Indian Institute of Technology Kharagpur

School of Bioscience Class Test (SLOT-1)

	Class Test (SLOT-1)	
Sub: Science of Living Systems	Sub No: BS20001	Full Marks: 20
Session: Spring, 2018-19	Date: 13-02-2019	Time: 30 minutes
NAME:	ROLLNO:	DEPT:
Choose (tick) the (O!	NE) correct answer OR wri	te in a few words
1. During DNA replication, helicase enzyme un DNA. In a PCR reaction, we use an alternative (A) A special buffer with high salt concentration (C) A special DNA polymerase that can denature	mechanism for DNA unwind on (B)High t	ding. What is that?
2. In gel electrophoresis different sized DNA m (A) DNA is positively charged and hence migrate (B) Different DNA molecules separate according (C) Smaller molecules migrate faster (D) DNA is visualized in the gel by staining with	rates towards the negative terning to mass	minal in the applied electric field gradient
3. Griffith's experiment proving DNA as the go (A) Termination (B) Transform		the principle of (D) Translation
4. The accepted theory for DNA replication is	_	
(A) conservative theory (B) dispersive	theory (C) semi-conservation	tive theory (D) evolutionary theory
5. Write True/False against each statement:(A) If the GC content of a single stranded DNA(B) If we replace the DNA polymerase of our beautiful and the content of the polymerase of our beautiful and the content of the polymerase of the pol	oody with Taq polymerase, it	can work just fine.
6. In classic Sanger DNA sequencing technique of the following is the correct combination?(A) All four ddNTPs and four dNTPs in same r(B) Each tube with one type of ddNTP and one	reaction tube	
2 and so on) (C) Each tube will have one type of dNTP and (D) Each tube will have one type of ddNTP and	• 1	
7. Following is the protein coding part of the D 5' ATG GCC CAA TAC TGG TGC ACG ACG What will happen to the protein product of the (A) Protein length will be unaffected (C) Amino acid composition will be changed	G TGC GAA GTC TGC ATA	A TTT TAA 3' the 10th codon from GAA into TAA? e shorter in length
8. Estimate the length of the protein coded by t introns are underlined. 5' GCACATATGGCGATACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGGACGAAGGGAAGGGAAGGGACGAAGGGAAGGGAAGGGAAGGAAGGGAAGGAAGGGAAGGAAGGGAAGAAGGAAGAAGGAAGAAGGAAGAAGAAGGAAGAAGAAGAAGAAGAAAGAAAA	<u>GCGGT</u> TGAGGCCGTTGT(-

9. Theoretically, a vast number of different proteins can be assembled from 20 different amino acids. How many

(D) $20^{10} \text{ X } 10^{20}$

(C) 10^{20}

polypeptide chains are possible that are 10 amino acids long? (B)20¹⁰

(A) 20 X 10

	ou remove the lac ope	erator (the repressor bi	nding site) what w	ill be the effect on the meta	bolic
state of the bacteria? (A) Lactose metaboliz	zing enzymes will be	produced irrespective	of the presence or	absence of lactose	
(B) Glucose metaboli	•				
(C) Lactose will neve(D) RNA Polymerase		· ·	never be synthesize	ed	
(D) It WIT olymeruse	will not be usic to on	na the promoter			
11. Anticodon is pres		(C) DNA	Œ		
(A) mRNA	(B)tRNA	(C) rRNA	(L	D) amino acid	
12. Tetracycline is an (A) lysing the bacteris		acteria by			
(B) interfering with b		by binding to RNA po	lymerase		
C) blocking bacterial	· · · · · · · · · · · · · · · · · · ·	~			
(D) blocking bacteria	I DNA replication by	binding to DNA polyi	nerase		
13. During transcripti	on, RNA polymerase	reads the template DN	IA strand in:		
(A)3'-5' direction	(B) 5'- 3' direction	(C) in both direct	ctions (D) does n	not require a DNA template	
14. Write True/False	against the following	statements:			
(A) The σ subunit is r	-		ymeras <u>e fro</u> m <i>E. co</i>	oli. T	
(B) Ribosomal RNA	(rRNA) is used as a te	mplate for protein syr	nthesis. F	_	
15. Which of the follo	owing can be an exam	ple of tertiary structur	e of a protein?		
(A) A multi-subunit p		•	_	A globular domain	
16. How are seconder	wy structures stabilized	Lin protoins?			
16. How are secondar (A) Through ionic bo	•	-	side chains		
(B) Through covalent	bonds joining differe	nt parts of the peptide	backbone		
(C) Through hydroge					
(D) Through hydroge	n bonds joining differ	ent parts of the peptid	e backbone		
17. What is a protein	domain?				
(A) The α-helical or β		•	, ,	it protein complex	
Part of a protein i	folded into a self-conta	ained 3D structure	(D) An unfolded p	part of a protein	
18. Hydrogen bonds i	n α-helices are				
(A) more numerous th			present at Phe resid		
(C) analogous to the s	steps in a spiral stairca	ise (D) roug	ghly parallel to the	helix axis	
19. In a folded protein	n, the nonpolar (hydro	phobic) amino acids t	end to be		
(A) hidden inside the	•	· · · -	osed on the outside	e of the protein	
(C) distributed randor	mly throughout the pro	otein (D) can	not be predicted		
20. Peptidyl transfera	se enzyme				
(A) is rRNA	(B) forms peptide	bonds (C) com	nponent of ribosom	de Dall of the above	