- 12. (a) Explain similarities and differences between BLAST and FASTA tools for sequence alignment.
 - (b) Layout and analysis of Latin Square Design.
- 13. Write short notes on any four of the following: 2½ each
 - (a) Cumulative frequency curves
 - (b) Coefficient of variation
 - (c) Randomized block design
 - (d) Role of internet in Bio-informatics
 - (c) NCBI

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NP-3332 M.Sc. (Bio-Tech.) Examination, Dec.-2024 Statistical Methods and Bioinformatics

in Biology

(H-103)

M.Sc. (Bio-Tech.)

Time: Three Hours | [Maximum Marks: 50

Note: Attempt questions from all sections as per instructions. Use of calculator is allowed.

Section - A

(Very Short Answer Questions)

Note: Answer all five questions. Each question carries 2 marks. Very short answer is required. 2x5 = 10

P.T.O.

- Explain Histogram and frequency polygon.
- 2. Explain Mean Deviation.
- 3. Explain paired t-test.
- What are the basic principles of experimental design and explain one of them.
- 5. What are the components of Bio-informatics.

Section - B

(Short Answer Questions)

Note: Answer any two questions out of the following three questions. Each question carries 5 marks. Short answer is required not exceeding 200 words.

6. Define Mean, Median and mode with their simple properties.

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- 7. Define the following terms:
 - (a) Type I and Type II errors.
 - (b) Large sample test
- 8. Explain Database accession and searching.

Section - C

(Detailed Answer Questions)

Note: Answer any three questions from this section. Each question carries 10 marks. Answer is required in detail.

3x10=30

- What is dispersion? What are various measures of dispersion? Explain them.
- Discuss the scope and applications of Bio-informatics in molecular biology and computers.
- 11. (a) How will you test the independence of two attributes in 2×2 contingency table.
 - (b) Explain the meaning of "Test for goodness of fit".

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P.T.O.