| Reg. No | Name | 23P2022 |
|---------|------|---------|
| | | |

M. Sc. DEGREE END SEMESTER EXAMINATION : MARCH 2023 SEMESTER 2 : ZOOLOGY

COURSE: 21P2ZOOT06: GENETICS AND BIOINFORMATICS

| | (For Regular - 2022 Admission and Supplementary - 2021 Admission |) |
|------------------------|--|----------------------------------|
| Duration : Three Hours | | Max. Weights: 30 |
| | PART A | |
| | Answer any 8 questions | Weight: 1 |
| 1. | R-banding is the reverse of G-banding. Comment. | (An, CO 5) |
| 2. | What is Maternal inheritance? | (U) |
| 3. | What is PEV? | (U) |
| 4. | What is Expressivity? | (U) |
| 5. | What is Telomere? | (U) |
| 6. | What is Loss of function mutation? | (U) |
| 7. | Explain Banklt. | (R) |
| 8. | Differentiate between Orthologous and Paralogous sequences | (An, CO 7) |
| 9. | Explain any two phylogenetic analysis programs? | (R, CO 7) |
| 10. | What are DNA Microarrays? | (R) (1 x 8 = 8) |
| | PART B | |
| | Answer any 6 questions | Weights: 2 |
| 11. | Explain briefly on pedigree analysis | (U) |
| 12. | Brief on mini and micro satellites | (U) |
| 13. | Brief on eukaryotic transposable elements | (U) |
| 14. | What is Complementation test? | (U) |
| 15. | What is meant by Annotation? Explain its importance | (U, CO 7) |
| 16. | What are the different methods for sequence alignment? Explain | (R) |
| 17. | Explain the procedure of Phylogeny inference from molecular sequence data. | (U, CO 7) |
| 18. | Explain data mining in proteomics. | (U, CO 8) (2 x 6 = 12) |
| | PART C | |
| | Answer any 2 questions | Weights: 5 |
| 19. | Elaborate on chromatin modifications and their mechanism of action. | (U, CO 5) |
| 20. | Brief on epigenetics of yeast | (U) |
| 21. | Elaborate on the data types and organization of major biological database | es (E) |
| 22. | Explain the methodology of deciphering evolutionary relationships from | (U, CO 7) |
| | molecular sequence data. | (5 x 2 = 10) |
| | | (= 10) |

OBE: Questions to Course Outcome Mapping

| СО | Course Outcome Description | CL | Questions | Total Wt. |
|------|--|----|---------------------|--------------|
| CO 5 | Develop the concepts of Human Genetics, Extra-chromosomal Inheritance, Epigenetics, Quantitative and Population Genetics | U | 1, 19 | 6 |
| CO 7 | Interpret the idea of sequence similarity search and sequence analysis methodology | U | 8, 9, 15, 17, 22 | 11 |
| CO 8 | Analyse the basic ideas of Genomics, Proteomics, systems biology and metabolomics | An | 18 | 2 |

Cognitive Level (CL): Cr - CREATE; E - EVALUATE; An - ANALYZE; A - APPLY; U - UNDERSTAND; R - REMEMBER;