

Register No:

Date:



ST. JOSEPH'S COLLEGE (AUTONOMOUS), BANGALORE

III SEMESTER EXAMINATION, OCTOBER 2019

MSc IN BIG DATA ANALYTICS

BDA 3618: BIOINFORMATICS

TIME 2 1/2 HRS

MAX MARKS 70

This Question Paper Contains One Printed Page And One Part

ANSWER ANY SEVEN QUESTIONS

7 X10 = 70

1. What is genetic information? How it is used by the cells and propagated?
2. Write a note on the importance of sequence analysis. Discuss various features of BLAST tool and list important parameters for choosing the best alignment.
3. Write a note on five eukaryotic cell organelles along with their major functions.
4. What are the various applications of next generation sequencing in research and healthcare?
5. Differentiate between reference and *de-novo* NGS assembly. Write a note on the steps involved in reference alignment along with the type of input required and output generated in each step.
6. Discuss in detail about the applications of multiple sequence alignment along with its importance. Write a note on various features of ClustalOmega tool.
7. Define genomics, transcriptomics, and proteomics. Write a note on various high-throughput data generated in these fields and their applications.
8. Write a detailed note on BLAST algorithm.
9. Discuss the BOWTIE algorithm in detail.