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**Department of Biotechnology**

**M.Sc. 4th Semester, 2023**

**Paper XVIII - Practical Examination**

**Time 7 hrs (10.00 AM to 5.00 PM)**

**Total marks 40**

1. Download the following gene sequence from NCBI  
Accession number: **XM\_003614410.3**
  - a. Using BLAST identify the 5 best homologs. Do the multiple alignments with default parameters.
  - b. Translate this gene sequence to corresponding protein sequence using any freely available bioinformatic software.
  - c. Using appropriate BLAST, identify the 5 best protein homologs. Do the multiple alignments with default parameters.
  - a. Find whether this protein has any crystal structure submitted in the PDB.  
6+2+6+1=15
  
2.
  - a. Design a pair of primers for the amplification of the given gene sequence using any free primer designing software.
  - b. Discuss the important parameters that you should consider while designing the primer pair.  
7+3=10
  
3. Viva voice (10)
  
4. Practical copy (5)