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63/2 (SEM-1) BIT 104

2021

(held in 2022)

**BIOTECHNOLOGY**

(Theory Paper)

Paper Code : BIT-104

(Genetics)

Full Marks – 80

Time – Three hours

The figure in the margin indicate full marks  
for the questions.

1. Answer any *eight* of the following questions :

1×8=8

(a) Split genes are characteristic of

(i) Eukaryotic Cells Genome

(ii) Prokaryotic Cell Genome

(iii) PPLO Genome

(iv) Random Occurrence

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(b) The proteins which are attached with eukaryotic DNA are

- (i) Transferrin and Elastin
- (ii) Protamine and Histone
- (iii) Transferring and Histone
- (iv) Histone and Transferrin

(c) Nucleosome comprise of \_\_\_\_\_ base pair of DNA on it in absence of H1.

- (i) 166 bp
- (ii) 146-147 bp
- (iii)  $200 \pm 40$
- (iv) All of these (in different conditions)

(d) Criss-cross inheritance is generally observed in

- (i) X-Linked Inheritance
- (ii) Y-Linked Inheritance
- (iii) Sex-Linked Inheritance
- (iv) Pseudoautosomal Inheritance

(e) Heteroplasmy refers to presence of

- (i) More than one mtDNA in the same mitochondria or different mitochondria within the same cell
- (ii) Only one mtDNA in the same mitochondria or different mitochondria within the same cell
- (iii) Same type of mtDNA in the same mitochondria
- (iv) More than one chloroplast DNA in the same chloroplast or different chloroplast within the same cell

(f) What is transition ?

(g) The relation between linkage and appearance of new trait is

- (i) Linkage is directly proportional to appearance of new trait
- (ii) Linkage is inversely proportional to appearance of new trait
- (iii) Linkage and appearance of trait is a coincidence
- (iv) None of these

(h) F pili are tubes for conjugation in bacteria and are present in

- (i) F+ (b) F+ and Hfr  
(iii) F- (iv) Hfr

(i) The phenotypic ratio of a dihybrid cross is 9:7 in F<sub>2</sub> generation which is

- (i) Complementary Gene Interaction  
(ii) Supplementary Gene Interaction  
(iii) Epistatic Gene Interaction  
(iv) Pleiotropic Gene Interaction

(j) What are tumour suppressor gene ?

(k) Fingerprinting experiment with 16SrRNA and 23sRNA is also known as

- (i) Chlorotyping (ii) Mitotyping  
(iii) Ribotyping (iv) None of these

2. Write short note on :

2×1=2

- (a) Genetic Drift Or (b) Migration Or  
(c) Bottleneck effect.

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(4)

3. Distinguish between any five : 2×5=10

- (a) Polygenic Trait and Monogenic Trait  
(b) Sterility and Infertility  
(c) rRNA and rDNA  
(d) Population and Community  
(e) F+ and F- Bacterial strain  
(f) Nucleotide Excision Repair and Base Excision Repair  
(g) Organelle Genome and Nuclear Genome.

4. Write short notes on any four : 5×4=20

- (a) Endosymbiont theory  
(b) 16S rRNA typing  
(c) Molecular Taxonomy  
(d) DNA Repair  
(e) Metastasis  
(f) Sex Linked Disorders  
(g) DNA Packaging  
(h) DNA Barcoding  
(i) Comparative Genomics.

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(5)

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5. Answer any *two* of the following questions :

8×2=16

(a) Describe the role of:

2×4=8

(i) Homeotic Gene

(ii) Hox Gene

(iii) Gap Gene

(iv) Pair rule Gene.

(b) Define p and q arm of chromosome. Differentiate chromosome on the basis of centromere. Give a brief nomenclature of human chromosome.

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(c) 'Gene flow and Migration are the wheels of population genetic'. – Justify.

8

(d) Describe the different pathways of apoptosis.

8

6. Answer any *two* of the following questions :

12×2=24

(a) Describe different types of DNA repair.

12

(b) Describe Haldane's Rule with suitable examples.

2+10=12

(c) What are markers ? Describe different types of molecular marker.

2+10=12

(d) What is Hardy-Weinberg law ? Mention the attributes of the law. Mention the importance of Hardy-Weinberg law with special reference to the allelic distribution in a population.

2+2+8=12

(e) Describe different types of non-mendelian inheritance with suitable examples.

12

(f) What is molecular taxonomy ? Mention its role in the study of phylogenetic relationships.

5+7=12

(g) What is mutation ? What are the different types of mutation ? What is the significance of mutation in evolution of characters.

1+8+3=12