Registration Number:

Date & session:



ST. JOSEPH'S UNIVERSITY, BENGALURU -27 BDA (BIG DATA ANALYTICS) – III SEMESTER SEMESTER EXAMINATION: OCTOBER 2023 (Examination Conducted in November/December 2023) BDADE 3621 – BIOINFORMATICS (For current batch students only)

Time: 2 Hours This paper contains TWO printed pages and THREE parts Max Marks: 50

PART A

Answer All the Questions 5 X 2 = 10

- What are the functions of the cell?
- 2 What is the central dogma of molecular biology?
- 3 Describe briefly the applications of sequence alignment.
- 4 Name the amplification methods used in Roche and Illumina technologies.
- 5 Name any two tools used for the alignment of NGS sequences.

PART B

Answer any FIVE questions

5 X 4 = 20

- 6 What is PDB and discuss its importance.
- 7 What are the techniques involved in the Human Genome Project?
- 8 What is the role of a Data Analyst in Bioinformatics?
- 9 Write a note on various types of BLAST.
- 10 Differentiate between RefSeq and GenBank databases.
- 11 What is a FASTA format? How do we obtain the gene sequence in FASTA format using NCBI Gene?
- What are the applications of multiple sequence alignment? Write a note on the usage of ClustalOmega.

PART C

Answer Any TWO questions 2 X 10 =20 13.(a) What is reference assembly? Discuss in detail the methodology of NGS technologies. 5 13.(b) Suggest a broad strategy for developing a computational method: There are 100 DNA samples from normal individuals and another 100 from patients with a genetic disorder. How would you compare the two sets of genomic DNA and find any existing difference that is associated with the specific genetic disorder? 5

14.(a)	Define genomics, transcriptomics, and proteomics along with their applications.	6
14.(b)	Write a note on various types of high-throughput data that are generated in these fields.	4
15.(a)	Describe the homology modeling of protein structure. What are its applications?	10