



DCT-003-1141002

Seat No. _____

M. Sc. Botany (Sem. I) (CBCS)

(W.E.F. 2016) Examination

August - 2022

Molecular Biology, Genetics & Evolution : BOT-102

Faculty Code : 003

Subject Code : 1141002

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

[Total Marks : 70

Instruction : Answer the following (any **five**).

1 Answer the following : **7×2=14**

- (a) Define leading and lagging strand.
- (b) What is speciation?
- (c) What is test cross? Write its significance.
- (d) Write Chargaff's rule.
- (e) What is the role of p factor in Transcription termination?
- (f) What is Shine-Dalgarno sequence?
- (g) Give difference between homozygous and heterozygous.

2 Answer the following : **7×2=14**

- (a) Write the functions of aminoacyl synthetase enzyme
- (b) What is frameshift mutation? Give one example
- (c) What is chromosomal aberration?
- (d) Enlist various initiation factors required for bacterial protein synthesis.
- (e) What are modified bases?
- (f) Write the function of topoisomerase and helicase enzyme in DNA replication.
- (g) What is theory of biogenesis?

- 3** Answer the following : **2×7=14**
- (a) Give brief account of C- value paradox.
 - (b) Explain the structure of DNA double helix and its types.
- 4** Answer the following : **2×7=14**
- (a) Explain the process of DNA replication.
 - (b) Write short note : DNA methylation.
- 5** Answer the following : **2×7=14**
- (a) Describe the theories of organic evolution.
 - (b) Describe briefly natural selection.
- 6** Answer the following : **2×7=14**
- (a) Write a note on RNA polymerase.
 - (b) Describe the different properties of genetic code.
- 7** Answer the following : **2×7=14**
- (a) Write in detail the process of transcription.
 - (b) What is dihybrid cross? Explain with suitable example.
- 8** Write the short note of the following : **2×7=14**
- (a) Types of mutation
 - (b) Polyploidy
- 9** Write the short note of the following : **2×7=14**
- (a) Mutagen
 - (b) DNA repair
- 10** Briefly describe : **2×7=14**
- (a) Translation process in prokaryotes.
 - (b) Construction of linkage map.