Registra	ation No :	
	Imber of Pages : 02 210	B.Tech PBT4I10
	4 th Semester Regular / Back Examination 2017-18	FD14IIV
	MOLECULAR BIOLOGY	
	BRANCH : BIOTECH	
	Time : 3 Hours	
	Max Marks : 100 Q.CODE : C1003	
210	Answer Part-A which is compulsory and any four from Part-B.	210
	The figures in the right hand margin indicate marks.	
	Answer all parts of a question at a place.	
	Part – A (Answer all the questions)	
Q1	Answer the following questions: <i>multiple type or dash fill up type:</i>	(2 x 10)
a)	Replication in prokaryotes differs from replication in eukaryotes for which of the following reasons?	
210		210
	than in eukaryotes.	
	 Prokaryotic chromosomes have a two origins of replication, whereas eukaryotic chromosomes have 100s. 	
	(iii) Prokaryotes produce Okazaki fragments during DNA replication, but	
	eukaryotes do not.	
	 (iv) Prokaryotic chromosomes have histones, whereas eukaryotic chromosomes do not. 	
210 b)	What is meant by the description "antiparallel" regarding the strands that	210
	make up DNA? (i) The twisting nature of DNA creates nonparallel strands.	
	(ii) Base pairings create unequal spacing between the two DNA strands.	
	(iii) One strand contains only purines and the other contains only pyrimidines.	
	(iv) The 5' to 3' direction of one strand runs counter to the 3' to 5' direction of the other strand.	
C)	The leading and the lagging strands differ in that	
210	 (i) the lagging strand is synthesized continuously, whereas the leading strand is synthesized in short fragments that are ultimately stitched 	210
	together	
	(ii) the leading strand is synthesized in the same direction as the movement	
	of the replication fork, and the lagging strand is synthesized in the opposite direction	
	(iii) the leading strand is synthesized at twice the rate of the lagging strand	
	(iv) the leading strand is synthesized by adding nucleotides to the 3' end of the growing strand, and the lagging strand is synthesized by adding	
210	nucleotides to the 5' end	210
d)	A new DNA strand elongates only in the 5' to 3' direction because	
	(i) replication must progress toward the replication fork(ii) DNA polymerase begins adding nucleotides at the 5' end of the template	
	(iii) can add nucleotides only to the free 3' end	
	(iv) the polarity of the DNA molecule prevents addition of nucleotides at the 3	
e)	end A particular triplet of bases in the template strand of DNA is 5' AGT 3'. The	
210	corresponding codon for the mRNA transcribed is	210
	(i) 3' UGA 5' (ii) 5' TCA 3'	

		210 210 210	210	210		210
	£)					
	f)	In the process of transcription, (i) proteins are synthesized (ii) mRNA attached	os to ribosomos			
g)		(ii) RNA is synthesized (iv) DNA is replica				
	a)	Codons are part of the molecular structure of .				
	5/	(i) mRNA (ii) tRNA				
		(iii) rRNA (iv) a protein				
210 h)	h)	A particular triplet of bases in the coding sequence of I	DNA is AAA. The	210		210
		anticodon on the tRNA that binds the mRNA codon is				
		(i) UUA (ii) AAA				
		(iii) UUU (iv) TTT				
	i)	How does termination of translation take place?				
		(i) The 5' cap is reached.				
		(ii) end of the mRNA molecule is reached.(iii) A stop codon is reached.				
		(iv) The poly A tail is reached				
210 j)		A single base substitution mutation is least likely to be	deleterious when t	210 he		210
	"	base change results in				
		(i) an amino acid substitution that alters the tertiary st	ructure of the prote	in		
		(ii) stop codon				
		(iii) n amino acid substitution at the active site of an en	zyme			
		(iv) codon that specifies the same amino acid as the or				
Q2 ₂₁₀		Answer the following questions: Short answer type			(2 x 10)	
QZ210	a)	What is Central Dogma?	210	210	(2 × 10)	210
	b)	What is Lampbrush chromosome?				
c) d) e)		How Okazaki segments are formed?				
		What is Selfish DNA?				
	e)	Write the different enzymes involved in post translation	al modification of			
		protein.				
	f)	What are the different proteins involved in Trp-operon?)			
210	g)	What is Gene silencing? 210 210	210	210		210
i)	h)	What is mutagenesis?	orintian process			
	j)	Differentiate between eukaryotic and prokaryotic trans What do you understand by overlapping genes?	subtion process.			
	3/					
~~		Part – B (Answer any four questi			(10)	
Q3	a)	Discuss the various experiments suggesting that 'DNA	as the genetic		(10)	
		monto vial?			(10)	
	b)	material'.				
210	b)	material'. Explain split genes and its significance.	210	210	(10)	210
		Explain split genes and its significance.			(5)	210
	b) a) b)	Explain split genes and its significance.			(5) (10)	210
	a)	Explain split genes and its significance. Discuss the Genome complexity. Write the significance Write a note on Cot curve analysis.	e of C- value Para		(5)	210
Q4	a)	 Explain split genes and its significance. Discuss the Genome complexity. Write the significance Write a note on Cot curve analysis. Explain the Gene structure in prokaryotes and Eukaryotes 	e of C- value Para		(5) (10)	210
Q4	a) b) a)	 Explain split genes and its significance. Discuss the Genome complexity. Write the significance Write a note on Cot curve analysis. Explain the Gene structure in prokaryotes and Eukaryo schematic diagram. 	e of C- value Para		(5) (10) (5) (10)	210
Q4	a) b)	 Explain split genes and its significance. Discuss the Genome complexity. Write the significance Write a note on Cot curve analysis. Explain the Gene structure in prokaryotes and Eukaryotes 	e of C- value Para		(5) (10) (5)	210
Q4 Q5	a) b) a) b)	 Explain split genes and its significance. Discuss the Genome complexity. Write the significance Write a note on Cot curve analysis. Explain the Gene structure in prokaryotes and Eukaryo schematic diagram. Discuss briefly the Translation process. 	e of C- value Para		(5) (10) (5) (10) (5)	210
Q4 Q5	a) b) a) b) a)	 Explain split genes and its significance. Discuss the Genome complexity. Write the significance Write a note on Cot curve analysis. Explain the Gene structure in prokaryotes and Eukaryo schematic diagram. Discuss briefly the Translation process. Briefly, discuss the different Models of DNA²¹ replication 	e of C- value Para otes with suitable	idox.	(5) (10) (5) (10) (5) (10)	
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