

Total number of printed pages-4

3 (Sem-5/CBCS) ZOO HE 1

2023

ZOOLOGY

(Honours Elective)

Paper : ZOO-HE-5016

(Computational Biology and Biostatistics)

Full Marks : 60

Time : Three hours

The figures in the margin indicate full marks for the questions.

1. Fill in the blanks : 1×7=7

(a) Gold biotechnology is also known as _____.

(b) DDBJ is a _____ sequence database.

(c) _____ is a computer-annotated protein sequence database.

(d) The information retrieval tool of NCBI GenBank is _____.

(e) Proteomics refers to the study of _____.

Contd.

(f) _____ has been referred to as the mother and father of Bioinformatics.

(g) The identification of drugs through the genomic study is called _____.

2. Answer the following : $2 \times 4 = 8$

(a) Write the differences between local alignment and global alignment.

(b) Write about the different branches of bioinformatics.

(c) What is systems biology ?

(d) Write about the ddNTP structure and its use.

3. Answer the following questions : (*any three*)
 $5 \times 3 = 15$

(a) Describe the different sequence submission tools available at NCBI.

(b) Describe the Sanger's di-deoxy method of DNA sequencing.

(c) Briefly explain essential aspects of primary and secondary database.

(d) Write the differences between PAM and BLOSUM matrices.

- (e) Explain co-efficient of variance and its importance.

Answer the following questions : $10 \times 3 = 30$

4. (a) Describe the scope and applications of bioinformatics in frontier areas of biology.

OR

- (b) What is phylogeny ? Discuss the various methods of phylogenetic analysis.

5. (a) Describe various types of BLAST with their applications. Briefly explain the algorithm of BLAST.

OR

- (b) What is sequence alignment ? Explain the various parameters used for optimum sequence alignment.

6. (a) How can you predict the structure of protein sequence ? Explain in detail one of the knowledge-based method to predict the protein structure.

OR

(b) Calculate the standard deviation and standard error from the following data :

X:	10-20	20-30	30-40	40-50	50-60	60-70	70-80
Y:	4	6	10	18	15	12	5