

2203000206026006
EXAMINATION FEBRUARY-MARCH 2024
BACHELOR OF SCIENCE (SIXTH SEMESTER)
MICROBIOLOGY-XXII (MB-606-GENOMICS PROTEOMICS
AND BIOINFORMATICS) – LEVEL 2

[Time: As Per Schedule]

[Max. Marks: 50]

Instructions:

1. Fill up strictly the following details on your answer book

- a. Name of the Examination : **BACHELOR OF SCIENCE (SIXTH SEMESTER)**
 - b. Name of the Subject : **MICROBIOLOGY-XXII (MB-606-GENOMICS PROTEOMICS AND BIOINFORMATICS)-LEVEL 2**
 - c. Subject Code No : **2203000206026006**
2. Sketch neat and labelled diagram wherever necessary.
 3. Figures to the right indicate full marks of the question.
 4. All questions are compulsory.

Seat No:

--	--	--	--	--	--

Student's Signature

Q.1 Give answers in brief

8

- a. Give full form of: i) NGS ii) ORF
- b. State the difference between dNTPs & ddNTPs.
- c. Define genome annotation
- d. Name the three data retrieval systems.
- e. What is metabolomics
- f. Give full form of i) NCBI ii) INSD
- g. State various forms of BLAST.
- h. Give two differences between global Vs local alignment.

Q.2 Explain/Comment on ANY TWO of the following: - 14

- a. Explain how DNA is sequenced by chain termination method.
- b. Functional genomics links genes to phenotypes.
- c. Biological databases can be classified into various categories.

Q.3 Discuss ANY TWO of the following: - 14

- a. Write a detailed note on present bioinformatics scenario in India
- b. Discuss in detail Multiple sequence alignment and the calculation of score for similarity search.
- c. Discuss how two-dimensional gel electrophoresis allow visualization of many more cellular proteins.

Q.4 Write short note on ANY TWO of the following: - 14

- a. Multiple Displacement Amplification Method.
- b. Systems Biology
- c. Compare FASTA & BLAST
