# STELLA MARIS COLLEGE (AUTONOMOUS), CHENNAI 600 086 (For candidates admitted during the academic year 2015 – 2016 & thereafter)

**SUBJECT CODE: 15BI/PE/FB34** 

# M.Sc. DEGREE EXAMINATION NOVEMBER 2017 BIOINFORMATICS THIRD SEMESTER

COURSE : ELECTIVE FUNDAMEN

PAPER : FUNDAMENTALS OF BIOINFORMATICS

TIME : 3 HOURS MAX. MARKS: 100

SECTION A (20x1=20)

## Answer ALL questions.

- 1. Entrez
- 2. File format
- 3. BLAST
- 4. Sequence Alignment
- 5. FASTA
- 6. Repeat sequence
- 7. PAUP
- 8. Dendrogram
- 9. UNIGENE
- 10. Primary database
- 11. Microarray
- 12. Introns
- 13. Restriction Endonuclease
- 14. Unrooted phylogenetic tree
- 15. Hybridization
- 16. Bootstrapping
- 17. EST
- 18. E value
- 19. Similarity and Identity
- 20. Conserved Regions

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#### **SECTION-B**

## Answer any FOUR questions in 200 words

(4x10=40)

- 21. Biological Data Formats.
- 22. Phylogenetic Analysis.
- 23. Scoring Matrices.
- 24. Local and Global Alignment.
- 25. Multiple sequence alignment
- 26. Classification of Biological Databases.
- 27. Application of Bioinformatics in various fields.

## **SECTION-C**

## Answer any TWO questions in 1000 words

(2x20=40)

- 28. Explain in detail about the data retrieval systems in Bioinformatics.
- 29. Analysis and Prediction of Gene Regulatory regions.
- 30. Discuss the concept of BLAST sequence alignment tool.
- 31. What are Microarrays? Explain its Types and application.

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