

**2016**

**P.G. Diploma Examination in  
Quality Control and Assurance in  
Microbial Technology**

**1st Semester Examination**

**PAPER—QUA-101**

*Full Marks : 50*

*Time : 2 Hours*

*The figures in the right-hand margin indicate full marks.*

*Candidates are required to give their answers in their  
own words as far as practicable.*

*Illustrate the answers wherever necessary.*

**Group—A**

[ Marks : 25 ]

Answer any five questions.

1. Mention the various methods used in the identification of an unknown bacteria? 5
2. How working distance is related to the resolution of light microscope? Justify. 5

*(Turn Over)*

3. In an experiment a researcher found two DNA sample, DNA-1 and DNA-2. The % of Thymine (T) is 10% in DNA-1 and 30% in DNA-2. Establish with proper justification which DNA will have high melting temperature ( $T_m$ ).  
5
4. Write down the various methods used in the enumeration of bacteria.  
5
5. Write down the differences between :
- (a) Viroids and Prions ;
  - (b) Cell wall of gram -ve bacteria with gram +ve bacteria.
- $2 \times 2 \frac{1}{2}$
6. Write a short note on DNA homology experiment. How it can be used in molecular characterization of bacteria?  
3+2
7. State the principles of Gram staining. Why 'negative staining' is so named?  
4+1
8. Define  $T_m$ . State the importance of G + C% in bacterial classification.  
1+4

**Group—B**

[ Marks : 25 ]

Answer any *five* questions.

1. What is Bioinformatics ? Write the different components and field of applications in Bioinformatics. 1+2+2
2. What do you mean by database ? Write in brief about various types of biological databases. 1+4
3. Write the short note (any *two*) of the following :  $2 \times 2 \frac{1}{2}$ 
  - (i) BLAST ;
  - (ii) EMBL ;
  - (iii) NCBI.
4. Write the role of Bioinformatics in drug discovery. 5
5. What is Phylogenetic tree ? Write its significance. 2+3
6. How sequence alignment helpful to study the interrelationship of the species ? What is gap penalty ? 3+2
7. How structure database has played a significance role to predict the unknown protein structure ? 5

8. Write the full form of the following :

5

- (a) ITAA ;
  - (b) HTML ;
  - (c) URL ;
  - (d) ExPASy ;
  - (e) FASTA.
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