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**PRESIDENCY UNIVERSITY  
BENGALURU**

**SCHOOL OF ENGINEERING  
MID TERM EXAMINATION - OCT 2023**

**Semester :** Semester V - 2021

**Course Code :** CSE3069

**Course Name :** Sem V - CSE3069 - Introduction to Bioinformatics

**Program :** B. TECH

**Date :** 2-NOV-2023

**Time :** 9:30AM - 11:00AM

**Max Marks :** 50

**Weightage :** 25%

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**Instructions:**

- (i) Read all questions carefully and answer accordingly.
  - (ii) Question paper consists of 3 parts.
  - (iii) Scientific and non-programmable calculator are permitted.
  - (iv) Do not write any information on the question paper other than Roll Number.
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**PART A**

**ANSWER ALL THE QUESTIONS**

**(10 X 1 = 10M)**

1. Proteomics refers to the study of:
  - a. Set of proteins in a specific region of the cell
  - b. Biomolecules
  - c. Set of proteins
  - d. The entire set of expressed proteins in the cell

(CO1) [Knowledge]
2. Which tool is best suited for finding Locus Specific Mutations?
  - a. Sequence Retrieval Systems
  - b. Entrez
  - c. Sequence Tagged Site
  - d. Expressed Sequence Tags

(CO1) [Knowledge]
3. The identification of drugs through the genomic study is called\_\_\_\_\_.
  - a. Genomics
  - b. Pharmacogenomics
  - c. Pharmacogenetics
  - d. Metabolomics

(CO1) [Knowledge]
4. Analysing or comparing the entire genome of species is called? (CO1, Knowledge)
  - a. Bioinformatics
  - b. Genomics
  - c. Proteomics
  - d. Genetics

(CO1) [Knowledge]

5. DNA and RNA are examples of which FAMILY of BIOLOGICAL MOLECULES?  
 a. Proteins  
 b. Carbohydrates  
 c. Lipids  
 d. Nucleic Acid  
 (CO1) [Knowledge]
6. FASTA format starts with \_\_\_\_\_ symbol. (CO2, Knowledge)  
 a. / (Backslash)  
 b. \* (Asteric)  
 c. @ (At Symbol)  
 d. > (Greater than symbol)  
 (CO2) [Knowledge]
7. Identification of genes and their respective functions are called \_\_\_\_\_  
 a. Functional Genomics  
 b. Structural Genomics  
 c. Comparative Genomics  
 d. None of the above  
 (CO2) [Knowledge]
8. Entrez, a life science search engine used to search across databases is maintained by \_\_\_\_\_.  
 a. Swiss prot  
 b. EMBL  
 c. DDBJ  
 d. NCBI  
 (CO2) [Knowledge]
9. Which of the following is a primary function of carbohydrates?  
 a. Storing genetic information  
 b. Providing short-term energy storage  
 c. Acting as a long-term energy source  
 d. Speeding up chemical reactions  
 (CO2) [Knowledge]
10. PIR was established by \_\_\_\_\_.  
 a. NBRF  
 b. NCBI  
 c. SIB  
 d. DDBJ  
 (CO2) [Knowledge]

## PART B

**ANSWER ALL THE QUESTIONS**

**(2 X 10 = 20M)**

11. A. Bioinformatics has variety of application fields. Explain any 5 fields of application in detail. (5Marks)  
 B. Ankit wants to study the types of databases associated with bioinformatics. Explain the types with examples. (5 Marks)  
 (CO1) [Comprehension]
12. Describe the following in detail  
 (i) Global and Local Alignments with examples. (5 Marks)  
 (ii) FASTQ & FASTA with examples. (5 Marks)  
 (CO2) [Comprehension]

**PART C**

**ANSWER THE FOLLOWING QUESTION**

**(1 X 20 = 20M)**

13. A. Reconstruct the sequence using the below given k-mers. (10 Marks)

'CATC', 'CAAG', 'CCAA', 'TCCA', 'AGGT', 'CGAT', 'GTAC', 'TACG', 'ACGA', 'CGAT', 'GATC', 'ATCG', 'ATCG', 'TCGA', 'GATC', 'AAGG', 'TCGG', 'CGGA', 'GGAT', 'GGTA', 'GATC', 'ATCA', 'TCAG', 'CAGC', 'AGCA', 'GCAT', 'TCCA', 'ATCC'

B. Calculate the alignment Score for the given Sequences S1 & S2 using the below Substitution matrix by ignoring the gap penalties. (10 Marks)

| Substitution Matrix |    |    |    |   |
|---------------------|----|----|----|---|
|                     | T  | A  | C  | G |
| T                   | 4  |    |    |   |
| A                   | -2 | 3  |    |   |
| C                   | -3 | -2 | 4  |   |
| G                   | -1 | -4 | -1 | 5 |

**S1 = CGTAG-ATAG-TGCTAG-AGAAT-GGG-CCACT**

**S2 = GTAGCTGATG-ATCGATCGTACGTAGC-GCTGA**

(CO2) [Application]