

M. SC. (BIOTECHNOLOGY) SEM-II (2012 COURSE)(CHOICE

BASED CREDIT SYSTEM) : WINTER - 2017

SUBJECT : GENETIC ENGINEERING & APPLICATIONS

Day : Thursday

W-2017-0969

Time : **10.00 AM TO 01.00 PM**

Max. Marks : 60

N.B.:

- 1) **Q.No.1 and Q.No.5 are COMPULSORY.** Out of the remaining questions attempt **ANY TWO** questions from each section.
- 2) Answers to both the sections should be written in **SEPARATE** answer books.
- 3) Draw neat and labeled diagrams **WHEREVER** necessary.
- 4) Figures to the right indicate **FULL** marks.

SECTION – I

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|------------|--|-------------|
| Q.1 | a) What are exo and endonucleases? Explain with suitable examples. Add a note on importance of class II restriction enzymes. | [05] |
| | b) What are special purpose vectors? Explain vectors used to maximize recombinant protein production. | [05] |
| Q.2 | Write short notes on the following: | [10] |
| | a) Methylases c) Genomic library | |
| | b) M13 vectors d) Direct gene transfer methods | |
| Q.3 | a) Explain principle and applications of PCR. Add a note on primers used in PCR. | [05] |
| | b) Explain in detail different methods used for screening genomic / cDNA library. | [05] |
| Q.4 | Explain in detail: | [10] |
| | a) Radioactive and non-radioactive DNA labeling methods. | |
| | b) Different λ based vectors. Add a note on cosmids. | |

SECTION – II

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|------------|--|------------------------------------|
| Q.5 | Explain in detail: | [10] |
| a) | Importance of <i>Saccharomyces cerevisiae</i> in genetic engineering. Add a note on vectors based on 2μ plasmid. | |
| b) | Applications of genetic engineering in agriculture. | |
| Q.6 | Write short notes on the following: | [10] |
| a) | Automated sequencing | c) Analysis of translation product |
| b) | Restriction mapping | d) Reporter genes |
| Q.7 | a) What is site directed mutagenesis(SDM)? Explain different methods for SDM. | [05] |
| | b) Compare and contrast recombinant protein expression form <i>E.coli</i> , yeast and animal cells. | [05] |
| Q.8 | Elaborate: | [10] |
| a) | Any two methods of transcript analysis. | |
| b) | Sanger's method of sequencing with suitable diagram. | |

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