Indraprastha Institute of Information Technology Delhi (IIITD) Department of Computational Biotechnology

BIO213 - Introduction to Quantitative Biology

ASSIGNMENT-2 (April 06, 2023)

Instructions:

- 1. You are required to submit the assignments by next **Saturday**, **15 April 2023**.
- 2. Use Python to write your program.
- 3. Compile everything into a single PDF. Use <roll no.-name> to label and save the file.
- 4. You are also required to submit a well commented working code.
- 5. Requests for extension of submission deadline will NOT be entertained.

Question 1. Write a program to implement Chou and Fasman method of secondary structure prediction (Helix and Beta strand) for the following protein sequence.

SGFRKMAFPSGKVEGCMVQVTCGTTTLNGLWLDDTVYCPRHVICTAEDMLNPNYEDL LIRKSNHSFLVQAGNVQLRVIGHSMQNCLLRLKVDTSNPKTPKYKFVRIQPGQTFSVLA CYNGSPSGVYQCAMRPNHTIKGSFLNGSCGSVGF

a) Provide a well commented code for the same.

(30 marks)

b) Display the output using the notation – H: Helix, S: Beta strand.

(10 marks)

Chou and Fasman parameters to be used for the prediction are as following:

| Residue | Ρα | Residue | Рβ |
|---------|------|---------|------|
| Glu | 1.53 | Met | 1.67 |
| Ala | 1.45 | Val | 1.65 |
| Leu | 1.34 | Ile | 1.60 |
| His | 1.24 | Cys | 1.30 |
| Met | 1.20 | Tyr | 1.29 |
| Gln | 1.17 | Phe | 1.28 |
| Trp | 1.14 | Gln | 1.23 |
| Val | 1.14 | Leu | 1.22 |
| Phe | 1.12 | Thr | 1.20 |
| Lys | 1.07 | Trp | 1.19 |
| Ile | 1.00 | Ala | 0.97 |
| Asp | 0.98 | Arg | 0.90 |
| Thr | 0.82 | Gly | 0.81 |
| Ser | 0.79 | Asp | 0.80 |
| Arg | 0.79 | Lys | 0.74 |
| Cys | 0.77 | Ser | 0.72 |
| Asn | 0.73 | His | 0.71 |
| Tyr | 0.61 | Asn | 0.65 |
| Pro | 0.59 | Pro | 0.62 |
| Gly | 0.53 | Glu | 0.26 |