

JBG-003-1181004

Seat No.

M. Sc. (Sem. I) (CBCS) (W.E.F. 2016) Examination

December - 2019

Zoology: ZOOL - 104

(Biostatistics & Bioinformatics)

Faculty Code: 003

Subject Code: 1181004

Time : $2\frac{1}{2}$ Hours]

[Total Marks: 70

1 Answer the following: (any seven)

- $2 \times 7 = 14$
- (a) Differentiate sample mean and population mean.
- (b) What is cumulative frequency and relative cumulative frequency?
- (c) What is paired and unpaired t- test?
- (d) What is the difference between mathematical and positional averages ?
- (e) What is Partial Correlation Analysis?
- (f) Explain gene prediction.
- (g) What is Biological Database?
- (h) Name software / tools for eukaryotic gene prediction.
- (i) Differentiate analogous and orthologous gene.
- (i) What is GenBank format?
- 2 Answer of the following: (any two)

 $7\times2=14$

- (a) Describe different measures of dispersion with suitable examples.
- (b) What is bivariate Regression analysis? Describe the steps involved it.
- (c) Describe any non-parametric test you have studied with suitable example.

3 Answer the following:

 $7 \times 2 = 14$

- (a) Distinguish between:
 - (a) Null hypothesis and Alternate hypothesis.
 - (b) Type I Error and Type II Error.
- (b) What is "F" ratio? Describe the method of calculating it and write its application in biological science.

OR

3 Answer the following:

 $7 \times 2 = 14$

- (a) What is sequence alignment? Distinguish between pairwise and multiple sequence alignment with significance in molecular biology.
- (b) Discuss applications of bioinformatics in molecular biology and medicine / drug designing.
- 4 Answer the following:

 $7 \times 2 = 14$

- (a) Write notes on Structural modeling of Proteins.
- (b) Define Transcriptomics and their importance.
- 5 Write short notes on any two of the following: $7\times2=14$
 - (a) Describe Chi-Square Test and their importance giving suitable example.
 - (b) What is the difference between 't' test and ANOVA? Explain Giving examples.
 - (c) Write note on BLAST.
 - (d) Write note on Functional Genomics.