## Introduction to Biology - Monsoon 2022

## **End Semester Examination**

(Course outcomes: CO-2, CO-4 and CO-5)

| Max. Time: 3.0 hrs  | Max. Marks: 65                                  |
|---|---|
| 1. The following is the DNA sequence for the transcription initiation of the promoter region is boxed. Transcription begins at and including A/T base pair. |   |
| 5'TGGACTGCTA TAATAGCAGG GCTGCCGAAT GTGCTGCCAT 3'ACCTGACGAT ATTATCGTCC CGACGGCTTA CACGACGGTA   | ACGGCCATGG TTCTTAAAGT3' TGCCGGTACC AAGAATTTCA5' |
| (A) Which DNA strand (choose from top or bottom) serves as the transcription? Explain. [2 marks]  | e template strand for                           |
| (B) Fill in the first 6 nucleotides of the primary/ nascent mRNA [1 mark]   | transcribed from Gene A.                        |
| 5'  | 3'  |
| (C) Fill in the first four amino acids of Protein A encoded by Ge provided on the last page. [1 mark]   | ne A. Note: A codon chart is                    |
| N — — — — — — — — — — — — — — — — — — —   | C   |
| (D) The last 5 amino acids (amino acid105- amino acid109) at the Protein A are indicated below. Each of these amino acids is this protein.                  |   |
| N - pro <sup>105</sup> -asn <sup>106</sup> -ser <sup>107</sup> -met <sup>10</sup>   | <sup>8</sup> -leu <sup>109</sup> -C             |
| The DNA sequence encoding the above 5 amino acids is inc  | cluded within the sequence below                |

You isolate and sequence the following two different mutant alleles of Gene A that encode the above 5 amino acids. Each mutant allele is due to a point mutation that is bold and underlined. Which of these mutants will ALTER the folding of Protein A

Wild-type

Mutant 1 5'-AACCAAATTCCATGTTATAGC-3' Mutant 2 5'-AACCGTATTCCATGTTATAGC-3' 3'-TTGGTTTAAGGTACAATATCG-5'

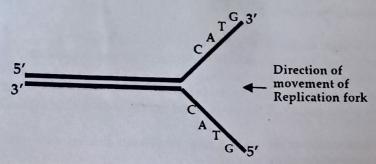
5'-AACCGAATTCCATGTTATAGC-3'

3'-TTGGCTTAAGGTACAATATCG-5'

Explain, in terms of the change in the reading frame and/ or amino acid sequence, why you selected this mutant and NOT the other. [3 marks]

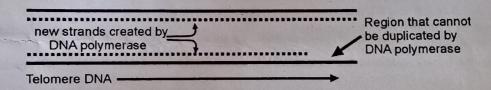
| 2.         | What is the  | e sequence (           | (1 to 4) is           | n which these                   | proteins function                                    | on during DNA                    | replication       | on                       |          |
|------------|--|------------------------|-----------------------|---------------------------------|--|----------------------------------|-------------------|--------------------------|----------|
| (          |  | primase                | (1104)1               |                                 | A helicase   |                                  |                   |                          |          |
|            | the second secon | ligase                 |                       | Di                              | A Helicuse   |                                  |                   |                          |          |
|            | A STATE OF THE PARTY OF THE PAR | polymeras              | e                     |                                 |  |                                  |                   |                          |          |
|            |  |                        |                       |                                 |  |                                  | ant of DN         | A from                   |          |
|            | Briefly exp<br>super-coili   | olain their fi<br>ng?  | unctions.             | Which enzy                      | ne relieves a re                                     | plicating segm                   | [5 n              | narks]                   |          |
| 3          | sequence o   | f events tha           | t will oc             | cur when the                    | odons and antic<br>in-coming tRNA<br>er the next pep | A sets into its I                | oing sit          | e. Reulaw                | 2        |
|            |  |                        |                       |                                 | 8/-1   |                                  |                   |                          |          |
|            |  |                        | (                     |                                 |  | )                                |                   |                          |          |
|            |  |                        |                       | -                               |  | er.                              |                   |                          |          |
|            | The diagra   | m below sh             | ows an i              | mRNA moleci                     | ale with various                                     | regions labell                   | led:              |                          |          |
| 1.         | The diagra   |                        |                       |                                 |  |                                  |                   |                          |          |
|            |  | _                      |                       |                                 |  |                                  | UAG               |                          | PPP-     |
|            | AAAAAAA  |                        |                       | AUG                             |  |                                  | 5                 | 6                        | 7        |
|            | 1  |                        | 2                     | 3                               | 4  |                                  |                   |                          |          |
|            | (A) Th   | nere is a pro          | blem wi               | th above diag                   | am. True or Fa                                       | lse. Explain y                   | our choice        | e [1.5 mar               | rks]     |
|            | (B) Ide  | ntify by nu            | mber the              | e region(s) tha                 | is/are:  |                                  |                   | [2.5 mar                 | ks]      |
|            | ( <i>b</i> ) ide   | coding (i.e            | . contair             | ns codons that                  | are part of the                                      | peptide)                         |                   |                          |          |
|            | b)   | non-codin              |                       |                                 |  |                                  |                   |                          |          |
|            | c)   | 3' end                 |                       |                                 |  |                                  |                   |                          |          |
|            | d)   | 5' end                 |                       |                                 |  |                                  |                   |                          |          |
|            | e)   | ribosome l             | oinding s             | site                            |  |                                  |                   |                          |          |
| 5.)        | Draw and o   | explain the regulation | (a) patte<br>of recon | rn of chromos<br>nbination duri | ome segregation                                      | on in mitosis a<br>meiotic progr | nd meiosi<br>ams. | s and (b) ti<br>[4 marks | he<br>i] |
| d          | Explain the  | underlying             | princip               | ole of: [4                      | marks]   |                                  |                   |                          |          |
| <b>y</b> . | (a) Poly   | merase cha             | in reacti             | ion                             |  |                                  |                   |                          |          |
|            |  | ger Sequen             |                       |                                 |  |                                  |                   |                          |          |
| 9.         | In response  | to hypoxia             | (lack o               | f oxygen), ma                   | mmalian cells<br>or regulating th                    | induce expression? [3 marks      | sion of gro       | oup of gen               | es.      |
|            | Explain the  |                        |                       |                                 | ATT BELLEVIE   |                                  |                   |                          |          |

## 8. Shown below is a segment of replicating DNA



- (A) On the schematic, draw the elongating DNA strands and label their 5' and 3' ends
- (B) To which strand (choose from top, bottom or both) can primer 5'CATG3' bind during replication?
- (C) Which strand (choose from top or bottom) is the template for discontinuous (lagging) strand synthesis? [3 marks]
- 19. Explain how telomerase and DNA polymerase operate together to lengthen the chromosomes. Label the 3' and 5' ends of the strands and modify this diagram to show where DNA polymerase and telomerase will lengthen the strands. Also, explain why DNA polymerase alone cannot accomplish the task of telomere DNA synthesis.

  [3 marks]



- 11. During protein synthesis, the thermodynamics of base paring between tRNAs and mRNAs sets the upper limit for the accuracy with which protein molecules are made. True or False. Explain your choice. [3 marks]
- One indication of the relative importance of various ATP-producing pathways is the  $V_{\text{max}}$  of certain enzymes of these pathways. The values of  $V_{\text{max}}$  of several enzymes from chest muscles used for flying of pigeon and pheasant are listed below.

|                        | $V_{\rm max}$ ( $\mu$ mol substrate/min/g tissue) |          |  |  |  |  |  |
|------------------------|---|----------|--|--|--|--|--|
| Enzyme                 | Pigeon  | Pheasant |  |  |  |  |  |
| Hexokinase             | 3,0   | 2.3      |  |  |  |  |  |
| Glycogen phosphorylase | 18.0  | 120.0    |  |  |  |  |  |
| Phosphofructokinase-1  | 24.0  | 143.0    |  |  |  |  |  |
| Citrate synthase       | 100.0   | 15.0     |  |  |  |  |  |
| Triacylglycerol lipase | 0.07  | 0.01     |  |  |  |  |  |

- (a) Discuss the relative importance of glycogen metabolism and fat metabolism in generating ATP in the chest muscles of these birds.
- (b) Compare oxygen consumption in the two birds.
- (c) Judging from the data in the table, which bird is the long-distance flyer? Justify your answer. [3 marks]

| 13.         | Which of the follow function? Explain yo  | ing r                    | nutat                                  | ional<br>r. Ra  | chan                     | iges v<br>iem.                         | voul                     | i you                    | pred                     | ict to                   | be t              | he mos                                 | t deleterio<br>3 marks] | us to gene          |  |
|-------------|---|--------------------------|--|-----------------|--------------------------|--|--------------------------|--------------------------|--------------------------|--------------------------|-------------------|--|-------------------------|---------------------|--|
|             | <ol> <li>Insertion of a single nucleotide near the end of the coding sequence.</li> <li>Removal of a single nucleotide near the beginning of the coding sequence.</li> <li>Deletion of three consecutive nucleotides in the middle of the coding sequence.</li> <li>Deletion of four consecutive nucleotides in the middle of the coding sequence.</li> <li>Substitution of one nucleotide for another in the middle of the coding sequence.</li> </ol> |                          |  |                 |                          |  |                          |                          |                          |                          |                   |  |                         |                     |  |
| 64.         | 14. What are the three classes of cell-surface receptors? Discuss different ways he adapted to an extracellular signal molecule.  |                          |  |                 |                          |  |                          |                          |                          |                          |                   | ow cells b<br>[2 mar                   |                         |                     |  |
| 15.         | 5. DNA polymerization happens in 5' to 3' direction while proof reading happens in 3'to 5' direction. What will be the consequence(s) if the directions are interchanged? [2 marks]   |                          |  |                 |                          |  |                          |                          |                          |                          |                   |  |                         |                     |  |
| <b>16</b> . | Place the following e   | event                    | s in t                                 | heir c          | orre                     | ct sec                                 | quenc                    | e:                       |                          |                          |                   |  | [2 marks]               |                     |  |
|             | Translation RNA processing  | <u> </u>                 | Trans<br>_ Nuc                         | cripti<br>clear | ion<br>expo              | rt —                                   | _ Po                     | lyade                    | nylat                    | ion                      |                   | Саг                                    | oping                   |                     |  |
| 17,         | Wobble base pairing A. AUG a B. AAA a What does wobble h  | nd U                     | UU<br>UU                               |                 |                          | I                                      | C. GC                    | the fo<br>GA ar<br>AG ar | nd GC                    | GC                       | airs              | codons                                 |                         | 2 marks]            |  |
| 18.         | Explain the basic me  | chan                     | ism 1                                  | for re          | pair (                   | of UV                                  | V ind                    | uced                     | pyrin                    | nidin                    | e dir             | ner.                                   | [2                      | 2 marks]            |  |
| (9)         | What is genomic imp   | print                    | ing?                                   |                 |                          |  |                          |                          |                          |                          |                   |  | Ţ                       | 2 marks]            |  |
| 29.         | Why does mitochone  | dria r                   | need 1                                 | to ope          | erate                    | the (                                  | 2 cyc                    | le?                      |                          |                          |                   |  | [                       | 2 marks]            |  |
| 21./        | Although oxygen do<br>when O <sub>2</sub> is present. \   | es no<br>Why             | ot par                                 | ticipa          | ite di                   | rectly                                 | y in t                   | he cit                   | ric a                    | eid c                    | ycle,             | the cyc                                |                         | es only<br>2 marks] |  |
| Codon table |   |                          |  |                 |                          |  |                          |                          |                          |                          |                   |  |                         |                     |  |
|             |   | GCA<br>GCC<br>GCG<br>GCU | AGA<br>AGG<br>CGA<br>CGC<br>CGG<br>CGU | GAC<br>GAU      | AAC                      | UGC                                    | GAA<br>GAG               | CAA<br>CAG               | GGA<br>GGC<br>GGG<br>GGU | CAC                      | AUA<br>AUC<br>AUU | UUA<br>UUG<br>CUA<br>CUC<br>CUG<br>CUU |                         |                     |  |
|             |   | Ala                      | Arg                                    | Asp             | Asn                      | Cys                                    | Glu                      | Gln                      | Gly                      | His                      | lle               | Leu                                    |                         |                     |  |
|             |   | AAA<br>AAG               | AUG                                    | UUC             | CCA<br>CCC<br>CCG<br>CCU | AGC<br>AGU<br>UCA<br>UCC<br>UCG<br>UCU | ACA<br>ACC<br>ACG<br>ACU | Uaa                      | UAC                      | GUA<br>GUC<br>GUG<br>GUU |                   | JAA<br>JAG<br>JGA                      |                         |                     |  |
|             |   | Lys                      | Met                                    | Phe             | Pro                      | Ser                                    | The                      | Tro                      | Tvr                      | Val                      | A CONTRACTOR      | utop                                   |                         |                     |  |