Name: MIDTERM EXAM ANSWER KEY [ANSWERS IN RED]

Biology 200 - Section 921

Midterm Examination - June 29, 2006 Duration of Examination: 50 minutes Total marks: 50 (25 % of course grade)

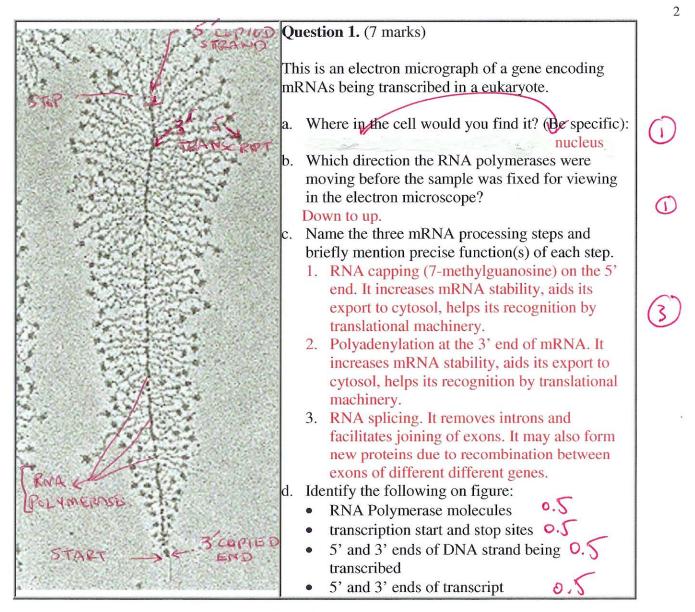
IMPORTANT INSTRUCTIONS:

- 1. ANSWER ALL QUESTIONS ON THE EXAMINATION BOOKLET
- 2. Read carefully, think, then begin to answer the question.
- 3. Fill in your Name and Registration Number at the top of every page.
- 4. Please use **pen** only, **not** pencil.
- 5. Please make sure that the examination booklet has a total of **7 questions** and **5 pages**.

Question	1	2	3	4	5	6	7	Total
Student mark								
Marks available	7	4	15	10	6	2	6	50

Genetic Code Table

		U	C second	A position	G		
		UUUPhe	UCUSer	UAUTyr	UGUCys	U	
	U	UUCPhe	UCCSer	UACTyr	UGCCys	С	
		UUALeu	UCASer	UAAstop	UGAstop	Α	
		UUGLeu	UCGSer	UAGstop	UGGTrp	G	
1		CUULeu	CCUPro	CAUHis	CGUArg	U	3
S	С	CUCLeu	CCCPro	CACHis	CGCArg	С	r
t		CUALeu	CCAPro	CAAGln	CGAArg	А	d
		CUGLeu	CCGPro	CAGGln	CGGArg	G	
Р		AUUIle	ACUThr	AAUAsn	AGUSer	U	Р
0	А	AUCIle	ACCThr	AACAsn	AGCSer	С	0
S		AUAIle	ACAThr	AAALys	AGAArg	А	S
i		AUGMet	ACGThr	AAGLys	AGGArg	G	i
t		GUUVal	GCUAla	GAUAsp	GGUGly	U	Т
i	G	GUCVal	GCCAla	GACAsp	GGCGly	С	i
0		GUAVal	GCAAla	GAAGlu	GGAGly	А	0
n		GUGVal	GCGAla	GAGGlu	GGGGly	G	n
							J



Question 2. (4 marks). The following mRNA sequence was used as a template to translate into a if this is template, then protein. 5' CCUAUGGGUAGGCUAC 3'

3'-CAUCGGAUGGGUAUCC 5'-GUAGCCUACCCAUAGG A) If you were told that this segment of RNA was close to the 3' end of an mRNA that encoded a large protein, would you know which reading frame was used [see Genetic code table on page 1]?

> CC UAU GGG UAG GCU AC Tyr Gly Stop

3

B) If you were told that this segment of RNA was close to the 5' end of an mRNA that encoded a large protein, would you know which reading frame was used?

CCU AUG GGU AGG CUA C Met Gly Arg Leu

2

Question 3. (15 marks). Short answers (attempt five):

a. Without a continual input of energy, animal cells will burst. Why?

Animal cells contain high concentration of many molecules that will cause the osmotic influx of water. Unless ions are constantly pumped out (e.g. Na+ pumped out by Na+-K+ pump) to maintain an osmotic balance, cells will eventually burst.

b. How do histones interact with DNA to form nucleosome?

Histones have a high proportion of positively charged amino acids (lysine and arginine). These positive charges help the histones bind tightly to the negatively charged sugar-phosphate backbone of DNA.

c. What is the role of flippases in generating the lipid asymmetry in cell membranes? Flippases are the enzymes, which catalyze the <u>selective</u> transfer of <u>specific</u> phospholipid molecules from one monolayer to the other opposite monolayer in a lipid bilayer membrane. This creates asymmetry in the bilayer membranes.

(Some students may draw a figure to illustrate the mechanism of flippases action)

d. Plant cells, fungi and bacteria do not have the Na⁺-K⁺ pumps in their plasma membrane. Explain how the symport-mediated transport of ions takes place in plant cells, fungi and bacteria.

Plant cells, fungi and bacteria use H^+ - ATPase (H^+ -pump) on their plasma membrane for symportmediated transport. H^+ - ATPases pump H^+ out of the cell, thus creating an electrochemical proton gradient (proton conc. higher outside than inside). The import of many molecules (sugars and amino acids) is driven by H^+ symports, which use the electrochemical gradient of H^+ across the plasma membrane (similar to electrochemical gradient of Na⁺ in case of animal cells).

e. Briefly describe the key structure-function relationship of aminoacyl-tRNA synthetases. Structure: are the enzymes with two binding sites, one for a specific amino acid and the other for a specific tRNA.

Function: catalyze the ATP-mediated covalent coupling of an amino acid to its appropriate set of tRNA molecules

f. Briefly describe the key structure-function relationship of spectrin protein in human red blood cells.

Structure: Long, thin, flexible rod-like proteins linked to specific transmembrane proteins in plasma membranne; main component of the cell cortex.

Function: Provides support for the plasma membrane; maintains cell shape.

Question 4. (10 marks). Name one major similarity and one major difference between the following (attempt 5):

A. The large and small subunits of ribosome

Similarity(ies): Both are made up of rRNA and proteins; participate in translation.

<u>Difference(s)</u>: Size: The large subunit (60S) and small subunit (40S)

Function: The small subunit matches the tRNA yo the codons of the mRNA. The large subunit catalyzes the formation of the peptide bond between amino acids to form a polypeptide chain.

B. Na⁺-driven Glucose Pump and Glucose uniport carrier

Similarity(ies): Both are carrier type of membrane-bound transporters. Both transport glucose.

<u>Difference(s)</u>: Na⁺-driven Glucose Pump uses energy in Na+ gradient to pump glucose against its concentration gradient. Glucose uniport carrier facilitates passive transport of glucose from its high to low concentration.

C. Scanning and transmission electron microscopy

<u>Similarity(ies)</u>: Both use a beam of electrons to visualize structures. Have high resolution power. Both require samples that have been chemically fixed.

<u>Difference(s)</u>: Scanning microscopy provides 3-dimensional images and greater details of surface of the structure(s). TEM provides greater resolution between two subcellular structures.

D. Proteasome and lysosome

<u>Similarity(ies)</u>: Both lysosome and proteasome degrade/hydrolyze proteins.

<u>Difference(s)</u>: Lysosome is a membrane-bounded organelle which contains a number of digestive enzymes including protein-degrading enzymes, proteases active at low pH. Proteasome is a large protein complex in the cytosol that degrades cytosolic proteins that have been marked for destruction by ubiquitin.

E. Protein domain and protein subunit:

<u>Similarity(ies)</u>: Both protein domain and subunit are parts of protein structure.

<u>Difference(s)</u>: A protein domain is a compact and stable folded region of a polypeptide. A protein subunit is a folded polypeptide that interacts with other subunits to form a quaternary protein structure in a multi-subunit protein.

F. The α helix and the β sheet

G. <u>Similarity(ies)</u>: Both α helix and the β sheet represent secondary protein structures.

<u>Difference(s)</u>: An α helix represent a linear sequence of amino acids folded into a right-handed helix stabilized by internal H-bonds between backbone atoms (H of amino and O of carboxyl). A β sheet represents a folding pattern in which polypeptide chains associate with each other through H-bonds.

Question 5. (6 marks). For the following carrier proteins, write the carrier type (e.g. uniport, symport or antiport), cellular location, energy source, and function(s):

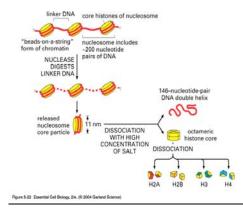
- a. <u>Na[±]-K[±] pump (Na[±]-K[±] ATPase)</u>
 - (i) Type of carrier: Antiport
 - (ii) Cellular location(s): plasma membrane of most animal cells
 - (iii) Energy source: ATP hydrolysis
 - (iv) Function(s): Active export of Na^+ and import of K^+
- b. Bacteriorhodopsin
 - (i) Type of carrier: Uniport
 - (ii) Cellular location(s): plasma membrane of some bacteria
 - (iii) Energy source: light
 - (iv) Function(s): active export of H^+ out of the cell
- c. <u>Ca²⁺</u>-pump (Ca²⁺ ATPase)
 - (i) Type of carrier: Uniport
 - (ii) Cellular location(s): plasma membrane (and ER membrane) of eukaryotic cells
 - (iii) Energy source: ATP hydrolysis
 - (iv) Function(s): active export of Ca^+

Question 6. (2 marks). Fill in the blanks:

- a. Molecular chaperones are large proteins which assist proper folding of other polypeptides.
- b. The most highly condensed form of interphase chromatin is called heterochromatin.
- c. All the lipids found in membranes are said to be <u>amphipathic</u> because they have one hydrophilic end and one hydrophobic end.
- d. Each group of three consecutive nucleotides in RNA is called a codon.

Question 7. (6 marks). <u>Name</u> and <u>briefly explain</u> the experimental methodology to study the following biological structures/processes (**attempt one**):

a. <u>Macromolecular composition of nucleosomes</u>



- 1. Nuclease digestion of linker DNA
- 2. Dissociation of proteins and DNA by high [Salt]
- 3. Dissociation of individual histone subunits
- 4. SDS-Page, amino acid sequencing or Mass Spec (MS)

b. Investigate the mobility of a lipid within plasma membrane

Its mobility in the membrane can be studied by the florescence dye or radiactive labelling and the FRAP technique. Labelling with florescence dye, \rightarrow photobleaching selective lipids in the membrane, \rightarrow recovery of fluorescence after certain time. Half marks if proteins are labelled.

