MOLECULAR BASIS OF INHERITANCE

STUDY-NOTES

NUCLEIC ACIDS

Deoxyribonucleic acid (DNA) and ribonucleic acid (RNA) are the two types of nucleic acids found in living systems. DNA acts as the genetic material in most of the organisms. RNA though it also acts as a genetic material in some viruses, mostly functions as a messenger.

THE DNA

DNA is a long polymer of deoxyribonucleotides. The length of DNA is usually defined as number of nucleotides (a pair of nucleotide referred to as base pairs) present in it. DNA (deoxyribonucleic acid) is double halical structure that was cracked by Watson and Crick based on the X-ray crystallography results. Each strand of a DNA helix is composed of repeating units of nucleotides.

Structure of Nucleotide

- A nucleotide has three components a nitrogenous base, a pentose sugar (ribose in case of RNA, and deoxyribose for DNA), and a phosphate group. There are two types of nitrogenous bases Purines (Adenine and Guanine), and Pyrimidines (Cytosine, Uracil and Thymine).
- All nucleotides are common in DNA and RNA. But uracil is found in RNA, and thymine is present only in DNA. DNA is negatively charged due to the presence of negatively charged phosphate groups.
- A nitrogenous base is linked to the pentose sugar through the N-glycosidic linkage. Two nucleotides are linked through 3'-5' phosphodiester linkage to form a dinucleotide. A polymer formed in such a manner has a free phosphate group at 5'-end of ribose sugar, which is referred to as 5'-end of polynucleotide chain. The other end of the polymer has a free 3'-OH group of the ribose sugar. This is referred to as 3'- end of the polynucleotide chain. The bonding between sugar and phosphates forms the backbone of a polynucleotide chain. The nitrogenous bases are linked to sugar moiety and project from the backbone.

The salient features of the double helix structure of DNA

- Two polynucleotide chains wrap around each other, where the backbone is constituted by sugar-phosphate, and bases project inside.
- The two DNA chains are **antiparallel** to each other. It means, if one chain has the polarity 5'-3', the other has 3'-5'.
- The bases in the two strands are paired through **hydrogen bonding** forming base pairs. Adenine form two hydrogen bonds with thymine whereas cytosine form three hydrogen bonds with guanine.
- The two strands are coiled in right handed pattern.
- The plane of one base pair stacks over the other in double helix. This, in addition to H-bonds, confers stability of the helical structure.

Packaging of DNA helix

- The length of DNA double helix in a typical mammalian cell is calculated (simply by multiplying the total number of bp with distance between two consecutive bp, that is, 6.6×109 bp $\times 0.34 \times 10$ -9m/bp), it comes out to be approximately 2.2 metres.
- The positively charged basic proteins that surround the DNA is known as histones. Histones are rich in basic amino acids such as lysine and arginine. Histones are organised to form a unit of eight molecules called histone

octamer. The DNA is negatively charged and is packaged by wrapping around the positively charged histone octamer. This forms a structure called **nucleosome**.

- A nucleosome typically contains 200 base pairs of DNA helix. Nucleosomes form the repeating unit of a structure called chromatin in nucleus.
- Chromatins are thread-like stained bodies seen in nucleus. The nucleosomes in chromatin appear as 'beads-on-string' structure when viewed under electron microscope (EM). The beads-on-string structure in chromatin is packaged to form chromatin fibres that are further coiled and condensed at metaphase stage of cell division to form chromosomes. At higher levels chromatin packaging requires additional proteins. These proteins are the Non-histone Chromosomal (NHC) proteins.
- In a typical nucleus, loosely packed region of chromatin stains light and are referred to as **euchromatin**. The densely chromatin stains dark are called **heterochromatin**. Euchromatin is said to be transcriptionally active chromatin, whereas heterochromatin is inactive.

DNA AS A GENETIC MATERIAL

Griffith performed an experiment known as a transforming experiment. He used two strains of *Pneumococcus*. These two different strains were used to infect the mice. The two strains used were type III-S (smooth), that contains outer capsule made up of polysaccharide and type II-R (rough) strain does not contain capsule. The capsule protects the bacteria from the host immune system.

Transforming Principle: Griffith Experiment

Frederick Griffith, in a series of experiments with *Streptococcus pneumoniae* (bacterium responsible for pneumonia), witnessed a miraculous transformation in the bacteria.

The Griffith experiment is explained below.

- Rough strain of *Pneumococcus* is injected in mouse. The mouse is alive.
- Smooth strain of *Pneumococcus* is injected in mouse. The mouse dies.
- When heat killed smooth strain of *Pneumococcus* is injected into mouse, the mouse is alive.
- In the last set of experiment, rough strain and heat killed smooth strain are injected into mouse. The mouse dies.

This proves that there is some transforming substance present in heat killed S strain that is converting or transforming the rough strain into virulent strain that is responsible for the death of the mouse. This transforming substance was later found to be DNA.

The Genetic Material is DNA: Hershey and Chase Experiment

The genetic material is DNA, Alfred Hershey and Martha Chase (1952) performed an experiment to prove that DNA is the genetic material. They worked on bacteriophages, which are viruses that infect the bacteria. When the bacteriophage attaches to the bacteria, its genetic material enters into the bacterial cell. The viral genetic material uses the bacterial cell to synthesise more viral particles. Hershey and Chase grew some viruses on a medium that contained radioactive phosphorus and other medium that contained radioactive sulfur. When radioactive phosphorus was present in the medium the viruses contained radioactive DNA but not radioactive protein. This is because DNA contains phosphorus, but protein does not. Similarly, when the growth medium contained radioactive sulphur the viruses contained radioactive protein but not radioactive DNA. This is because DNA does not contain sulphur.

This experiment proves that DNA is the genetic material.

Properties of Genetic Material

A molecule that can act as a genetic material must fulfill the following criteria:

- (i) It should be able to generate its replica (Replication).
- (ii) It should be chemically and structurally, stable.
- (iii) It should provide the scope for slow changes (mutation) that are required for evolution.
- (iv) It should be able to express itself in the form of 'Mendelian Characters'.

CENTRAL DOGMA OF MOLECULAR BIOLOGY

It is an explanation about how the flow of genetic information occurs in a biological system. This explains how DNA replicates and then gets converted into messenger RNA (mRNA) via transcription. Then this mRNA is translated to form proteins.

DNA REPLICATION

- DNA replication is a process of producing two identical copies of DNA from a single DNA molecule. It is a
 process of biological inheritance.
- Before the cell enters the mitosis, the DNA is replicated in S phase of interphase.
- Replication begins on the specific site on the DNA, known as origin of replication.
- DNA is a double helix in which two strands are complementary to each other. These two strands of a helix separate at the time of replication to form two new DNA molecules. Out of the two strands of DNA formed, one is identical to one of the strand and another strand is complementary to the parent strand. This form of replication is known as semi-conservative replication.
- DNA polymerase in the most important enzyme involved in DNA replication.
- DNA replication is an energy dependent process. During the process of replication, the two DNA strands do not separate completely. The replication occur within the small opening of the DNA helix known as replication fork. The DNA polymerase catalyse the reaction in 5' to 3'.
- Consequently, on one strand (the template with polarity 3'-5'), the replication is continuous, while on the other (the template with polarity 5'-3'), it is discontinuous.
- The enzyme DNA ligase later joins the discontinuously synthesised fragments. The strand which is synthesised
 continuously is known as leading strand whereas the one synthesised discontinuously is known as lagging strand.

TRANSCRIPTION

- The process of copying genetic information from one strand of the DNA into RNA is termed as transcription.
- It is a process of formation of RNA such as messenger RNA from DNA before gene expression or protein synthesis occurs.
- During transcription, one of the strand of DNA acts as template for mRNA formation. The synthesis of mRNA occurs via RNA polymerase enzyme.
- Transcription usually occurs for a particular DNA segment which is required further for gene expression. Other
 than the messenger RNA, other forms of RNA such as ribosomal RNA (rRNA), micro RNA, small nuclear RNA
 (snRNA) can also be transcribed in the similar manner.
- A transcription unit in DNA consists of following three regions- a promoter, structural gene and a terminator. DNA dependent RNA polymerase catalyses the polymerisation in 5'-3' direction.
- Promoter is the region where RNA polymerase binds. Terminator defines the end of transcription.
- Transcription consists of three steps- initiation, elongation and termination.
- Initiation involves the binding of the RNA polymerase to promoter. Single type of DNA dependent RNA polymerase catalyses the transcription of all types of RNA in bacteria.
- Elongation is the process of addition of nucleotides to form the RNA.
- Termination factor helps in termination of transcription.
- The RNA synthesised after transcription is known as primary transcript. The primary transcript undergoes modification such as splicing, capping, tailing etc. Primary transcript consist of introns and exons. The removal of introns is known as **splicing**. Addition of poly-A tail at the 3'-end of the RNA is known as **tailing**. In **capping** an unusual nucleotide (methyl guanosine triphosphate) is added to the 5'-end of RNA.
- Some viruses have a property of reverse transcription. They are able to convert RNA template into DNA. The enzyme used is known as reverse transcriptase.

TRANSLATION

Translation refers to the process of polymerisation of amino acids to form a polypeptide.

- The order and sequence of amino acids are defined by the sequence of bases in the mRNA. The amino acids are joined by a bond which is known as a peptide bond
- This is the process of gene expression or protein synthesis that occurs in cytosol.
- Ribosomes are the cell organelles that are involved in protein synthesis. The messenger RNA formed by the process of transcription is decoded by ribosomes to form a polypeptide made up of amino acids.
- Messenger RNA is composed of a polymer of nucleotides or codons. Each codon consists of 3 nucleotides that will code for a single amino acid.
- Some important components that are involved in protein synthesis- ribosomes, messenger RNA and transfer RNA (tRNA).
- Transfer RNA is involved in physically linking mRNA and the amino acid sequence of proteins.

4 major steps in Translation

- Activation of amino acids: amino acids bind to specific tRNA molecules.
- Initiation of polypeptide synthesis: In capping an unusual nucleotide (methyl guanosine triphosphate) is added to the 5'-end of RNA.
- Elongation of polypeptide synthesis: It involves the addition of amino acids to the growing polypeptide chains.
- Termination of polypeptide synthesis: It involves the end of the translation of protein synthesis.

GENETIC CODE

The set of rules by which information encoded in genetic material is translated into proteins in the living cells. Salient features of genetic code are as follows:

- The codon consists of three nucleotides. 61 codons code for 20 different amino acids.
- One codon codes for one amino acid.
- One amino acid can be coded by more than one codon.
- The code is universal.

REGULATION OF GENE EXPRESSION

All the genes in the living cells are not active all the time. They become active when needed. Regulation in eukaryotes can occur at the following levels:

- Transcriptional level
- Processing level
- Transport of mRNA from nucleus to the cytoplasm
- Translational level

Lac operon or lactose operon

- An operon consists of structural genes, operator genes, promoter genes, regulator genes, and repressor.
- Lac operon consists of lac Z, lac Y and lac A genes.
- Lac Z codes for galactosidase, lac Y codes for permease and lac A codes for transacetylase. When repressor molecules bind the operator, genes are not transcribed.
- When repressor does not bind the operator and inactivated by interaction with the inducer, transcription is switched on.
- In case of lac operon, lactose is an inducer. So, binding the lactose switch on the transcription.

HUMAN GENOME PROJECT

Human genome project was a mega project that aimed to sequence every base in human genome. This project has yielded much new information. Many new areas and avenues have opened up as a consequence of the project.

The salient features of the human genome project are as follows:

- Human genome consists of a large portion of repeated sequences.
- Chromosome 1 has the most genes (2968), and the Y has the fewest (231).

- Humans are said to have about 30,000 genes.
- The functions are unknown for over 50 per cent of the discovered genes.
- The human genome contains 3164.7 million nucleotide bases.
- The average gene consists of 3000 bases, but size varies.
- Less than 2 percent of the genome codes for proteins.

DNA Fingerprinting

- DNA Fingerprinting is a technique to find out variations in individuals of a population at DNA level. It works
 on the principle of polymorphism in DNA sequences. It has immense applications in the field of forensic science,
 genetic biodiversity and evolutionary biology.
- This technique is used to determine the probable identity of a person based on the nucleotide sequences of certain regions of human DNA that are unique to individuals.
- It involves identifying differences in some specific regions in DNA sequence is called repetitive DNA.
- Since DNA from every tissue (such as blood, hair-follicle, skin, bone, saliva, sperm etc.), from an individual show
 the same degree of polymorphism, they become very useful identification tool in forensic applications.
- The technique of DNA Fingerprinting was initially developed by Alec Jeffreys. He used a satellite DNA as probe that shows very high degree of polymorphism. It was called **Variable Number of Tandem Repeats** (VNTR).

QUESTION BANK

4 6.7	MULTIPLE CHOICE QUESTIONS	Mr (40) is
1.	. Select the two correct statements out of the four (i -iv) statements given below about lac operon.	POTON PARTIE
	(i) Glucose or galactose may bind with the repressor and inactivate it.	
	(ii) In the absence of lactose the repressor binds with the operator region.	
	(iii) The z gene codes for permease.	
	(iv) This was elucidated by Francois Jacob and Jacques Monod.	
	The correct statements are	
	(a) (ii) and (iii) (b) (i) and (iii) (c) (ii) and (iv) (d) (i) and (ii).	
2.	. The 3' 5' phosphodiester linkages inside a polynucleotide chain serve to join	
	(a) one DNA strand with the other DNA strand (b) one nucleoside with another nucleoside	
	(c) one nucleotide with another nucleotide (d) one nitrogenous base with pentose sugar	
3.	. The lac operon consists of	
	(a) four regulatory genes only (b) one regulatory gene and three structural gene	es
	(c) two regulatory genes and two structural genes (d) three regulatory genes and three structural genes	enes
4.	. In eukaryotic cell transcription, RNA splicing and RNA capping take place inside the	
	(a) ribosomes (b) nucleus (c) dictyosomes (d) ER	
5.	. Which one of the following statements about the particular entity is true?	

6. Whose experiments cracked the DNA and discovered unequivocally that a genetic code is a "triplet"?

(a) Centromere is found in animal cells, which produces aster during cell division.

(b) The gene for producing insulin is present in every body cell.

(a) Hershey and Chase

(c) Nucleosome is formed of nucleotides.(d) DNA consists of core of eight histones.

(b) Morgan and Sturtevant

(c) Beadle and Tatum

(d) Nirenberg and Matthaei

7.	An octamer of 4 histones complexed with DNA forms			
	(a) endosome (b) nucleosome	(c)	mesosome	(d) centromere
8.	In the genetic dictionary, there are 64 codons as			
	(a) 64 amino acids are to be coded			
	(b) 64 types of tRNAs are present			· /
	(c) there are 44 nonsense codons and 20 sense codons			
	(d) genetic code is triplet			
9.	DNA replication is	(1,)	semiconservative and ser	midiscontinuous
	(a) conservative and discontinuous	` '	conservative and ser	maiscommuous
	(c) semiconservative and discontinuous	(u)	Collect vative	
10.	Genetic code consists of	(h)	cytosine and uracil	
	(a) adenine and guanine	'	all of these	
	(c) cytosine and guanine	, ,		
11.	The final proof for DNA as the genetic material came		Avery, MacLeod and Mo	-Carty
	(a) Hershey and Chase (c) Hargobind Khorana	()	Griffith	Carty
	If there are 999 bases in an RNA that code for a pro-	'	52	and the base at position
12.	901 is deleted such that the length of the RNA become	es 99	98 bases, how many code	ons will be altered?
	(a) 11 (b) 33		333.	(d) 1
13	During DNA replication, Okazaki fragments are used	to e	longate	
	(a) the lagging strand towards replication fork	(b)	the leading strand away	from replication fork
	(c) the lagging strand away from the replication fork	(d)	the leading strand toward	ds replication fork
14.	Which of the following RNAs should be most abundan	nt in	animal cell?	
	(a) tRNA (b) mRNA		miRNA	(d) rRNA
15.	Spliceosomes are not found in cells of			
	(a) fungi (b) animals	(c)	bacteria	(d) plants
16.	The association of histone H 1 with a nucleosome indi	icate	s that	
	(a) DNA replication is occurring	. ,	the DNA is condensed in	
	(c) the DNA double helix is exposed	(d)	transcription is occurring	5
17.	In operon concept, regulator gene functions as	(-)	latan	(d) all of these
	(a) inhibitor (b) repressor	(c)	regulator	(u) an or these
18.	Initiation codon in eukaryotes is	(a)	AUG	(d) UAG
	(a) GAU (b) AGU	(c)	AUU	(u) 0710
19.	DNA is mainly found in	(a)	cytoplasm only	(d) none of these
	(a) nucleolus (b) nucleus only	(0)	Cytopiasiii omy	(a) none of these
20.	In prokaryotes, the genetic material is	(h)	circular DNA without hi	istones
	(a) linear DNA without histones(c) linear DNA with histones	` /	circular DNA with histo	
21	In DNA, when AGCT occurs, their association is as p	(-)		
41.	(a) ATGC (b) AGCT	(c)	ACGT	(d) All of these
22	The eukaryotic genome differs from the prokaryotic g	` /		
44.	(a) the DNA is complexed with histone in prokaryotes			
	(b) the DNA is circular and single stranded in prokaryo			
	(c) repetitive sequences are present in eukaryotes			
	(d) genes in the former case are organized into operons	S		

23.	what base is responsible for not			
	(a) 5bromouracil (b) 5	imethylcytosine (c)	Guanine	(d) Adenine
24.	Genes that are involved in turning	ng on or off the transcrip	otion of a set of structura	l genes are called
	(a) redundant genes (b) re	egulatory genes (c)	polymorphic genes	(d) operator genes
25.	DNA elements, which can switch	their position, are called		. J.
	(a) cistrons (b) tr	ransposons (c)	exons	(d) introns
26.	The codons causing chain termin	ation are		
	(a) AGT, TAG, UGA (b) U	JAG, UGA, UAA (c)	TAG, TAA, TGA	(d) GAT, AAT, AGT
27.	The equivalent of a structural ge	ene is		
	(a) muton (b) c	istron (c)	operon	(d) recon
28.	Which of the following rRNAs ac	cts as structural RNA as	well as ribozyme in bacte	eria?
	(a) 5S rRNA (b) 1	8S rRNA (c)	23S rRNA	(d) 5.8S rRNA
29.	A molecule that can act as a general (a) it should be able to express it (b) it should be able to generate (c) it should be unstable structurate (d) it should provide the scope for	tself in the form of 'Mend its replica ally and chemically	elian characters'	cept . A
30.	DNA dependent RNA polymerase	catalyses transcription of	on one strand of the DNA	which is called the
				(d) antistrand
31.	There are special proteins that h	elp to open up DNA dou	ble helix in front of the	replication fork. These
	proteins are		9	-
	(a) DNA ligase (b) D	NA topoisomerase I (c)	DNA gyrase	(d) DNA polymerase I
32.	In protein synthesis, the polymeri	isation of amino acids in	volves three steps. Which	one of the following is
	not involved in the polymerisation	•		
	(a) Termination (b) In	nitiation (c)	Elongation	(d) Transcription
33.	Nucleosome core is made of			
	(a) H1, H2A, H2B and H3		H1, H2A, H2B, H4	
	(c) H1, H2A, H2B, H3 and H4		H2A, H2B, H3 and H4	
34.	Initiation codon of protein synthe			
	(a) GUA (b) G	` '		(d) AUG
35.	The transforming principle of Pna			
	(a) mRNA (b) D	NA (c)	protein	(d) polysaccharide
36.	Which one of the following is the			
	(a) UAA (b) U.	AG (c)	AUG	(d) UGA
37.	Which of the following is required	d as inducer for the expr	ession of Lac operon?	
	(a) Lactose (b) La	actose and Galactose (c)	Glucose	(d) Galactose
38.	A complex of ribosomes attached	to a single strand of RN.	A is known as	
	(a) polypeptide (b) Ol	kazaki fragment (c)	polysome	(d) polymer
39.	Which one of the following is not	applicable to RNA?		/
	(a) Heterocyclic nitrogenous bases	s (b)	Chargaff's rule	
	(c) Complementary base pairing	(d)	5' phosphoryl and 3' hydr	oxyl ends
40.	Identify the correct order of organ	nisation of genetic mater	ial from largest to smalle	st.
	(a) Genome, chromosome, gene, r	nucleotide (b)	Chromosome, genome, nu	cleotide, gene

41.	Satellite DNA is important because it		
	(a) does not code for proteins and is same	in all members of the population	
	(b) codes for enzymes needed for DNA rej	•	
	(c) codes for proteins needed in cell cycle		
	(d) shows high degree of polymorphism in p		norphism in an individual,
	which is heritable from parents to child		
42.	Out of 64 codons, 61 codons code for 20 t	-	
	(a) degeneracy of genetic code(c) wobbling of codon	(b) overlapping of gene(d) universility of codons	F
43.	Which of the following reunites the exon s	segments after RNA splicing?	
	(a) RNA polymerase (b) RNA prima	ase (c) RNA ligase	(d) RNA proteoses
44.	Which of the following enzymes are used to	to join bits of DNA?	
	(a) Ligase (b) Primase	(c) DNA polymerase	(d) Endonuclease
45.	Exon part of mRNAs have code for		
	(a) protein (b) lipid	(c) carbohydrate	(d) phospholipid
46.	Change in sequence of nucleotide in DNA	is called	. 6
	(a) mutagen (b) mutation	(c) recombination	(d) translation
47.	mRNA is synthesised on DNA template in	the direction	
	(a) $5' \to 3'$ (b) $3' \to 5'$	(c) both (a) and (b)	(d) any one
48.	In negative operon,	a	
	(a) corepressor binds with repressor	(b) corepressor does not bin	d with repressor
	(c) corepressor binds with inducer	(d) campressor has negative	effect on lac operon
49.	Which one of the following pairs of codon	s is correctly matched with their function	on or the signal for the
	particular amino acid?	(b) UUA, UCA Leucine	
	(a) AUG, ACG Start/methionine(c) GUU, GCU Alanine	(d) UAG, UGA Stop	
50	One turn of the helix in a B-form DNA is	•	
50.	(a) 2 nm (b) 20 nm	(c) 0.34 nm	(d) 3.4 nm
51	Antiparallel strands of a DNA molecule me		(4) 217 222
J1.	(a) one strand turns clockwise	time time	
	(b) one strand turns anticlockwise		
	(c) the phosphate groups of two DNA stran	nds, at their ends, share the same position	
	(d) the phosphate groups at the start of two	DNA strands are in opposite position (po	ole)
52.	Which antibiotic inhibits interaction between	en tRNA and mRNA during bacterial p	rotein synthesis?
	(a) Tetracycline (b) Erythromyc	in (c) Neomycin	(d) Streptomycin
53.	Amino acid sequence, in protein synthesis	is decided by the sequence of	
	(a) rRNA (b) tRNA	(c) mRNA	(d) cDNA
54.	E. coli cells with a mustard z gene of the la	c operon cannot grow in medium contai	ning only lactose as the
	source of energy because		
	(a) the lac operon is constitutively active in		
	(b) they cannot synthesise functional beta g		
	(c) in the presence of glucose, <i>E. coli</i> cells		

55.	Telomerase is an enzyme which is a
	(a) simple protein (b) RNA (c) ribonucleoprotein (d) repetitive DNA
56.	Using imprints from a plate with complete medium and carrying bacterial colonies, you can select streptomycin resistant mutants and prove that such mutations do not originate as adaptation. These imprints
	need to be used
	(a) on plates with and without streptomycin (b) on plates with minimal medium
	(c) only on plates with streptomycin (d) only on plates without streptomycin
57	Gene and cistron words are sometimes used synonymously because
51.	(a) one cistron contains many genes (b) one gene contains many cistrons
	(c) one gene contains one cistron (d) one gene contains no cistron
58	Types of RNA polymerase required in nucleus of eukaryotes for RNA synthesis
50.	(a) 1 (b) 2 (c) 3 (d) 4
50	Method of DNA replication in which two strands of DNA separate and synthesise new strands
57.	(a) dispersive (b) conservative (c) semiconservative (d) non conservative
60	Gene regulation governing lactose operon of E. coli that involves the lac I gene product is
00.	(a) negative and repressible because repressor protein prevents transcription
	(b) feedback inhibition because excess of b-galactosidase can switch off transcription
	(c) positive and inducible because it can be induced by lactose
	(d) negative and inducible because repressor protein prevents transcription
61.	Which one of the following is wrongly matched?
	(a) Transcription—writing information from DNA to tRNA.
	(b) Translation—using information in mRNA to make protein.
	(c) Repressor—protein binds to operator to stop enzyme synthesis.
	(d) Operon—structural genes, operator and promoter.
62.	Transformation was discovered by
	(a) Meselson and Stahl (b) Hershey and Chase (c) Griffith (d) Watson and Crick
63.	Select the correct option. Direction of RNA synthesis and direction of reading of the template DNA strand.
	Direction of RNA synthesis Direction of reading of the template DNA strand
	(a) $5' - 3'$
	(b) $3' - 5'$ $5' - 3'$ $5' - 3'$
	(0) 3. 3
	(d) 3 3
64.	Which of the following statements is not true of two genes that show 50% recombination frequency? (a) The gene show independent assortment.
	(b) If the genes are present on the same chromosome, they undergo more than one crossovers in every meiosis.
	(c) The genes may be on different chromosomes.
	(d) The genes are tightly linked.
65	The diagram given below shows an important concept in the genetic implication of DNA. Fill in the blanks
£	A to C.
	$ \begin{array}{c} \hline DNA \xrightarrow{A} mRNA \xrightarrow{B} Protein \xrightarrow{Proposed by} C \end{array} $
	DNA - mrina - Protein - C
	(a) A = Transcription, B = Translation, C = Francis Crick
	(b) A = Translation, B = Extension, C = Rosalind Franklin
	(c) A = Transcription, B = Replication, C = James Watson
	(d) A = Translation, B = Transcription, C = Ervin Chargaff

66.	. Which enzyme will be produced in a cell if there is a	non	sense mutation in the l	ac Y gene?
	(a) Transacetylase		Lactose permease and	
	(c) b galactosidase	(d)	Lactose permease	
67.	Satellite RNAs are present in some			
	(a) viroids (b) prions	(c)	bacteriophages	(d) plant viruses
68.	In mutational event, when adenine is replaced by gua	anine	, it is a case of	
	(a) frame shift mutation	(b)	transcription	
	(c) transition	(d)	transversion	/
69.	The following ratio is generally constant for a given	speci	es	
	(a) $A + G / C + T$ (b) $T + C / G + A$	(c)	G + C / A + T	(d) $A + C / T + G$
70.	What would happen if in a gene encoding a polypept	tide (of 50 amino acids, 25th	codon (UAU) is mutated
	to UAA ?			
	(a) A polypeptide of 24 amino acids will be formed.			
	(b) Two polypeptides of 24 and 25 amino acids will b	e for	med.	
	(c) A polypeptide of 49 amino acids will be formed.			
	(d) A polypeptide of 25 amino acids will be formed		8.	. 6
71.	What does "lac" refer to in what we call the lac oper			
	(a) Lactose		Lactase	
	(c) Lac insect	` '	The number 1,00,000	
72.	During translation initiation in prokaryotes, a GTP n	nolec	ule is needed in	
	(a) formation of fornylmet-tRNA			
	(b) binding of 30S subunit of ribosome with mRNA			
	(c) association of 30S mRNA with formylmet- tRNA			
	(d) association of 50S subunit of ribosome with initiat			
73.	Which one of the following triplet codes, is correctly protein synthesis or as 'start' to 'stop' codon?	ly ma	atched with its specific	ity for an amino acid in
	(a) UCG – start (b) UUU – stop	(c)	UGU - leucine	(d) UAC - tyrosine
74.	During transcription, the DNA site at which RNA po	lyme	rase binds is called	
	(a) promoter (b) regulator	(c)	receptor	(d) enhancer
75.	Degeneration of a genetic code is attributed to the			
	(a) first member of a codon	` '	second member of a co	
	(c) entire codon	(d)	third member of a code	on
76.	In the genetic code dictionary, how many codons are	used	to code for all the 20	essential amino acids?
	(a) 20 (b) 64	(c)	61	(d) 60
77.	Which of the following is not a property of the genet	ic co	de?	
	(a) Nonoverlapping (b) Ambiguous	(c)	Degeneracy	(d) Universal
78.	Genes of interest can be selected from a genomic libr	ary	by using	× 1/
	(a) cloning vectors (b) DNA probes	(c)	gene targets	(d) restriction enzymes
79.	In an inducible operon, the genes are			
	(a) usually not expressed unless a signal turns them "c	on"		
	(b) usually expressed unless a signal turns them "off"			
	(c) never expressed			
	(d) always expressed			

80.			used techniques in DNA				
	(a)	VNTR	(b) SSCP	(c)) SCAR	(d)	AFLP
81.	Rem	noval of introns and jo	ining of exons in a defined	ord	er during transcription	is ca	lled
	(a)	looping	(b) inducing	(c)) slicing	(d)	splicing
82.		ne strand of DNA has A strand sequence?	the nitrogenous base sequ	ience	as ATCTG, what would	d be	the complementary
	(a)	TTAGU	(b) UAGAC	(c)	AACTG	(d)	ATCGU
83.	Ribo	osomal RNA is actively	synthesised in			1.	
	(a)	lysosomes	(b) nucleolus	(c)	nucleoplasm	(d)	ribosomes
84.	Whi	ch one of the following	g is not a part of a transcr	iptio	n unit in DNA?		
	(a)	The inducer	(b) A terminator	(c)	A promoter	(d)	The structural gene
85.	Rem	oval of RNA polymera	se III from nucleoplasm w	vill a	ffect the synthesis of		
	(a)	tRNA	(b) hnRNA	(c)	mRNA	(d)	rRNA
86.	Wha	at are the structures c	alled that give an appeara	ance	as 'beads on string' in	the	chromosomes when
	view	ed under electron mici	roscope?				. 1
	(a)	Genes	(b) Nucleotides	(c)	Nucleosomes	(d)	Base pairs
87.	The	unequivocal proof of I	ONA as the genetic materia	al car	me from the studies on	a	
	(a)	bacterium	(b) fungus	(c)	viroid	(d)	bacteriophage
88.	Whi	ch one of the following	does not follow the centra	al do	gma of molecular biolog	gy?	
	(a)	Pea	(b) Mucor	(c)	Chlamydomonas	(d)	HIV
89.	The	one aspect which is no	t a silent feature of geneti	c cod	le, is its being		
	(a)	degenerate	(b) ambiguous	(c)	universal	(d)	specific
90.	Semi	iconservative replication	n of DNA was first demon	strat	ed in		
	(a)	Escherichia coli		. ,	Streptococcus pneumoni		
	(c)	Salmonella typhimurium	n	(d)	Drosophila melanogaste	er	
91.	In L	ac-operon, the gene 'a'	codes for the enzyme				
	(a)	lactase	(b) beta-galactosidase	(c)	permease	(d)	transacetylase
92.	Wha	t is not true for geneti	c code?				
	` '	It is nearly universal.					
		It is degenerate.					
	` '	It is unambiguous.	and in a non-continuous Cont	L:			
0.2			ead in a non-contiguous fash	nion.			
93.		the properties of admi-	ma in malation to the mains on		-:41- 41		
		_	ne in relation to thymine va hich run antiparallel one in			1 3' t	0.5'
			ine nucleotides and pyrimid				1
		_	hich run parallel in the 5' to			cquai	
94			pairs of nitrogenous bases of			atcha	d with the category
		ioned against it?	Paris of mittogenous pases (, au	cicle acids, is wrongly in	acent	a min the category
		Guanine, Adenine = Pu	rines	(b)	Adenine, Thymine = Pu	rines	
	(c)	Thymine, Uracil = Pyri	mdines		Uracil, Cytosine = Pyrir		es

95. Polysome is formed by

- (a) a ribosome with several subunits
- (b) ribosomes attached to each other in a linear arrangement
- (c) several ribosomes attached to a single mRNA
- (d) many ribosomes attached to a strand of endoplasmic reticulum

INPUT-TEXT BASED QUESTIONS

Read the following paragraphs and answer the following questions.

- I. DNA as an acidic substance present in nucleus was first identified by Friedrich Meischer in 1869. He named it as 'Nuclein'. However, due to technical limitation in isolating such a long polymer intact, the elucidation of structure of DNA remained elusive for a very long period of time. It was only in 1953 that James Watson and Francis Crick, based on the X-ray diffraction data produced by Maurice Wilkins and Rosalind Franklin, proposed a very simple but famous Double Helix model for the structure of DNA.
 - 1. On the basis of above, some statements are given as follows:
 - (i) If the sequence of bases of one strand is known, the sequence in other strand can be predicted.
 - (ii) The backbone of DNA is constituted by sugar phosphate and bases.
 - (iii) The two chains of DNA are antiparallel and complementary to each other.
 - (iv) The bases in both the strands are joined by phosphodiester bond.

Which of the following is/are correct statement(s)?

- (a) (i) and (ii)
- (b) (i), (ii) and (iv)
- (c) (ii), (iii) and (iv)
- (d) (i), (ii) and (iii)
- 2. Four kinds of bases are found in DNA. Still it is called an acid. It is because of
 - (a) Presence of phosphodiester bonds between sugars
 - (b) Presence of hydrogen bonds
 - (c) Presence of phosphoric acid
 - (d) Presence of pentose sugar
- 3. In DNA, two hydrogen bonds are found between
 - (a) Adenine and thymine

(b) Adenine and gaunine

(c) Guanine and cytosine

- (d) Cytosine and uracil
- **4.** Which of the following statement(s) is/are correct?
 - (i) The two chains are coiled in a left sided fashion.
 - (ii) A purine comes opposite to the pyrimidine in DNA helix.
 - (iii) Adenine forms two hydrogen bonds between Thymine.
 - (iv) The two chains are coiled in a right sided fashion.
 - (a) (i), (ii) and (iv)

(b) (iii) and (iv)

(c) (ii), (iii) and (iv)

(d) (i) and (ii)

- 5. Nucleoside includes
 - (a) Nitrogenous base and pentose sugar
- (b) Phosphate group and pentose sugar
- (c) Nitrogenous base and phosphate
- (d) Phosphate group only
- II. The process of copying genetic information from one strand of the DNA into RNA is termed as transcription. The principle of complementarity governs the process of transcription, except the adenosine complements now forms base pair with uracil instead of thymine. However, unlike in the process of replication, which once set in, the total DNA of an organism gets duplicated, in transcription only a segment of DNA and only one of the strands is copied into RNA. There are three major types of RNAs: mRNA (messenger RNA), tRNA (transfer RNA), and rRNA (ribosomal RNA). All three RNAs are needed to synthesise a protein in a cell.

- 1. During transcription, both strands are not copied because
 - (a) This would prevent RNA from being translated into protein.
 - (b) Both strands would code for RNA molecule with different sequences.
 - (c) A double starnaded RNA would form.
 - (d) All the above
- 2. A transcription unit is defined by three regions of DNA. Which is not the correct region?
 - (a) Promoter
- (b) Structural gene
- (c) Operator
- (d) Terminator

3. Look at the diagram.



This stage is called

- (a) Elongation
- (b) Initiation
- (c) Termination
- (d) Translation

- 4. Which of the following is the function of t-RNA?
 - (a) It provides template for transcription.
 - (c) It has the catalytic role during translation.
- (b) It brings amino acids and reads the genetic code.
- (d) It helps in initiating the process of transcription.

- 5. Which of the following is/are true?
 - (i) Sigma factor is related to initiation phase.
 - (ii) DNA dependent RNA polymerase binds to promoter to start transcription.
 - (iii) Sigma factor has the role of both initiation and termination.
 - (iv) Messenger RNA provides the template for transcription.
 - (a) (i), (iii) and (iv)

(b) (i), (ii) and (iv)

(c) (ii), (iii) and (iv)

(d) (i), (ii) and (iii)

				ANS	WERS				
1. (c)	2. (c)	3. (b)	4. (b)	5. (b)	6. (d)	7. (b)	8. (d)	9. (b)	10. (d)
11. (a)	12. (b)	13. (c)	14. (d)	15. (c)	16. (b)	17. (b)	18. (c)	19. (b)	20. (b)
21. (a)	22. (b)	23. (a)	24. (d)	25. (b)	26. (b)	27. (b)	28. (c)	29. (c)	30. (a)
31. (b)	32. (d)	33. (d)	34. (d)	35. (b)	36. (c)	37. (a)	38. (c)	39. (b)	40. (c)
41. (c)	42. (a)	43. (c)	44. (a)	45. (a)	46. (b)	47. (a)	48. (a)	49. (d)	50. (d)
51. (d)	52. (a)	53. (c)	54. (b)	55. (c)	56. (c)	57. (c)	58. (c)	59 . (c)	60. (d)
61. (a)	62. (c)	63. (a)	64. (d)	65. (a)	66. (c)	67. (d)	68. (c)	69. (c)	70. (a)
71. (a)	72. (c)	73. (d)	74. (a)	75. (d)	76. (b)	77. (b)	78. (b)	79. (a)	80. (a)
81. (d)	82. (b)	83. (b)	84. (a)	85. (a)	86. (c)	87. (d)	88. (d)	89. (b)	90. (a)
91. (d)	92. (d)	93. (b)	94. (b)	95. (c)					

EXPLANATION

- 12. The deletion occurs at 901 position so that the remaining 98 bases specifying 33 codons of amino acids will be altered.
- 21. In DNA, AGCT is associated with pair ATGC because in a DNA molecule, the purine adenine in either chain is associated with the pyrimidine thymine on the other. Similarly, purine guanine in either chain is associated with pyrimidine cytosine on the other.

- 30. The DNA-dependent RNA polymerase catalyse the polymerisation in only one direction that is $5\rightarrow 3$, the strand with polarity $5\rightarrow 3$ act as template and known as template strand.
- 42. There are 64 codons. Out of 64, 3 are stop codons or nonsense codons, i.e., these do not code for any amino acid and rest 61 code for one of the 20 amino acids. Neither of them code for more than one amino acids except GUG which normally code for valine but in certain conditions it also codes for N-formyl methionine as initiation codon.
- 52. Neomycin is a broad spectrum antibiotic effective against gram positive as well as gram negative bacteria. It's mechanism of action is by selective inhibition of protein synthesis on the 70s (prokaryotic) ribosome by inhibiting the interaction of mRNA and tRNA during translation process.
- 63. RNA polymerase initiates and extends the RNA (chain elongation) in the direction always in 5' to 3'.
- 68. In case of transition, purine base is replaced by another purine (e.g, A G) and pyrimidine is replaced by another pyrimidine (e.g, C T)
- 70. If in a gene encoding a polypeptide of 50 amino acid, 25th codon is mutated to UAA or any of the termination codon, then the chain will be terminated at that place.
- 78. Molecular probes are small DNA segments that are used to detect the presence of complementary sequences in nucleic acid samples in genomic library. These are usually formed of 200–500 nucleotide sequences. These segments or probes are labelled either with radioactive or with nonradioactive compound.

Input-Text Based Answers						
I.	1. (d)	2. (c)	3. (a)	4. (c)	5. (a)	
II.	1. (d)	2. (c)	3. (a)	4. (b)	5. (b)	