TOTAL PAGE : 1 14BCV21 / 14BCU21

### PSG COLLEGE OF ARTS & SCIENCE

(AUTONOMOUS)

## **BSc DEGREE EXAMINATION MAY 2019**

(Fifth Semester)

#### **Branch - BIOCHEMISTRY**

# **COMPUTATIONAL TECHNIQUES IN BIOINFORMATICS**

Time: Three Hours Maximum: 75 Marks

# **SECTION-A (20 Marks)**

Answer **ALL** questions

**ALL** questions carry **EQUAL** marks  $(10 \times 2 = 20)$ 

- 1 What is HTML?
- What is FTP?
- 3 Mention any two important feature of NCB1.
- 4 Define Bioinformatics.
- 5 What is Sequin?
- 6 What is SRS?
- What is the statistical tool used for sequence alignment?
- What is consensus sequences called? Mention its significance.
- 9 What is RAM PAGE?
- 10 What is RasMol?

# **SECTION - B (25 Marks!**

Answer **ALL** Questions

**ALL** Questions Carry **EQUAL** Marks  $(5 \times 5 = 25)$ 

11 a BriefonXMLandPHYTHON.

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b Write a note on WWW and HTTP.

12 a Write a short note on applications of bioinformatics.

OR

b Give an account on accessing and saving files using PubMed.

13 a Write note on Gen Bank and its structure.

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b Give an account on protein database.

14 a Describe BLAST.

OR

b Give an outline on multiple sequence alignment.

15 a Elaborate on structure-bass drug designing.

OR

b Give a brief note on tool for Ramachandran plot.

## **SECTION - C (30 Marks)**

Answer any **THREE** Questions

**ALL** Questions Carry **EQUAL** Marks  $(3 \times 10 = 30)$ 

- Give an account on methods of connecting to internet.
- 17 Give a detailed account on NCB1.
- 18 Explain Sequence retrieval from NCB1 and Swiss-port.
- 19 Describe the computational approaches to sequence alignment.
- Write a not on structure-based drug designing.

Z-Z-Z END