

(University of Choice)

MASINDE MULIRO UNIVERSITY OF SCIENCE AND TECHNOLOGY (MMUST)

MAIN CAMPUS

UNIVERSITY EXAMINATIONS 2018/2019 ACADEMIC YEAR

FOURTH YEAR FIRST SEMESTER EXAMINATIONS

FOR THE DEGREE OF BACHELOR OF SCIENCE MEDICAL BIOTECHNOLOGY SUPPLEMENTARY/SPECIAL EXAM

COURSE CODE: BMB 421

COURSE TITLE: GENE EXPRESSION SYSTEMS AND

SEQUENCING

DATE:

TIME:

INSTRUCTIONS TO CANDIDATES

This paper is divided into three sections, **A B** and **C**, carrying respectively: Multiple Choice Questions (**MCQs**), Short Answer Questions (**SAQs**) and Long Answer Questions (**LAQs**).

TIME: 2 Hours

MMUST observes ZERO tolerance to examination cheating

This Paper Consists of 5 Printed Pages. Please Turn Over.

SECTION A: MULTIPLE CHOICE QUESTIONS (20 MARKS)

- 1. When scientists were attempting to determine the features of the genetic code, Crick and co-workers found that when three base additions or three base deletions occurred in a single gene, the wild type phenotype was sometimes restored. This observation supported the hypothesis that
 - A. The code is triplet.
 - B. AUG is the initiating triplet.
 - C. The code is overlapping.
 - D. There are three amino acids per base.
- 2. In 1964, Nirenberg and Leder used the triplet binding assay to determine specific codon assignments. A complex of which of the following components was trapped on the nitrocellulose filter?
 - A. Ribosomes and DNA
 - B. Free tRNAs
 - C. Charged tRNA, RNA triplet, and ribosome
 - D. Uncharged tRNAs and ribosomes
- 3. Which of the following is true?
 - A. Every amino acid is coded for by a single codon.
 - B. There are more amino acids than there are codons.
 - C. Every codon codes for an amino acid.
 - D. Each codon in a gene codes for no more than one single amino acid.
- 4. The genetic code is "degenerate" because
 - A. There are more codons than amino acids.
 - B. There are more amino acids than codons.
 - C. Different organisms use different codons to encode the same amino acid.
 - D. Some codons specify more than one amino acid.
- 5. What is the name given to the three bases in a messenger RNA that bind to the anticodon of tRNA to specify an amino acid placement in a protein?
 - A. Protein
 - B. Anti-anticodon
 - C. Cistron
 - D. Codon
- 6. When studying the initiation of transcription, one often finds consensus sequences located in the region of the DNA where RNA polymerase(s) bind. Which are common consensus sequences?
 - A. CAAT, TATA
 - B. GGTTC, TTAT
 - C. TTTTAAAA, GGGGCCCC
 - D. Any trinucleotide repeat
- 7. An intron is a section of
 - A. Protein that is clipped out post-translationally.
 - B. RNA that is removed during RNA processing.
 - C. DNA that is removed during DNA processing.

- D. Transfer RNA that binds to the anticodon.
- 8. Three posttranscriptional modifications often seen in the maturation of mRNA in eukaryotes are
 - A. 5'-capping, 3'-poly (A) tail addition, splicing
 - B. 3'-capping, 5'-poly (A) tail addition, splicing
 - C. Removal of exons, insertion of introns, capping
 - D. 5'-poly (A) tail addition, insertion of introns, capping
- 9. What is the initiator triplet in both prokaryotes and eukaryotes and what is the amino acid coded for by this triplet?
 - A. UAA, no amino acid coded.
 - B. UAA, methionine.
 - C. AUG, arginine.
 - D. AUG, methionine.
- 10. Which type of RNA molecule carries an amino acid to the ribosome?
 - A. tRNA
 - B. rRNA
 - C. mRNA
 - D. siRNA
- 11. The term "peptidyl transferase" relates to
 - A. Base additions during mRNA synthesis.
 - B. Peptide bond formation during protein synthesis.
 - C. Elongation factors binding to the large ribosomal subunit.
 - D. 5' capping of mRNA.
- 12.By their experimentation using the <u>Neurospora</u> fungus, Beadle and Tatum were able to propose the hypothesis that:
 - A. Prototrophs will grow only if provided with nutritional supplements.
 - B. Several different enzymes may be involved in the same step in a biochemical pathway.
 - C. The role of a specific gene is to produce a specific enzyme.
 - D. More than one codon can specify a given amino acid.
- 13. What is the term which refers to a contiguous set of bacterial genes which are under coordinate control?
 - A. Lysogen.
 - B. Prototroph.
 - C. Operon.
 - D. Allosteric.
- 14. Which term most appropriately refers to a *trans* acting regulatory factor?
 - A. Translation
 - B. RNA processing
 - C. DNA binding protein
 - D. Helicase activation
- 15. The *lac* operon

- A. Is under negative and positive control.
- B. Is under positive control only.
- C. Is normally expressed constitutively.
- D. Is an example of tissue-specific expression?
- 16. What is the function of cAMP in regulation of the *lac* operon?
 - A. Activates a repressor protein
 - B. Activates an activator protein
 - C. Inactivates a repressor protein
 - D. Inactivates an activator protein
- 17. When a repressor binds to an operator, what process is affected?
 - A. Transcription of the operon is prevented.
 - B. DNA replication of the operon is prevented.
 - C. mRNA made from the operon cannot be translated.
 - D. Protein made from the operon does not function.
- 18. A mutant *E. coli* strain, grown under conditions that normally induce the *lac* operon, produces high amount of β -galactosidase. What is a possible genotype of the cells? (I = lac repressor gene; Z, Y, A = lac operon structural genes; P = lac promoter; O = lac operator)
 - A. lacI+ lacP+ lacO+ lacZ- lacY+ lacA+
 - B. $lacI^{+}$ $lacP^{+}$ $lacO^{c}$ $lacZ^{+}$ $lacY^{+}$ $lacA^{+}$
 - C. lacF lacP+ lacO+ lacZ- lacY+ lacA+
 - D. lacI+ lacP- lacO+ lacZ+ lacY+ lacA+
- 19. Consider a bacterial strain that has a mutant *lac* operator region that cannot bind repressor. If you were to introduce into this strain a wild type operator region on an F'element, the cell would
 - A. Synthesize lactose.
 - B. Display inducible expression of the *lac* operon.
 - C. Display constitutive expression of the *lac* operon.
 - D. Probably not catabolize lactose.
- 20. With regards to the *trp* operon,
 - A. Tryptophan is an inducer.
 - B. Tryptophan is a co-repressor.
 - C. Attenuation can halt replication.
 - D. None of the above.

SECTION B: SHORT ANSWER QUESTIONS (40 MARKS)

1. Give a detailed structure of a gene.	[8 Marks]
2. a) List any four eukaryotic promoter elements.	[4 Marks]
b) Differentiate between splicing and alternative splicing.	[4 Marks]
3. Describe RNA processing.	[8 Marks]
4. Describe characteristics of the genetic code.	[8 Marks]
5. a) List requirements for translation.	[4 Marks]
b) Which factors control gene expression?	[4 Marks]

SECTION C: LONG ANSWER QUESTIONS (40 MARKS)1. Discuss the Sanger dideoxy-sequencing technique. 2. Describe the human genome project.

[20 Marks] [20 Marks]