Ahmedabad University

School of Arts and Sciences

BIO213 Bioinformatics

IMSc Programme
Mid Term Examination 2022 (Monsoon Semester)

Total Marks = 30 **Time – 1 hours 30 minutes**

Enrolment Number:..... **SET 2**
Date: 18-09-2022 (Exam date)
No. of printed pages: 1 (Pages of question paper)

Note: For problems 1 and 2, you should present your scoring/traceback matrices as well as the best alignment. For Problem 3 provide a screen shot of each step you performed. The screenshot should be of the whole screen with sufficient resolution read your results.

1) Perform global alignments between sequences ATACTGGG and TGACTGAG and give us the total score of alignment. Here assume the scoring scheme is +1 for match, -1 for mismatch and -2 for an alignment with a gap.

2) Perform local alignments between sequences TCAGTTGCC and AGGTTG and give us the total score of alignment. Here assume the scoring scheme is +1 for match, -2 for mismatch and -2 for an alignment with a gap.

3) Find a novel gene using BLAST suite for the accession number CAG46737.1

A BLAST hit is considered to be a novel gene hit if

(a) If it is not a 100% identity match to a protein in the database from the same species corresponding to the accession number provided to you.
(b) If there is a match with >95% identity to a different species than the one you started with.
(c) Which other BLAST parameters did you check? What were the values of these parameters.
(d) Perform multiple sequence alignment of the bottom five of 500 BLAST hits.