# BOT/V/CC/13

# Student's Copy

# 2022

(CBCS)

## (5th Semester)

### BOTANY

### SEVENTH PAPER

### (Cytogenetics, Plant Breeding and Bioinformatics)

Full Marks: 75

Time : 3 hours

The figures in the margin indicate full marks for the questions

(SECTION : A—OBJECTIVE)

(Marks: 10)

Tick ( $\checkmark$ ) the correct answer in the brackets provided :

**1.** An inversion where the inverted segment does not include the centromere is

- (a) pericentric inversion ( )
- (b) metacentric inversion ( )
- (c) paracentric inversion ( )
- (d) reciprocal translocation ( )
- **2.** When the centromere is situated exactly at one end, the chromosome will be having only one long arm, it is called
  - (a) sub-metacentric ( )
  - (b) telocentric ( )
  - (c) acrocentric ( )
  - (d) metacentric ( )

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[ Contd.

 $1 \times 10 = 10$ 

- **3.** The change in chromosome number which involves one or few chromosome(s) of the genome is called
  - (a) euploidy ( )
  - (b) polyploidy ( )
  - (c) amphiploidy ( )
  - (d) aneuploidy ( )
- **4.** When there is addition of one chromosome in two different pairs  $(2n \ 1 \ 1)$ , it is called
  - (a) simple trisomics ( )
  - (b) secondary trisomics ( )
  - (c) double trisomics ( )
  - (d) tertiary trisomics ()
- **5.** A diagrammatic, graphical representation of relative distances between linked genes of a chromosome, is called
  - (a) ideogram ( )
  - (b) karyotype ( )
  - (c) karyogram ( )
  - (d) linkage map ( )
- **6.** A pattern of inheritance where a single phenotypic trait is governed by more than one pair of genes is
  - (a) extra-nuclear inheritance ( )
  - (b) self-sterility ( )
  - (c) qualitative inheritance ( )
  - (d) quantitative inheritance ( )
- **7.** The three codons which result in the termination of polypeptide chain synthesis are
  - (a) UAA, UAG, GUA
    (b) UAG, UAA, UGA
    (c) UAA, UGA, UUA
    (d) UGU, UAG, UGA
    (e) UGU, UAG, UGA

[ Contd.

- **8.** The process involving inspection, fumigation and growing the introduced plant material in isolation is known as
  - (a) quarantine ( )
  - (b) acclimatization ( )
  - (c) adaptation ( )
  - (d) All of the above ()

9. GenBank is a nucleic acid database maintained by

(a) National Center for Biotechnology Information (NCBI) ( )

)

- (b) DNA Data Bank of Japan (DDBJ) (
- (c) European Molecular Biology Laboratory (EMBL) ( )
- (d) European Bioinformatics Institute (EBI) ( )
- 10. The term 'bioinformatics' was coined by
  - (a) Frederick Sanger ()
  - (b) Paulien Hogeweg ( )
  - (c) Margaret Dayhoff ()
  - (d) J. D. Watson ()

# (SECTION : B-SHORT ANSWER)

(Marks: 15)

Write short notes on the following :

 $3 \times 5 = 15$ 

- UNIT—I
- 1. Functions of chromosome

OR

- 2. Consequences of duplication
- UNIT—II

**3.** Aneuploidy

OR

4. Polyploidy in plant breeding

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[ Contd.

UNIT—III

5. Karyotype

OR

6. Cytoplasmic male sterility

UNIT—IV

7. Physical mutagens

OR

8. Heterosis

UNIT—V

9. BLAST

OR

**10.** Bytes

# (SECTION: C-DESCRIPTIVE)

( Marks : 50 )

Answer the following questions :

Unit—I

 What is cytoskeleton? Elucidate the components and functions of a cytoskeleton. 2+8=10

### OR

- **2.** Write accounts of the following : 5+5=10
  - (a) Deletion and its consequences
  - (b) Chemical composition of chromosome

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[ Contd.

10×5=50

### Unit—II

**3.** What are numerical changes in chromosome? Give an account on sources and consequences of chromosomal anomalies. 2+8=10

#### OR

- 4. Write accounts of the following :
  - (a) Trisomy
  - (b) Segmental allopolyploidy

### UNIT—III

**5.** What is mapping of chromosomes? Describe an account of physical and genetic maps. 2+8=10

#### OR

- **6.** Write short notes on the following :
  - (a) Multiple allelism
  - (b) Kappa particles in Paramecium

#### UNIT—IV

**7.** What is mutation? Describe the types and molecular basis of mutation.

2+8=10

5+5=10

5+5=10

### OR

- **8.** Write short notes on the following : 5+5=10
  - (a) Hybrid vigour
  - (b) Chemical mutagens

### UNIT-V

**9.** What is bioinformatics? Give a basic concept about data and information.

2+8=10

#### OR

- **10.** Write accounts of the following : 5+5=10
  - (a) DNA sequence alignment

(b) Search tools

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