

**RAN-2103001106020002**

**B.Sc. (Biotechnology) (Sem. VI) Examination March - 2026**  
**BT-18 : Introduction to Bioinformatics (Old)**

**Time: 2 Hours ]****[ Total Marks: 50**

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| <p><b>સૂચના : / Instructions</b></p> <p>(1) ઉપરોક્ત દર્શાવેલ વિગતો ઉત્તરવહી પર અવશ્ય લખવી.<br/>Fill up strictly the above details on your answer book.</p> <p>(2) Figures to the right indicate full marks.</p> <p>(3) Draw neat and labeled diagrams wherever necessary.</p> | <p>Seat No.:</p> <table border="1" style="width: 100%; height: 20px; border-collapse: collapse;"> <tr> <td style="width: 15%;"></td> <td style="width: 15%;"></td> <td style="width: 15%;"></td> <td style="width: 15%;"></td> <td style="width: 15%;"></td> <td style="width: 15%;"></td> </tr> </table> <div style="border: 1px solid black; height: 80px; margin-top: 10px; position: relative;"> <div style="position: absolute; bottom: 5px; right: 5px; font-size: small;">Student's Signature</div> </div> |  |  |  |  |  |  |
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**Q. 1      Give Specific Answer (Any Four):** **(08)**

- A. State the objectives of bioinformatics.
- B. Mention any two databases related to metabolic pathways.
- C. Define the term "Topology" as used in the CATH classification system.
- D. What is meant by an accession number in biological databases?
- E. Explain the term "percentage identity" in BLAST output.
- F. What do you understand by iterative alignment?

**Q. 2      Attempt Any Two:** **(14)**

- A. Elaborate on the scope and significance of bioinformatics.
- B. Describe protein structure prediction servers available in India.
- C. Discuss protein sequence databases in detail.

**Q. 3      Explain in detail Any Two:** **(14)**

- A. Explain the role and features of Protein Data Bank (PDB) as a protein structure database.
- B. Describe the Smith-Waterman algorithm used for pairwise sequence alignment.
- C. Discuss the different types of gap penalties used in sequence alignment.

**Q. 4      Write Short note (Any Two):** **(14)**

- A. Explain the working and applications of BLAST.
- B. Compare global and local sequence alignment methods.
- C. Describe the Dot Plot technique for sequence comparison.