

P. P. SAVANI UNIVERSITY

Fifth Semester of B.Sc. Examination
December-2021

SSBT3150-Recombinant DNA Technology-II

13.12.2021, Monday Time: 12:30 p.m. to 03:00 p.m. Maximum Marks: 60

Instructions:

1. The question paper comprises of two sections.
2. Section I and II must be attempted in separate answer sheets.
3. Make suitable assumptions and draw neat figures wherever required.
4. Use of scientific calculator is allowed.

Section-I (Total Marks - 30)

- Q.1 Short Questions [10]**
- 1.1 Objectives [05]**
- 1.1a** cDNA stands for
- A Complex DNA
 - B Complementary DNA
 - C Corona DNA
 - D All of the above
- 1.1b** PACs vectors are based on:
- A Polyoma virus
 - B Polio virus
 - C P1 bacteriophage
 - D Polyoma virus
- 1.1c** Which of the following is/are applications of cDNA library
- A To study of gene function in-vitro
 - B To determine alternative splicing in cells/tissues
 - C Discovery of novel genes
 - D All of the above
- 1.1d** Digoxigenin is used for labeling of:
- A Proteins
 - B DNA
 - C RNA
 - D All of the above
- 1.1e** Which is true for Fosmids
- A it contains λ cos site
 - B it contains F-plasmid origin of replication
 - C it is a low copy number plasmid
 - D All the options are true
- 1.1f** During RNA extraction (at phenol chloroform step), at acidic PH i.e. PH 4.0
- A RNA will stay in aqueous phase and DNA will be in organic phase
 - B RNA will stay in organic phase and DNA will be in aqueous phase
 - C RNA will stay in the interphase and DNA will be in aqueous phase
 - D None of the options are correct
- 1.1g** Which of the following is potential challenge for genomic DNA library

construction

- A Desired gene might be cut internally
- B Gene size larger than vector capacity
- C vectors have limited capacity to accommodate large insert DNA
- D All of the options are true

1.1h BAC is

- A a protein
- B an enzyme
- C a vector
- D None of the above

1.1i Which of the following enzyme is used in end-filling reaction is

- A Klenow fragment
- B RNA Pol I
- C Helicase
- D Reverse Transcriptase

1.1j In blue white screening, WHITE colonies represent

- A Transformed but non-recombinant cells
- B Transformed and recombinant cells
- C Untransformed and recombinant
- D Untransformed and non-recombinant

1.2 Answer the Following: (MCQ/Short Question/Fill in the Blanks) [05]

1.2a What does UTR stands for?

1.2b M-MLV is a RNA dependent DNA polymerase (TRUE/FALSE)

1.2c During cDNA preparation genomic DNA contamination can be checked either by DNase or intron-exon crossing primers (TRUE/FALSE)

1.2d ____ binds to streptavidin and avidin with an extremely high affinity and high specificity

1.2e What are Riboprobes?

Q.2 Short Notes (Attempt any two) [06]

- A Manniatis strategy
- B mRNA purification
- C Blue white screening

Q.3 Explain in detail (Attempt any two) [14]

- A What are the various steps of cDNA library? How cDNA library is different from Genomic library?
- B Discuss three λ bacteriophage-based vectors in detail?
- C What is a genomic library? Discuss its construction in detail.

Section-II (Total Marks - 30)

Q.1 Short Questions

[10]

1.1 Objectives

[05]

- 1.1a** _____ measures the direct release of H⁺ (protons) from the incorporation of individual bases
- A Illumina sequencing
 - B Ion Torrent sequencing
 - C Maxam- Gillbert sequencing
 - D Solexa sequencing
- 1.1b** In _____ each base emits a unique fluorescent signal, which is being recorded
- A Ion Torrent sequencing
 - B Sanger sequencing
 - C Illumina sequencing
 - D Maxam- Gillbert sequencing
- 1.1c** In Maxam-Gilbert sequencing method, pyrimidines are hydrolyzed using:
- A Dimethylsulphate
 - B Hydrazine
 - C Formic acid
 - D Hot piperidine
- 1.1d** What is the current prevalent method for genome sequencing projects
- A Chain Termination Method
 - B Next-Generation Sequencing Method
 - C Maxam-Gilbert Method
 - D All of the above
- 1.1e** The role of Urea in the sequencing gel is
- A to minimize DNA secondary structure which affects electrophoretic mobility
 - B to decrease the viscosity of gel which affects electrophoretic mobility
 - C To decrease the surface tension which affects electrophoretic mobility
 - D All of the above
- 1.1f** Which of the following is true for Sanger sequencing
- A ddNTPs are used, which lack the 3' hydroxyl group
 - B DNA fragment to be sequenced are amplified by DNA polymerase and modified nucleotides are incorporated
 - C single-stranded DNA molecules that differ in length by just a single nucleotide can be separated from one another by polyacrylamide gel electrophoresis
 - D All the options are true
- 1.1g** Pyrosequencing is an example of
- A Maxam-Gilbert Method
 - B Chain Termination Method
 - C Next-Generation Sequencing Method
 - D Sanger Sequencing Method
- 1.1h** nucleobase-specific partial chemical modification of DNA is used in:
- A Lambert Beer's Method

- B NGS method
- C Maxam Gilbert Method
- D Pyrosequencing Method

1.1i Chain-terminating ddNTPs used in Sanger sequencing perform the chain termination by:

- A 3'H prevents strand extension
- B 5'P prevents strand extension
- C 3'OH prevents strand extension
- D All the options are correct

1.1j According to Clarke and carbon probability formula, if organism A has larger genome size than organism B (P=95% for both organisms); then minimum number of clones with 45 kb DNA insert size, to be screened will be:

- A More for Organism A than B
- B More for Organism B than A
- C Equal for both the organisms A and B
- D None of the options are correct

1.2 Answer the Following: (MCQ/Short Question/Fill in the Blanks) [05]

1.2a For 20 Kb insert, λ replacement vector can be used (T/F)

1.2b A vector digested with Sau3A enzyme can be ligated to an insert digested with XbaI, as the sticky end produced by both is same (T/F)

1.2c Full form of ddNTPs is _____

1.2d _____ is a method to detect a polypeptide produced from a cloned gene

1.2e In nick translation, DNA to be processed is treated with _____ to produce single-stranded nicks.

Q.2 Short Notes (Attempt any two) [06]

- A DIG labelling
- B Plaque hybridization
- C Manual Sanger sequencing

Q.3 Explain in detail (Attempt any two) [14]

A What is high throughput sequencing? Explain two most widely used Next generation sequencing methods in detail

B Explain Maxam Gilbert Sequencing in detail?

C What do you understand by visual screening? Explain with three examples