(103)輔仁大學碩士班招生考試試題

考試日期:103年3月7日第3節

本試題共 3 頁 (本頁為第 1 頁)

科目:	分子生物學
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系所組:營養科學

- I. Explain the following terms: (5 points for each sentence, 20 points)
 - (a) RNA editing
 - (b) MicroRNA
 - (c) Chromatin remodeling
 - (d) Wobble hypothesis
- II. What are three mechanisms of translational repression that are known to exist in eukaryotes? (15 points)
- III. Name four general types of postsynthetic processing reactions that are observed in RNA. (10 points)

 Briefly (one sentence or less) point out an example of each type. In your example, identify the type of RNA molecule involved (tRNA, mRNA, rRNA, etc.), the type of "processing" involved, and whether the example is characteristic of eukaryotes or prokaryotes, or both. Do not describe specific genes, sequences, complicated structures, or enzymes.
- IV. Choice questions (3 points for each question, 45 points)
- (1) An mRNA whose translation is controlled by binding of a metabolic end product such as flavin adenine dinucleotide or adenosylcobalamin is called a .
 - (a) Ribozyme
 - (b) miRNA
 - (c) riboswitch
 - (d) regulatory RNA
- (2) DNA methylation in eukaryotes plays a role in _____
 - (a) protection of restriction sites
 - (b) cancerous cell growth
 - (c) gene silencing
 - (d) methyl-directed mismatch repair
- (3) In a histone protein, what modification to lysine marks the nucleosome as a transcription target?
 - (a) physical separation of transcription from translation
 - (b) the presence of chromatin instead of naked DNA
 - (c) the larger size of the chromosomes
 - (d) the presence of introns
- (4) What mRNA modification, just 5' to the start codon, is used by the eukaryotic cell to enable location of the first AUG codon?
 - (a) a sequence of UAUAUA
 - (b) a GC "box" that is recognized by eIFI
 - (c) 7-methylguanine cap on the 5' end of the mRNA
 - (d) a short sequence that forms a hairpin loop
- ※ 注意:1.考生須在「彌封答案卷」上作答。
 - 2.本試題紙空白部份可當稿紙使用。
 - 3.考生於作答時可否使用計算機、法典、字典或其他資料或工具,以簡章之規定為準。

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肝目	:	分子生物學		系所組:營養科	-學
(5)	mito	ochondria?	sp70 proteins play in delive		of action such as the
	` '		olding to allow protein to pro lysis of the unfolded protein	perly traverse membranes	
	(c)	keeps the protei	n in an unfolded state to allo		
	(a)	transport	on of an intermediate fold	ed protein that is present	only during membrane
(6)		vivo, which of the TA box?	the following transcription	initiation factors is requ	ired for binding to a
		TFIID			
	` '	TFIIE TFIIF	•		
	` '	TFIIH			
	_		DNIA		II I DAY DAY
(7)		merase tra	, RNA polymeraseanscribes the major structu		· · · · · · · · · · · · · · · · · · ·
		I; III; II			
	` '	II; I; III			
	, ,	III; II; I I; II; III			
	` ,				
(8)		seudogene is a _ nonexpressed g	• ana		
			ple promoter regions that cau	uses the start of transcription	to be mistaken
	(c)	mutated gene			
	(d)	gene that codes	for a non-functional protein		
9)	Ban	aHI has a restr	iction site of G↓GATCC w	vhile <i>Hpa</i> I has a restrictio	n site of GTT↓ACC.
-			mHI produces	and <i>Hpa</i> I produces	•
	` '	a 3' overhang;	a 5' overhang		
	` '	a 5' overhang; a 5' overhang;	a 3' overhang blunt ends		
	` '	a 3' overhang;	blunt ends		
	(u)	a 5 Overnang,	orunt ends		
10)		at is a polyribo			
			synthesize different subunits		
	, ,		nes all attached to the same mare covalently bonded to each		
			only produces proteins that	- ·	

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科目:	分子生物學	系所組:營養科學	
(a) (b) (c)	hat technique is used to DNA sequencing restriction digestion polymerase chain rea electrophoresis	to amplify portions of DNA, usually about 500 base pairs in siz	e?
(a) (b) (c)		o describe genetic variation that occurs among individuals of or loss of restriction sites that can be detected by Southern and	
(a) (b) (c)	factors ai	lenosyl methionine; ATP lavin; TPP ADH	and the
(a) (b) (c)	segment of DNA that segment of mRNA sy	s a: ulting from endonuclease action. is an intermediate in the synthesis of the lagging strand. rnthesized by RNA polymerase. t is a subunit of the 30S ribosome.	
(a) (b) (c)	a region of DNA that the specific binding s the position of interna	has been damaged by mutation. ite of a repressor, polymerase, or other protein on the DNA. ally double-stranded regions in a single-stranded DNA molecule. cular gene of a chromosome.	
	tch the protein or stru points for each question	uctural feature on the left with one appropriate description on on, 10 points)	the right.
	activator helix-turn-helix leucine zipper repressor zinc finger	 (a) a positive regulator (b) a negative regulator (c) facilitates transcription only when bound to a signal molecule (d) a DNA-binding structural motif found in many prokaryotic regulatory proteins (e) a structural feature involved in protein-protein interactions between some regulatory protein monomers (f) a protein that dissociates from DNA when bound to a signal molecule (g) a DNA-binding structural motif found in many eukaryotic regulatory proteins 	

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