

G-4/404(B)/21

Roll No.....

## M.Sc. IV Semester Examination, 2021

BIOCHEMISTRY

### Paper IV

(Bioinformatics)

Time : 3 Hours ]

[Maximum Marks : 80

**Note :** All questions are compulsory. Question Paper comprises of 3 sections. **Section A** is objective type/multiple choice questions with no internal choice. **Section B** is short answer type with internal choice. **Section C** is long answer type with internal choice.

### SECTION 'A'

#### (Objective Type Questions)

Choose the correct answer :

8 × 1 = 8

1. Who coined the term Bioinformatics and when ?

- (a) Paulien Hogeweg, 1979
- (b) Dr. Margaret Oakley Dayhoff, 1976
- (c) Robert Ledley, 1978
- (d) David W. Mount, 1977

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2. Which one of the following is not a primary nucleic acid database ?

- (a) Gene Bank
- (b) DDBJ
- (c) EMBL
- (d) TREMBL

3. Which one of the following is a primary protein database ?

- (a) SWISS PROT
- (b) EMBL
- (c) DDBJ
- (d) NCBI

4. .... is a secondary database.

- (a) DDBJ
- (b) PROSITE
- (c) NRDB
- (d) OWL

5. .... is a composite database.

- (a) PROSITE
- (b) DDBJ
- (c) NRDB
- (d) EMBL

6. .... is a primary protein structure database.

- (a) PDB
- (b) PubChem
- (c) ChemBank
- (d) SCOP

7. Which one of the following is a secondary protein structure database ?

- (a) PubChem
- (b) PDB
- (c) ChemBank
- (d) SCOP

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\ 8. FASTA format starts with .....symbol.

- (a) /                      (b) \*  
(c) >                      (d) #

**SECTION 'B'**                      **4 × 6 = 24**  
**(Short Answer Type Questions)**

**Note :** Answer the following questions in 250 words.

**Unit I**

1. How knowledge of computer and specific software be helpful to store molecular data and its better use ?

Or

How can you generate large scale data in molecular biology with the use of gel electrophoresis.

**Unit II**

2. What are data base ?

Or

What is data ?

**Unit III**

3. What are meta data ?

Or

What is Gene bank ?

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**Unit IV**

4. How gene expression data are helpful to scientists and researchers ?

Or

What would be the data related to prokaryotic transcription you need to generate and store for further study ?

**SECTION 'C'**                      **12 × 4 = 48**  
**(Long Answer Type Questions)**

**Note :** Explain in detail with suitable flow chart/diagram.

**Unit I**

1. Write major steps to use BLAST.

Or

Write major steps to use FASTA.

**Unit II**

2. Write applications of bioinformatics.

Or

Explain role and use of Pubmed.

**Unit III**

3. Explain Gene Bank.

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Or

Write notes on search strategies in the headings of Indices, Boolean, fuzzy and neighbouring search.

**Unit IV**

4. What will you consider for data related to prokaryotic transcription ? Give lists of such data for prokaryotic transcription.

Or

What will you consider for data related to regulation of prokaryotic gene expression ? Give lists of such data for regulation of prokaryotic gene expression.

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