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K22U 0371

Reg. No. :

Name :

VI Semester B.Sc. Degree (CBCSS – OBE – Regular) Examination, April 2022
(2019 Admission)

CORE COURSE IN BOTANY/PLANT SCIENCE

6B11BOT/PLS : Genetics, Molecular Biology and Plant Breeding

Time : 3 Hours

Max. Marks : 40

Instruction : Draw diagrams wherever specified.

PART – A

Objective type questions. Answer **all**. (4x1=4)

- The function of β subunit of polymerase is
 - Template binding
 - Catalytic binding
 - Promoter binding
 - Cation binding
- Which of the following is not involved in classical plant breeding practices ?
 - Hybridisation of pure lines
 - Artificial selection of pure lines
 - Desirable traits of higher yield
 - Molecular Biology
- Mendel's findings were rediscovered by
 - Correns
 - De Vries
 - Tschermak
 - All
- cDNA is synthesised from RNA by the enzyme
 - DNA polymerase
 - DNA synthetase
 - Helicase
 - Reverse transcriptase

PART – B

Short Essay Questions. Answer **any eight**. (8x2=16)

- How does Mass selection differ from pure line selection ?
- Explain Pleiotropy.
- Discuss Chargaff's rule.
- Briefly explain the plant acclimatization.

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- What is Carcinogenesis ?
- What is the DNA repairing mechanisms occurring in organisms ?
- Comment on chromosome mapping.
- Name the enzymes used in DNA replication and briefly point out their function.
- Describe transposons.
- Write notes on dominance and incomplete dominance.
- What is pedigree analysis ?
- Why Mendel selected *Pisum sativum* as his experimental plant.

PART – C

Essay Questions. Answer **any four**. (4x3=12)

- Explain the Operon concept with diagrammatic representation.
- Give an account of Watson and Crick model of DNA.
- What is Heterosis ? Explain its importance in plant breeding.
- Explain dominant epistasis with example.
- Explain the morphology of chromosome with figure.
- Briefly discuss the sex determination in plants.

PART – D

Long Essay Questions. Answer **any one**. (1x8=8)

- Write an essay on structural aberrations.
- Discuss the human genetic syndromes with examples.
- Explain the post transcriptional modifications in eukaryotic mRNA.