

# BS20001: Spring2022 Midsem exam

Total Marks: 50  
25 X 2=50

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2

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19CS30031

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Which one of the following statements is INCORRECT?  
(2 Points)

- ☐ Leading strand is synthesized from 5' to 3' direction
- ☐ Lagging strand requires more primers than leading strand during replication
- ☒ Lagging strand is synthesized from 3' to 5' direction

- ☐ Okazaki fragments are observed during lagging strand synthesis

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Backbone of DNA and RNA is covalently linked through \_\_\_\_\_ and is \_\_\_\_\_ in nature.  
(2 Points)

- ☐ phosphoester bond, hydrophobic
- ☐ phosphoester, hydrophilic
- ☒ phosphodiester bond, hydrophilic
- ☐ phosphodiester bond, hydrophobic

5

In lac operon, if you remove the lac operator (the repressor binding site) what will be the effect on the metabolic state of the bacteria?  
(2 Points)

- ☐ RNA Polymerase will not be able to bind the promoter
- ☒ Lactose metabolizing enzymes will be produced irrespective of the presence or absence of lactose
- ☐ Lactose will never be metabolized because the enzymes will never be synthesized
- ☐ Glucose metabolism will be hampered

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On the ribosome, the mRNA is read from \_\_\_\_\_, and the polypeptide chain is synthesized from \_\_\_\_\_.  
(2 Points)

- ☐ 3' to 5'; C- to N-terminus
- ☐ 5' to 3'; C- to N-terminus

☐ 3' to 5'; N- to C-terminus

☒ 5' to 3'; N- to C-terminus

7

We have 64 codons, out of those three are stop codons. On the other hand, we have 20 amino acids that constitute a protein. As a result of that one amino acid can be coded by multiple codons. During translation, tRNA brings correct amino acid at the translation site and recognize codon through its anticodon sequence. Based on the above information, which one of the following statements is most plausible?

(2 Points)

☐ Should have 64 different types of tRNA

☐ Should have less than 20 different types of tRNA

☒ Should have more than 20 and less than 64 different types of tRNA

☐ Should have 20 different types of tRNA

8

A genetic analysis of an unknown infectious agent reveals that it contains only nucleotides G, A, T and C, in the proportion 30 %, 35 %, 15 % and 20 %, respectively. Based on this information, this infectious agent is most likely

(2 Points)

☐ Double-stranded DNA virus

☒ Single-stranded DNA virus

☐ Single-stranded RNA virus

☐ Not enough information is provided

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Arrange the following options in increasing order of protein structure hierarchy:

- A: alpha-helix
- B: amino acid sequence
- C: quaternary structure
- D: a protein domain

(2 Points)

- ☐ B, C, A, D
- ☒ B, A, D, C
- ☐ A, B, C, D
- ☐ A, C, B, D

10

100 template DNA molecules are PCR amplified for 25 cycles in a 100  $\mu$ l reaction mixture. How many amplified products will be there in 0.01  $\mu$ l of solution after amplification?

(2 Points)

- ☐ 3355443200
- ☐ 335544320
- ☐ 3355443
- ☒ 335544

11

A 100 amino acid protein has only polar and charged residues. It has no hydrophobic residues. Which of the following can be expected for its structure?

(2 Points)

- ☐ It will be all beta stranded structure.
- ☐ It will be alpha and beta mixed structure.
- ☒ It will not fold properly.
- ☐ It will be all alpha helical structure.

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What is common feature demonstrated during replication and transcription in *E. coli*?

(2 Points)

- ☐ Both follow the semi-discontinuous mode of synthesis
- ☒ Both processes involve making and breaking of hydrogen bonds
- ☐ Both require RNA primers for the synthesis of new strands which should be removed later
- ☐ Both require helicase to separate double stranded DNA into single strands

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In classic Sanger DNA sequencing technique, four types of ddNTPs are used along with the normal dNTPs. Which of the following is the correct combination?

(2 Points)

- ☒ Each tube will have one type of ddNTP and all four types of dNTP
- ☐ Each tube with one type of ddNTP and one type of dNTP (e.g., ddATP + dATP in tube 1, ddGTP + dGTP in tube 2 and so on)
- ☐ Each tube will have one type of dNTP and all four types of ddNTP
- ☐ All four ddNTPs and four dNTPs in same reaction tube

14

Peptide backbone has three torsion angles namely omega, phi and psi. However, Ramachandran map ignores one of these and plots only two. Which of the following statements gives the CORRECT reasoning for this? (2 Points)

- ☐ A two-dimensional plot can be made using only two variables. Hence one of the angles was ignored.
- ☒ One of the torsion angles is part of the peptide plane and in most cases has only one value. Hence, unnecessary to plot.
- ☐ It does not matter which two torsion angles are used to create the Ramachandran map. The same map can be created by choosing any two of the three torsion angles.
- ☐ Historically only two torsion angles were discovered at that time. Hence, Ramachandran and his student used only those two torsion angles for the map.

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The coding region of a gene is 132 nucleotides long, including both start and stop codons. Which of the following would be the most likely effect of a single nucleotide deletion at position 91 in the coding region? (2 Points)

- ☒ There would be changes only after the first 30 amino acids
- ☐ There would be no effect on the polypeptide
- ☐ The entire amino acid sequence of the polypeptide would change
- ☐ There would be change only in the 31st amino acid

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Theoretically, a vast number of different proteins can be assembled from 20 different amino acids. How many polypeptide chains are possible that are 10 amino acids long? (2 Points)

- ☒  $20^{10}$
- ☐  $20^{10} \times 10^{20}$
- ☐  $20 \times 10$
- ☐  $10^{20}$

17

Which of the following pairs of amino acids might contribute to protein conformation by forming electrostatic interactions?

(2 Points)

(Hints: Nonpolar: Glycine, Phenylalanine and Tyrosine; Positively charged: Lysine and Arginine; Negatively charged: Glutamate and Aspartate; Polar: Asparagine, Glutamine, Serine)

- ☐ Lysine and Asparagine
- ☐ Phenylalanine and tyrosine
- ☒ Glutamate and lysine
- ☐ Glycine and aspartate

18

In an alien species, there are only 2 types of nucleotides (instead of 4 types), but codons are 4 nucleotide long. If each type of codon specifies one unique amino acid, how many possible amino acids can be coded. Also consider that they have two stop codons.

(2 Points)

- ☒ 14
- ☐ 64
- ☐ 62
- ☐ 16

One undergrad student is repeating Anfinsen's experiment with an enzyme that has TEN cysteine residues and forms FIVE disulfide bonds. The enzyme is denatured by urea and reduced by BME. What is the expected activity of the enzyme if it is oxidized first and then urea is removed?

(2 Points)

- ☐ 0.5%
- ☐ 2%
- ☒ 0.1%
- ☐ 1%

In gel electrophoresis different sized DNA migrate at different rate. Which of the following statements is FALSE?

(2 Points)

- ☐ DNA is visualized in the gel by staining with ethidium bromide, which fluoresces under UV light
- ☐ DNA molecules get separated according to their size
- ☐ DNA is negatively charged and hence migrates towards the positive terminal in the applied electric field gradient
- ☒ Smaller DNA migrates slowly than larger DNA

Which of the following is in correct order:

(2 Points)

- ☐ Transcription, Transport of mRNA from nucleus to cytoplasm, Poly-adenylation, mRNA Splicing, Translation, Protein folding
- ☐ Transcription, 5'capping of mRNA, Transport of mRNA from nucleus to cytoplasm, mRNA Splicing, Translation, Protein folding



- ☒ Transcription, 5'capping of mRNA, mRNA Splicing, Transport of mRNA from nucleus to cytoplasm, Translation, Protein folding
- ☐ Transcription, mRNA splicing, Poly-adenylation, translation, Transport of mRNA from nucleus to cytoplasm, Protein folding

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Estimate the length of the protein coded by the following DNA sequence. Start and stop codons are in bold letters; introns are underlined.

5'CACAT**ATG**GCGATACGAAGGGGACGCATGGCGGACAGGGCCGTTGC  
**TAAGGTTGTG** 3'

(2 Points)

- ☐ 13
- ☐ 14
- ☐ 11
- ☒ 10

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A new strain of bacteria was isolated from a natural hot water geyser. Comment on the expected DNA base compositions in this thermophilic organism (living in high temperatures).

(2 Points)

- ☐ Insufficient data
- ☐ A+T > G+C
- ☐ Equal A+T and G+C composition
- ☒ G+C > A+T

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Choose the correct one from the following options that indicate the number of molecules present per *E. coli* cell in ascending order  
(2 Points)

- ☐ DNA<mRNA<protein<rRNA
- ☐ DNA<tRNA<mRNA<protein
- ☒ DNA<mRNA<tRNA<protein
- ☐ DNA<protein<mRNA<tRNA

25

The rate of protein synthesis in prokaryote is limited by the rate of mRNA synthesis. If mRNA synthesis occurs at the rate of 51 nucleotides/sec, then the rate of protein synthesis occurs at:  
(2 Points)

- ☐ 50 amino acids/sec
- ☐ 25 amino acids/sec
- ☒ 17 amino acids/sec
- ☐ 12 amino acids/sec

26

12 amino acids stretch of a protein forms an alpha-helix. What could be the expected phi and psi angles of the amino acids participating in forming alpha-helix?  
(2 Points)

- ☐ +60, -50
- ☒ -60, -50

☐ -60, +50

☐ +60, +50

27

What will be the transcript (i.e. RNA) of this gene (coding strand is underlined)?

(2 Points)

5'	G	C	T	C	A	G	C	<u>A</u>	<u>T</u>	G	G	G	G	G	C	G	T
3'	C	G	A	G	T	C	G	T	A	C	C	C	C	C	G	C	A

☐ 3' AUGGGGGCGUAA5'

☐ 5' UACCCCCCGCAUU3'

☐ 3' UACCCCCCGCAUU5'

☒ 5' AUGGGGGCGUAA3'

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