BSEH MARKING SCHEME

CLASS- XII Biotechnology (March-2024) Code: D

 The answer points given in the marking scheme are not final. These are suggestive and indicative. If the examinee has given different, but appropriate answers, then he should be given appropriate marks.

Q.	Answers	Marks
No.		
1.	b) Southern Hybridisation	1
2.	d) All of the above	1
3.	b) ApoE	1
4.	c) Both of the above	1
5.	c) Haberlandt	1
6.	b) Epitop <mark>es</mark>	1
7.	Finite	1
8.	polyhydroxybutyrate	1
9.	antifoams	1
10.	A curator is one who reviews and checks newly	1
	submitted data ensuring all mandatory information has	
	been provided.	
11.	Ionic bonds	1
12.	Lambda or λ	1
13.	c) A is true but R is false	1

14.	b) Both A and R are true, and R is not the correct explanation of A.	1
15.	b) Both A and R are true, and R is not the correct	1
10.	explanation of A.	
16.	The dideoxynucleotide chain termination method	2
	(1 mark)	
	Chemical degradation method	
	(1 mark)	
17.	1. Electrophoretic techniques, SDS/PAGE.	2
	2. Fingerprinting.	
	3. Two-dimensional gel electrophoresis.	
	4. Protein sequencing.	
	5. Mass spectrometry.	
	(Any four,1 mark each)	
18.	In Meta <mark>genomic, th</mark> e large number of microbial	2
	genomes is collected from the different environmental	
	niche like <mark>water, air s</mark> oil etc. This is called as	
	metagenomes.	
	(1 mark)	
	The collected DNA is processed for restriction	
	digestion by the enzymes restriction endonucleases	
	and then fragments are cloned. The cloned fragments	
	are screened for presence of variety of molecules.	
	These novel molecules are used for large scale	
	production.	

	(1 mark)	
	Or	
	 Presence of strong inducible promoter 	
	 It make post translational modifications 	
	 Downstream processing is simple as it does not 	
	secrete its own protein in fermentation medium	
	(any two, 1 mark each)	
19.	• Their function is to collect and maintain	2
	important and useful microbial cultures.	
	 They function to preserve and safeguard 	
	microbial diversity.	
	 It acts as a reliable source for culture as they 	
	preserve the authenticity of the culture.	
	 The cultures are available for taxonomical 	
	stud <mark>ies, industr</mark> ial research, and academic	
	purp <mark>ose.</mark>	
	 Another main function of culture collection 	
	centers is the preparation of informative	
	documents regarding the culture maintained by	
	them.	
	(Any four, ½ mark each)	
20.	1. They can be grown in glass or plastic vessels in	2
	the presence of nutrient media.	

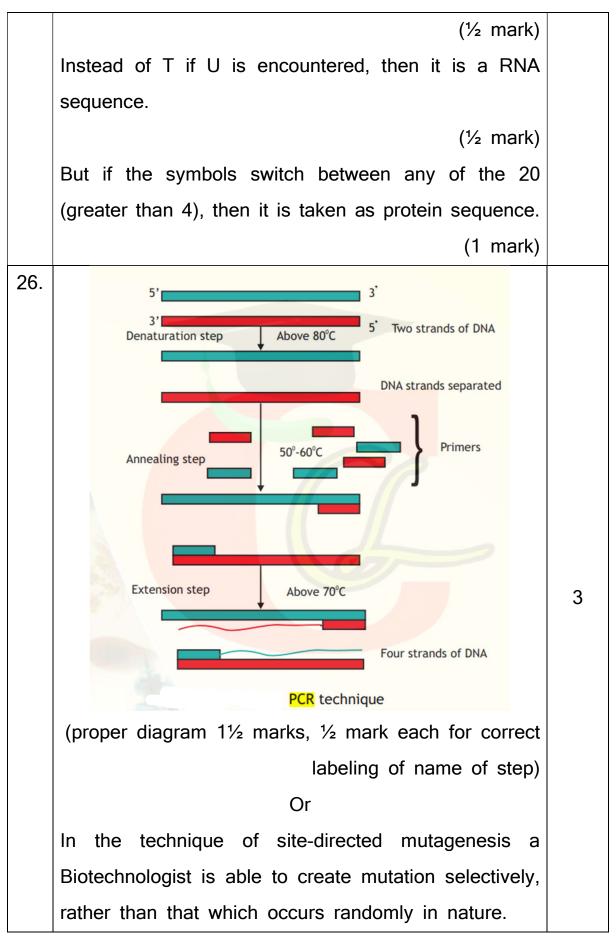
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	2. Cell can be grown till limited generation only in	
	nutrient media.	
	(1 mark each)	
	(1	
	Or	
	When the cells are cultured from the primary cell,	
	then it is called secondary culture or cell lines.	
	(1 mark)	
	The growth characteristics of cell lines is determined	
	by their doubling time, life span and formation of	
	the layer.	
	(1 mark)	
21.	Gene knock outs are defined as the removal of a	
	gene to c <mark>reate the p</mark> recise genetic modification of	
	mouse embryonic cells.	
	(½ mark)	
	Genes Knock out are created in animal and they are	
	carried to the next generation by breeding. Embryonic	2
	stem cells are used in this technology.	
	Knock out genes make the mouse model of the	
	human diseases by making the genetic modification	
	in the mouse embryonic stem cells. This is helpful for	

	various genetic diseases and their therapeutic	
	modalities.	
	(1½ mark each)	
22.	Stem cells are the cell which have capability to renew	
	themselves by mitotic division and can be	
	differentiated into diverse cell types.	
	(1 mark)	
	Application of embryonic stem cell technology are	3
	widely in medicinal field.	3
	1. Tissue repair	
	2. Gene therapy	
	3. Toxicological studies.	
	(Any two, ½ mark each)	
23.	Single cells can be isolated from either callus or any	
	other part of the plant (e.g. leaf) and cultured in liquid	
	medium.	
	(½ mark)	
	Both mechanical and enzymatic methods can be used	
	for isolation of plant cells.	
	(½ mark)	3
	Once the cells have been isolated, they may be	
	cultured by batch cultures or continuous cultures.	
	(½ mark)	
	The cell suspension cultures can be used for:	
	i) induction of somatic embryos/shoots.	

ii) in vitro mutagenesis and mutant selection. iii) genetic transformation. iv) production of secondary metabolites. (any three, $\frac{1}{2}$ mark each) Or Plant regeneration can be defined as the regeneration of a whole new plant by cultivating plant cells or explant an artificial nutrient medium. This on regeneration is possible because of the unique totipotency properties of the plant. (1 mark) There are two methods for plant regeneration which are as follow: 1. Organogenesis - It is defined as the techniques where plant organs like shoot, roots, etc. are directly developed from the explant or callus culture. (1 mark) 2. Somatic embryogenesis - It is a technique where totipotent somatic cells embark on the path of embryonic development to give somatic embryos which undergo regeneration to develop into a new plant. (1 mark)

24.	Batch culture is a closed culture system, which	
	contains an initial limited amount of nutrients. After	
	the medium is inoculated with the bacterial inoculum,	
	the organism will grow and show usual growth	
	phases.	
	(1 mark)	
	Growth results in the consumption of nutrients and	
	excretion of microbial products.	
	(½ mark)	
	At stationary phase, the growth is zero. This means,	3
	that in such a culture, growing cells are exposed to	
	continually changing environment due to gradual	
	consumption of nutrients and the accumulation of	
	metabolit <mark>es.</mark>	
	(1 mark)	
	Culturing microbes in the laboratory, in an ordinary	
	flask, is nothing but an example of batch culture.	
	(½ mark)	
25.	The usual approach taken by standard computer	
20.	programs like sequence search programs scan the	
	first 20 symbols.	2
	(1 mark)	3
	If the symbols encountered switch between any of the	
	4 bases only, then the sequence at hand is taken as	
	a DNA sequence.	



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The principle involves cloning the target gene into an	
M13 vector wherein it is presented as a single	
stranded part of the phage genome.	
(1 mark)	
A small oligonucleotide is added containing a	
complementary sequence to the gene but with one or	
more altered nucleotides.	
This allows the oligonucleotide to bind to a	
complementary portion in the target gene. This then	
acts like a primer in vitro to synthesise a double	
stranded replicative form.	
(1 mark)	
The duplex DNA molecule is then introduced into	
bacterial cells by transformation. Subsequent	
replication inside bacterial cells will produce either	
wild type or mutant gene containing plasmids. If	
appropriate expression signals are present altered	
protein can be expressed and studied.	
(1 mark)	

27.	Misfolded proteins denature easily and lose their	
	structure and function. Incorrect protein folding can	
	lead to many human diseases.	3
	(½ mark)	

's disease is an example of a	Alzhei
enerative condition caused by protein	neuro
J.	misfol
(½ mark)	
lding occurs in a cellular compartment called	Protei
plasmic reticulum.	the er
(½ mark)	
vital cellular process because proteins must	This is
ctly folded into specific, three-dimensional	be c
order to function correctly.	
(½ mark)	
or misfolded proteins contribute to the	Unfold
of many diseases.	
	P - - - - - - - - - -
(½ mark)	
isfolding is the primary cause of Alzheimer's	Protei
Parkinson's disease, Huntington's disease,	
It-Jakob disease, cystic fibrosis, Gaucher's	
, 3	diseas
enerative disorders.	neuroo
(½ mark)	

28.	This technique involves the generation and 2-D	
	analysis of peptides from a protein.	
	(½ mark)	
	Each protein has a unique peptide map (2-D analysis)	
	and hence serves as a fingerprint for the protein.	
	(½ mark)	
	The steps involved in generating a peptide	
	map/fingerprint are as follows:	
	1. Pure Hb and scHb are taken separately into test	
	tubes.	
	(½ mark)	
	2. The Hb and scHb are digested with the proteolytic	
	enzyme trypsin which cleaves the protein after basic	5
	amino ac <mark>id residues A</mark> rg and Lys.	
	(½ mark)	
	3. Two separate strips of Whatman filter paper are	
	spotted with Hb and scHb tryptic peptides and the	
	peptides allowed to separate using the technique of	
	paper electrophoresis at pH 2.0. Highly charged	
	peptides will migrate more towards the anode/cathode.	
	(½ mark)	
	4. The paper strips are dried, attached to larger	
	squares of Whatman paper and chromatographed at	
	right angles to the electrophoretic direction using a	
	solvent system Butanol:Water:Acetic acid. In such a	

	system peptides will separate based on their partition	
	coefficient between the solvent and paper which is	
	dependant on the relative hydrophobicity of the	
	peptides. More hydrophobic peptides will move with	
	the solvent to longer distances.	
	(1 mark)	
	5. The chromatograms are dried and stained with a	
	suitable visualisation reagent like Ninhydrin wherein	
	peptide containing regions appear as orange yellow	
l	spots.	
	(½ mark)	
l	6. The peptide map for Hb and scHb are compared	
	and it was found that one peptide was differently	
l	placed in the scHb map.	
	(½ mark)	
	7. On examining this peptide and determining its	
	amino acid <mark>sequence,</mark> Ingram found that it had a	
	valine substitution for glutamic acid in the peptide.	
	(½ mark)	
	Or	
	From the commercial point of view, proteins may be	
	classified into the following categories:	
	1. Blood products and vaccines.	
	2. Therapeutic antibodies and enzymes.	
	3. Therapeutic hormones and growth factors.	

4. Regulatory factors.

5. Analytical application.

6. Industrial enzymes.

7. Functional non-catalytic proteins.

8. Nutraceutical proteins.

(Any four, $\frac{1}{2}$ mark each) 1. Blood and vaccines: Α better products understanding of haematapoiesis (formation of blood cells) as well as factors responsible for blood coagulation has led to the discovery of several useful proteins. Several proteins from blood and plasma have been commercially available for decades. While these products have traditionally been obtained from blood donated by human volunteers, some are now produced by recombinant DNA technology. For example Factor VIII for treatment of Haemophilia A, Factor IX for treatment of Haemophlia B, Hepatitis B vaccine for prevention of hepatitis etc.

(1½ marks)

2. Therapeutic antibodies and enzymes: Polyclonal antibodies have been used for more than a century for therapeutic purposes. More recently monoclonal antibody preparations as well as antibody fragments produced by recombinant DNA technology have found therapeutic use. For example tissue

	-	
	plasminogen activator (t-PA) is a proteolytic enzyme	
	used to digest blocks in arteries following myocardial	
	infarctions.	
	(1½ marks)	
	Or explanation of any other categories.	
29.	Database retrieval tools include ENTREZ,	
	TAXONOMY BROWSER, and LOCUS Link.	
	(¹ / ₂ mark each)	
	Entrez is an integrated database retrieval system.	
	Through this system one can access literature (in the	
	form of abstracts), sequences and structures. Entrez	
	is an excellent system for obtaining comprehensive	
	information on a given biological question.	
	(1½ marks)	
	Locus link carries information on the official gene	_
	names and other descriptive information about genes.	5
	Additionally, through Locus link one can access	
	information on homologous genes. For example, it is	
	very convenient to obtain information on the mouse	
	homologue of a given human gene. Homologues from	
	other organisms are also available.	
	(1½ marks)	
	Or explanation of any other categories.	
	Or	
L		

Many kinds of analysis can be made using various bioinformatics tools. These include: 1. Processing raw information: The experimentally determined sequence (raw information) is processed using bioinformatics tools into genes, the proteins encoded and their function, the regulatory sequences, and inferring phylogenetic relationships. 2. Genes: Gene prediction can be done by using computer programs like GeneMark for bacterial genomes and GENSCAN for eukaryotes. 3. Proteins: Protein sequences can be inferred from the predicted genes by using simple computer programs. 4. Regulatory sequences: Regulatory sequences can also be identified and analysed by using bioinformatics tools. Inferring phylogenetic relationships: 5. Information regarding the relationships between organisms can be obtained by aligning multiple sequences, calculating evolutionary distance and constructing phylogenetic trees. 6. Making a Discovery: Using the bioinformatics tools and databases, the functions of unknown genes can be predicted.

	(Any five, 1 mark each)	
30.	1. Micropropagation	
	2. Virus-free plants	
	3. Artificial seeds	
	4. Embryo rescue	
	5. Haploids and triploids	
	6. Somatic hybrids and cybrids	
	7. Production of secondary metabolites	
	8. Somaclonal variation	
	9. In vitro plant germplasm conservation	
	10. Gametoclonal variation	
	(½ mark each)	
	Or	5
	 Social implication - There are several issues that 	0
	are raised against genetic engineering. One	
	majo <mark>r concern i</mark> s the acceptance of genetic	
	engine <mark>ering and</mark> GMOs. In most country	
	products obtained through the use of GMOs are	
	labeled.	
	(1½ marks)	
	Economical implication - It is of great concern	
	that the economic benefits from GMOs and	
	genetic engineering will be derived by the giant	
	multinational company only. Mainly GMOs are	
	greatly influenced by commercial interest rather	

than the balance between social, environmental, and economic. (1½ marks) • Environmental implication - There is great concern that GMOs developed can turn invasive and thereby causing loss of biodiversity. It can lead to environmental pollution as the use of insecticides and herbicides is increased. It can also lead to the development of resistance in insects and weed due to excessive use of insecticides and herbicides.

