| NEET : CHAPTER WISE TEST- 5 |   |        |   |  |  |
|-----------------------------|---|--------|---|--|--|
| SUBJECT :- BIOLOGY          |   |        | DATE  |  |  |
| CLASS :- 12 <sup>th</sup>   |   |        | NAME  |  |  |
| CHAP                        | PTER :- MOLECULAR BASIS OF INHERITANCE  |        | SECTION   |  |  |
|                             | (SECT   | ION-A) |   |  |  |
| 1.                          | Find the incorrect match.<br>(A) $\varphi \times 174$ phage – 5383 deoxyribose<br>sugar in its genetic material<br>(B) $\lambda$ - phage-2 × 48502 nucleotides<br>(C) E. coli-4.6 × 10 <sup>6</sup> base pairs<br>(D) Human-3.3 x 10 <sup>9</sup> nucleotides in its  |        | <ul> <li>(A) Structure (a) is H<sub>1</sub> protein, which is rich in arginine and aspartic acid.</li> <li>(B) Structure (b) is the histone core with two molecules each of four histone proteins.</li> <li>(C) Structure (c) is single-stranded DNA</li> </ul>   |  |  |
| 2.                          | Genome         Select the incorrectly matched pair.         (A) Purine nucleosides       1'-9 glycosidic linkage         (B) 3D model of B-DNA       Watson and Crick         (C) C-valve of B-DNA       Total amount of DNA in diploid cell         (D) Adenine and guanine       Nine-membered double-ring structures | 8.     | <ul> <li>(D) Structure (b) is called Nu-Body, which has five types of histone proteins.</li> <li>Euchromatin</li> <li>(A) Is transcriptionally active</li> <li>(B) Stains lightly</li> <li>(C) Is loosely packed</li> </ul>   |  |  |
| 3.                          | The force that holds DNA together in a<br>double helix is<br>(A) Hydrogen bonds   | 9.     | (D) All of the above<br>The unequivocal proof that DNA is the   |  |  |
| 4.                          | (B) The force of the twist<br>(C) N-glycosidic linkage<br>(D) lonic interactions<br>How many N-glycosidic linkage are<br>present in $\varphi$ 174 bacteriophage?<br>(A) 48502 (B) 5386<br>(C) 10772 (D) 4.6 * 10 <sup>6</sup>   |        | <ul> <li>genetic material came from the experiment of</li></ul>   |  |  |
| 5.                          | In a DNA molecule, two polynucleotide<br>chains are different in<br>(A) Number of purine and pyrimidine<br>bases and pyrimic<br>(B) Polarity<br>(C) Molecular weight<br>(D) All of the above  | 10.    | <ul> <li>RNA is not a better genetic material because</li> <li>(A) It is chemically and structurally unstable</li> <li>(B) It is able to generate its replica</li> <li>(C) 2'-OH group present at every nucleotide in RNA as an additional moiety, which makes RNA labile and easily degradable.</li> </ul> |  |  |
| 6.                          | <ul> <li>What is incorrect for DNA?</li> <li>(A) The pitch of the helix is 3.4 nm.</li> <li>(B) Phosphate moiety is at 5' end of deoxyribose sugar,</li> <li>(C) Diameter of DNA is 2 nm.</li> <li>(D) The charge on the DNA is positive.</li> </ul>  | 11.    | <ul> <li>(D) All except (B)</li> <li>Select the incorrect statement.</li> <li>(A) RNA was the first genetic material.</li> <li>(B) Some essential life processes like metabolism, translation, and splicing evolved around DNA.</li> </ul>  |  |  |
| 7.                          | Refer to the given figure of nucleosome,<br>and select the correct option:  |        | <ul><li>(C) DNA has evolved from RNA with chemical modification.</li><li>(D) RNA acts as genetic material as well as catalyst.</li></ul>  |  |  |
|                             | Core of histone molecules   | 12.    | RNA is not genetic material in<br>(A) TMV<br>(B) QB bacteriophage<br>(C) HIV<br>(D) φ × 174   |  |  |

| 13. | How many amino acids make up a protein?<br>(A) 10 (B) 20 (C) 30 (D) 50  | 21. | replicate twice in culture medium containing N <sup>15</sup> , the percentage of hybrid |
|-----|---|-----|---|
| 14. | Which of the chemical groups makes RNA labile and easily degradable?  |     | DNA will be<br>(A) 25% (B) 75%  |
|     | (A) 3'-OH (B) 5'-PO <sub>4</sub><br>(C) 2'-OH (D) 3'-PO <sub>4</sub>  |     | (C) 50% (D) 17.5%   |
| 15. | Teminism is concerned with  | 22. | lsotopes used to prove semiconservative   |
|     | (A) Reverse transcriptase   |     | mode of DNA replication of DNA here   |
|     | (B) DNA-dependent RNA polymerase  |     | (A) $N^{14}$ , $C^{14}$ (B) $S^{35}$ , $p^{32}$   |
|     | (C) RNA-dependent DNA polymerase<br>(D) Both (A) and (C)  |     | (C) $N^{14}$ , $N^{15}$ (D) $N^{14}$ , $p^{32}$   |
| 16. | How many phosphodiester bonds are<br>present in the following DNA sequence?                                       | 23. | Discontinuously synthesized fragments of  |
|     | 5'ATCGATG 3' 3'TAGČTAC'   |     | DNA are joined by   |
|     | (A) 12 (B) 10 (C) 14 (D) 16   |     | (A) RNA ligase (B) Helicase   |
| 17. | DNA replication is<br>(A) Polymerization of nucleotides by DNA-<br>dependent DNA polymerase activity              |     | (C) DNA polymerase II (D) DNA ligase  |
|     | (B) Energetically very expensive process<br>(C) Polymerization of ribonucleotides in $5' \rightarrow 3$ direction | 24. | Which of the following is not the<br>component of transcription unit?                   |
|     | (A) Only (A) is correct.  |     | (A) Promoter (B) Terminator   |
|     | (B) Only (A) and (B) are cor <mark>rect.</mark>   |     | (C) Structural gene (D) Inducer   |
|     | (C) Only (A) and (C) are correct.<br>(D) (A) (B) and (C) are correct.   | 25  | The functional unit of gene that specifies  |
|     | (D) $(A)$ , $(D)$ , and $(C)$ are correct.  |     | synthesis of one polypeptide is   |
| 18. | DNA replication is  |     | (A) Muton (B) Cistron   |
|     | (A) Semiconservative  |     | (C) Intron (D) Transposon   |
|     | (C) Bidirectional   |     |   |
|     | (D) All of the above  | 26. | A fully processed hnRNA is called mature  |
| 10  | A failure in cell division offer DNA  |     | (A) Methyl quanidine trinhosphate at 5  |
| 13. | replication results into  |     | end   |
|     | (Å) Gene mutation   |     | (B) Tail of adenylate residue   |
|     | (B) Chromosomal aberration  |     | (C) Introns   |
|     | (D) All except (C)  |     | (D) More than one option is correct   |
| 20. | Read the given statements with respect to   | 27. | Read the following statements with  |
|     | DNA replication.  |     | respect to transcription.   |
|     | catalyzes the polymerization in only one  |     | (A) There is a single RNA polymerase  |
|     | direction that is $5' \rightarrow 3'$ .   |     | enzyme that catalyzes transcription of all  |
|     | B. In eukaryotes, the replication of DNA  |     | (B) Termination is controlled by a factor   |
|     | C The discontinuously synthesized   |     | (C) In eukaryotes, the polycistronic genes  |
|     | fragments are later joined by the enzyme  |     | have interrupted coding sequences.  |
|     | ligase.   |     | (D) Additional regulatory sequence may be   |
|     | D. Deoxyridonucleoside triphosphate serves dual purpose in addition to acting                                     |     | present further upstream or downstream to   |
|     | as substrates, they provide energy for  |     | the promotor.   |
|     | polymerization reaction during DNA  |     | Correct statement Is/are  |
|     | replication.  |     | (A) Only (A) and (D) $(B)$ Only (A) and (D)   |
|     | (A) Two (B) One   |     | (C) Only (A) and (B)  |
|     | (C) Four (D) Three  |     | (D) (A), (B), (C), and (D)  |
|     |   | 1   |   |

| 28. | If coding strand of DNA has the nitrogenous         |     | (SECTION-B)   |
|-----|---|-----|---|
|     | base sequence as 5'ATGCTA3'. What would             | 36. | Which of the following option is not correct                  |
|     | be the mRNA strand sequence?                        |     | with respect to soluble RNA?                                  |
|     | (A) 3'AUGCUAG3' (B) 5'AUGCUAG 3'                    |     | (A) It has structural and catalytic roles in                  |
|     | (C) 5'UACGAUC3' (D) 3'UACGAUC5'                     |     | protein synthesis.  |
|     |   |     | (B) It constitutes 15% of total cellular RNA.                 |
| 29. | In prokarvotes, the initiation and                  |     | (C) It possesses three unpaired nucleotides                   |
|     | termination factors of transcription are            |     | (D) It is soluble in 1 mole of NaCl solution                  |
|     | respectively  |     |   |
|     | $(A) \sigma$ and $\sigma$ (B) $\sigma$ and $\sigma$ | 37. | The tRNA is associated with all the                           |
|     | (A) $\theta$ and $\beta$ (B) $\beta$ and $\theta$   |     | features except   |
|     | (C) $\sigma$ and 15 (D) $\rho$ and y                |     | (A) It has clover leaf 2D structure                           |
|     |   |     | (B) There are no tRNAs for stop codons                        |
| 30. | Post-transcriptional modification of primary        |     | (C) At its 5' end, unpaired CCA sequence                      |
|     | transcript in eukaryotes does not involve           |     | IS present<br>(D) It has an anticodon loop that reaches       |
|     | (A) Addition of 7-methyl guanosine                  |     | the genetic code  |
|     | triphosphate at 5" end of hnRNA                     |     |   |
|     | (B) Tail ring at 3' end                             | 38. | Select the incorrectly matched pair.                          |
|     | (C) Removal of introns and joining of               |     | (A) Soluble RNA-tRNA  |
|     | exons in a coordinated manner                       |     | (B) Shine-Dalgarno sequence-mRNA                              |
|     | (D) Removal of RNA primers from hnRNA               |     | (C) Smallest RNA-TRNA<br>(D) Ribozyme-23S rRNA in prokaryotes |
|     |   |     | (D) Ribbzyme-256 RivA in provaryotes                          |
| 31. | The role of sigma factor in bacterial RNA           | 39. | Which of the following techniques is used                     |
|     | polymerase is                                       |     | to determine the protein structures?                          |
|     | (A) To catalyze RNA syntheses                       |     | (A) X-ray crystallography                                     |
|     | (B) to terminate transcription process              |     | (B) Kryptonics X-ray vision                                   |
|     | (C) To unwind the DNA template                      |     | (C) Magnetic resonance imaging (MRI)                          |
|     | (D) To bind RNA polymerase enzyme                   |     | (D) None of the above   |
|     | correctly on the promotor region                    |     |   |
| ~~  |   | 40. | Movement of ribosome on mRNA is                               |
| 32. | If the genetic code is tetraplet, then what is      |     | called  |
|     | the possible number of codons which code            |     | (A) Translocation (B) Rotation                                |
|     | 20 amino acids?                                     |     | (C) Transfection (D) Translation                              |
|     | (A) 261 (B) 64                                      |     |   |
|     | (C) 256 (D) 43                                      | 41. | Which of the following is a precursor of                      |
| 22  |   |     | amino acid synthesis?   |
| 33. | (A) Derwined for efficient translation              |     | (A) fatty acids   |
|     | (A) Required for efficient translation              |     | (B) $\alpha$ -ketoglutaric acid                               |
|     | (D) Dresent at 5' and before start addar            |     | (C) mineral salts   |
|     | (B) Present at 5 end before start codori            |     | (D) volatile acids  |
|     | (D) All of the above                                | 40  | Which of the following serves provides                        |
|     |   | 42. | attachment site for RNA polymerase                            |
| 24  | Degeneration of a gapatic adda is attributed        |     | enzyme?   |
| 34. | begeneration of a genetic code is attributed        |     | (A) Promoter gene (B) i gene                                  |
|     | (A) First pusicatida of a cadar                     |     | (C) Operator gene (D) Regulator gene                          |
|     | (R) Second nucleotide of a codon                    |     |   |
|     | (C) Third nucleotide of a codon                     | 43. | Which of the following is not correct with                    |
|     | (D) Entire codon                                    |     | respect to lac operon?  |
|     |   |     | (A) It is inducible operon system.                            |
| 35. | Which of these proves the degeneracy of             |     | (B) It exactly negative control through                       |
|     | genetic code?                                       |     | repressor protein.  |
|     | (A) Substitution                                    |     | (C) Regulator gene functions all the time                     |
|     | (B) Transition of nucleotides                       |     | constitutively.   |
|     | (C) Wobble hypothesis                               |     | (D) The repressor proteins bind to the                        |
|     | (D) Frameshift mutation                             |     | promoter region.  |
|     | · · · · · · · · · · · · · · · · · · ·               |     | PC #3   |

| 44. | <ul> <li>Read the following statements with respect to lac operon, and select the correct option.</li> <li>A. It has three structural genes.</li> <li>B. Glucose can act as inducer.</li> <li>C. Repressor protein is acting in presence of lactose.</li> <li>D. Repressor is synthesized by regulator gene.</li> <li>(A) Only (B) is incorrect.</li> <li>(B) (A) and (D) are correct.</li> <li>(C) (A) and (D) correct.</li> <li>(D) (A), (B), and (D) are correct.</li> </ul> | 48. | <ul> <li>48. Study the following statements with respect to DNA fingerprinting.</li> <li>A. The process of DNA fingerprinting involves specific sequence of DNA known as repetitive DNA.</li> <li>B. VNTR belongs to a class satellite DNA referred to as microsatellite.</li> <li>C. In addition to the application in forensi science, it has much wider application such as in determining population and genetic diversities.</li> </ul> |
|-----|---|-----|--|
| 45. | β-Galactosidase is synthesized by E. coli<br>to catalyze hydrolysis ofA<br>into B and<br>C<br>(A) A-Lactose; B-Glucose; C-Galactose<br>(B) A-Lactose; B-Glucose; C-Fructose<br>(C) A-Galactose; B-Glucose; C-Fructose   | 49. | <ul> <li>(A) Only (B) and (C) are correct.</li> <li>(B) Only (B) is incorrect.</li> <li>(C) Only (A) is correct.</li> <li>(D) All are incorrect.</li> <li>Select the most commonly used vector in the Human Genome Project.</li> </ul>   |
| 46. | All the non-human genome models were<br>sequenced at the times of Human<br>Genome Project except<br>(A) Drosophila (B) Penicillium  |     | <ul><li>(A) Bacteriophage</li><li>(B) Bacteriophages</li><li>(C) Plasmids only</li><li>(D) BAC and YAC</li></ul>   |
| 47. | <ul> <li>(C) Arabidopsis</li> <li>(D) Caenomabditis</li> <li>The technique of DNA fingerprinting was initially developed by</li> <li>(A) Alec Jeffreys</li> <li>(B) Hargovind Das Khurana</li> <li>(C) Miescher</li> <li>(D) F.Crick</li> </ul>   | 50. | Phospholipids are<br>(A) monoglycerides<br>(B) diglycerides<br>(C) triglycerides<br>(D) Any of the above   |