

STELLA MARIS COLLEGE (AUTONOMOUS) CHENNAI 600 086
(For candidates admitted from the academic year 2015 – 2016)

SUBJECT CODE: 15BI/PC/GP34

M. Sc. DEGREE EXAMINATION, NOVEMBER 2016
BIOINFORMATICS
THIRD SEMESTER

COURSE : CORE
PAPER : GENOMICS AND PROTEOMICS
TIME : 3 HOURS

MAX. MARKS: 100

SECTION – A

ANSWER ALL THE QUESTIONS:

(20x1=20)

1. Molecular clocks.
2. Characteristic features of Gene.
3. Positive selection.
4. How do you locate a gene in a genome?
5. Synteny
6. ORF
7. Viral Genomes
8. MZEF
9. SAGE
10. EST
11. KEGG
12. UNIGene
13. 2D-SDS-PAGE
14. Tandem LC
15. Preparative IEF
16. Y2H
17. Protein-Protein Interaction
18. Restriction enzymes
19. Give examples for proteolytic enzymes
20. How do protein undergo modifications?

SECTION – B

ANSWER ANY FOUR QUESTIONS IN 300 WORDS EACH. ALL QUESTIONS CARRY EQUAL MARKS: (Draw Diagrams wherever necessary) (4x10=40)

21. Explain the experimental techniques for locating a gene.
22. Phylogenetic Analysis.
23. Application of microarray in Medicine.
24. Explain in detail the concept of mapping protein modifications.
25. What are the different methods for digesting a protein?
26. Explain the protein – protein Interaction with yeast hybrid model.
27. Comparative Genomics.

SECTION – C

ANSWER ANY TWO QUESTIONS IN 300 WORDS EACH. ALL QUESTIONS CARRY EQUAL MARKS: (Draw Diagrams wherever necessary) (2x20=40)

28. How do you analyze an unknown gene?
29. How do you predict a gene and list the applications of genome analysis?
30. Explain Human Genome Project in detail.
31. Explain Mass Spectrometry.
