

School:

School of Science

Program/s:

Biomedical Science

Year: 2nd

Semester: 3rd

Examination:

End Semester Examination

Examination year:

December - 2021

Course Code: BM203

Course Name: Molecular Biology-II

Date: 06/12/2021

Time:

08:30 am to10:30 am

Total Marks: 40

Total Pages: 02

Instructions:

→ Write each session/answer on a new page.

→ Use of a calculator is permitted/not permitted:NA

→ Any other relevant instructions if any: Provide neat and clean diagram wherever necessary.

Q. No.	Details			Marks	CO	BTL
Q.1	A. Choose the correct answer. (5 x 1)				CO1	BTI
	1. The part of DNA coding for a protein is:				CO2	BT2
	a. Gene	b. Allele			002	D12
	c. Genome	d. Cistron	п		CO3	
	2. One of the nucleotides in a genic segment of	DNA underwent a mutation:				
	a. Point mutation	b. Pseudogene				
7,	c. Polymorphism	d. Missense mutation				
	3. With reference to human genome		2			
: 2	a. 98% of DNA is non-coding	b. 50% of DNA is mobile ele	ments			
	c. 3% of DNA is simple repeats	d. All are true	inents			
0	4. Splicing in prokaryotes is specifically observed in					
	a. rRNA genes	b. tRNA genes	TO AND ADDRESS OF THE			
	c. all genes expressing proteins	d. All of the above				
,=	5. The nucleophilic attack triggering the initiation					
	a. 3' Guanine	b. 5' Guanine				
	c. Adenine at branch point site	d. None of the above				
	B. Identify the below statements as True or Funderlined word(s) to make the statement to	False. If the answer is False, c rue.	hange the			
	1. The sugar found in RNA is called deoxy					
	2. The DNA molecule is double stranded and the RNA molecule is single stranded.					
	3. The process of translation occurs at the ribosome.					
	4. The job of mRNA is to pick up amino acids and transport them to the					
	ribosomes.	cids and transport them to the				
	5. Transcription must occur before translat	ion may occur				

		1 W/h - 4 '		Ar	1y 5	10	CO1	BT
		5. Differentiate between6. Explain C value par7. What are non-codin	n factors? stand by degeneracy of geneton Genetic Code and its uniquen monocistronic and polycis adox. g RNAs? Enlist non-coding I	de properties. Stronic mRNAs.		2 x 5)	CO2	вт
Q.3	Match A with B and C							
		A	В		1	.0	CO1	BT1
	_	Prokaryotic gene		C			CO2	BT2
	1	promoter	Poly U site	Transcriptional elongation			CO3	
	2	RNA splicing	Hairpin loop streture	holoenzyme				
	3	Apoenzyme	Consensus sequence	Precursor rRNA and tRNA				
	4	RNA processing	Phosphorylation	Rho-independent termination				
	5	Non-coding RNAs	Trans-esterification	RNA-interference		*		
	6	Eukaryotic gene promoter	Sigma factor	Pribnow box				
	7	Spliceosome	Methylation	Rho-dependent transcription termination				
	8	C-terminal domain	RNA:protein.	Self-splicing				
	9	Prokaryotic transcription termination	Gene-expression regulation	TATA binding protein				(
	10	RNA:DNA helicase activity	TATA box	small nuclear ribonucleoproteins (snRNPs/snurps)				
4 E	Explain in detail. (Provide figures if necessary)							
	1. Explain eukaryotic gene organization				10 (5 x 2)	CO2	? В	T1 T2