BLAST.			

- (b) Name the database of literature from
- MEDLINE.
- (c) What does PIR stand for?
- (d) Write the full form of DDBJ.

5. Answer any one of the following:

- (a) What is the procedure for sequence submission to NCBI?
 - (b) Write two salient features of SWISS-Port.
- 6. Answer any one of the following:
 - (a) What are the different steps for performing BLAST?

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(b)	Write	notes	on	YTH's	
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- (i) EMBL Bank
- (ii) Protein Information Resource

UNIT-III

7. Answer any three of the following:

 $1 \times 3 = 3$

- (a) Which alignment is suitable for aligning closely related sequence?
- (b) Write a name of sequence alignment tool.
- (c) Global alignment is based on which algorithm?
- (d) Which alignment method is suitable for finding out conserved patterns in DNA?
- 8. Answer any one of the following:
 - (a) What is multiple sequence alignment?
 - (b) Write about BLOSUM.

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- 9. Answer any one of the following: 5
 (a) Write a note on MSA by CLUSTAL W.
 (b) Write a note on the different types of sequence alignment methods.
 UNIT—IV
 10. Answer any three of the following: 1×3=3
 - (a) What are population trees?
 - (b) Write the name of a software used for phylogenetic analysis.
 - (c) What is phylogeny?
 - (d) What do you mean by common ancestor in phylogenetic tree?
- 11. Answer any one of the following:
 - (a) What are the different types of phylogenetic trees?
 - (b) What limitation do molecular phylogenies have?

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(11)

12.	Answer	any	one	of	the	following		5.00
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- Write a note on the methods employed in phylogenic analysis.
- (b) What is the significance of molecular phylogenic analysis?

UNIT-V

13. Answer any three of the following:

 $1 \times 3 = 3$

- Write the full form of QSAR.
- What are the two main different aspects (b) of drug discovery?
- Name a docking software.
- What does ADMET stand for?

14. Answer any one of the following:

- (a) What is Lipinski rule of five?
- (b) Write two applications of bioinformatics in drug discovery.

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15. Answer any one of the following:

(a) How are bioinformatics techniques used for crop improvements?

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(b) Write a note on computer-aided drug designing.

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