ZOOLOGY IMPORTEN INSIDE ONE MARK QUESTIONS

UNIT:II MOLECULAR GENETICS

- 1. The concept of the gene was first explained by----- (Gergor Mendel) in 1860's.
- 2. Each gene occupies a specific position called -----(Locus).
- 3. Genes may exist in several alternate forms called-----(Alleles).
- 4. Genes are capable of----- (Self-duplication) producing their own copies.
- 5. Wilhelm Hofmeister, had observed that cell bodies during mitosis called----- (Chromosome).
- 6. Friedrich Miescher, isolated a substance from the cell nuclei and called it as----- (Nuclein).
- 7. Altman(1889) was renamed a nuclein acid, and is now known as -----(DNA).
- 8. -----(Griffith's) experiment proved that DNA is the Genetic material.
- 9. Have double carbon-nitrogen ring structure and are called-----(Purines).
- 10. Have single ring structure and these are called-----(Pyrimidines).
- 11. Nucleic acids are a long chain or polymer of repating subunits called-----(Nucleotides).
- 12. Those containing Deoxyribose sugar are called-----(Deoxyribo Nucleic acid(DNA)).
- The chemical bond that link the sugar components of adjacent nucleotides are called-----(Phosphodiester bond).
- 14. The double helix model for DNA was proposed by----- (James Watson and Francis Crick)in 1953.
- 15. -----(Fraenkel-Conrat and Singer) 1957 first demonstrated that RNA is the genetic material.
- 16. The term 'RNA world' first used by----- (Walter Gilbert)in 1986.
- 17. The catalytic RNA is known as----- (Ribozyme).
- 18. -----(Hershey and Chase) clearly indicates that it is DNA that acts as a genetic material.
- Informtion storage should be able to express itself in the form of------ (Mendelian Characters).
- 20. The distance between two consecutive base pairs is -----(0.34nm or 0.34×10⁻⁹m).
- 21. The length of E.coli DNA is -----(1.36mm).
- 22. The number of base pairs of E.coli is----- $(4\times10^6 \text{m}(1.36\times10^3 \text{m}/0.34\times10^{-9}).$
- 23. DNA is held with some proteins in a region called the -----(Nucleoid).
- 24. DNA of prokaryotes is almost circular and lacks chromatin organization, hence termed -----(Genophore).
- 25. Chromatin is formed by series of repeating units called----- (Nucleosomes).
- 26. The four histone proteins are organized to form a unit of eight molecules called-----(Histone Octamere).
- 27. The negatively charged DNA is wrapped around the positively charged histone octamere to form a structure called-----(Nucleosome).
- 28. Semi-conservative replication was proposed by----- (Watson and Crick) in 1953.
- 29. The mode f DNA replication was determined in 1958 by------(Meselson and Stahl).
- The heavy DNA could be distinguished from light DNA with a technique called-----(Cesium Chloride(CsCl)) density gradient centrifugation.
- 31. DNA polymerase I also known as -----(Kornberg Enzyme).
- 32. E.coli that has 4.6×10⁶bp completes its replication process within------ (38 minutes).
- 33. Replication begins at all initiation site called the site of -----(Origin of replication or Ori).

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υ¬.	The replication is continuous and is known as the (Leading Strana).
35.	The replication is discontinuous known as the (Lagging strand).
36.	The discontinuously synthesized fragments of the lagging strand called the (Okazaki
	fragments) are joined by the enzyme DNA ligase.
37.	Y-shaped structure called the (Replication fork).
	Francis Crick proposed the(Central Dogma) in molecular biology.
39.	The process of copying genetic information from one strand of DNA into RNA is termed
	(Transcription).
40.	In eukaryotes the promoter has AT rich regions called(TATA box).
41.	In prokaryotes this region is called(Pribnow box).
42.	The structural gene may be(monocistronic (eukaryotes) or polycistronic(prokaryotes)).
43.	In prokaryotes, there are three major types of RNAsand(mRNA, tRNA, and
	rRNA).
44.	The polymerase binding sites are called(Promoters).
45.	The monocistronic structural genes are interrupted coding sequence known as
	(Exons)and non- coding sequence called(Introns).
46.	The introns are removed by a process called(Splicing).
	hnRNA undergoes additional processing called as(Capping)and(tailing).
	That mediate transfer of genes between organisms(Horizontal Gene Transfer(HGT)).
	The specific order of base pairs is called(Genetic code).
	Marshall Nirenberg, Severo Ochoa Enzyme polynucleotide phosphorylase called
	(Ochoa's enzyme).
51.	(Hargobind Khorana, Francis Crick) and many others have contributed significantly
	to decipher the genetic code.
52.	The genetic codon is a(Tripple code) and 61 codons code for amino acids.
53.	3 codons do not code for any amino acid and function as(Stop codon).
	The code is always read in a fixed direction i.e.from5' ——>3' direction called(Polarity).
	Codons are designated as termination codons and also are known as(Non-sense
	codons).
56.	Wobble Hypothesis proposed by(Crick) 1966.
	The third base of the codon is called(Wobble base).
	And Wobble base position is called(Wobble position).
	The two dimensional clover leaf model of tRNA was proposed by(Robert Holley).
	The secondary structure of tRNA depicted in looks like a(Clover leaf).
	In actual structure, the tRNA is a compact molecule which looks like an(Inverted I).
	It is called an(Adapter molecule) this term was postulated by Francis Crick.
	In additional it also shows a small lump called(Variable loop or Extra arm).
	The process of addition of amino acid to tRNA is known as(Aminoacylation or charging).
	The resultant product is called(Aminoacyation – tRNA)charged tRNA.
	Without aminoacylation tRNA is known as(Uncharged tRNA).
	The prokaryotic ribosome(70 S).
	Consists of two subunits the(Larger subunit 50 S)and(Smaller subunit 30 S).
	The ribosomes of eukaryotes(80 S).
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70.	Larger consisting ofand(60 Sand 40 S) subunits.
71.	'S' denotes the sedimentation efficient which is expressed as(Svedberg unit).
72.	One of the alternative ways of dividing up a sequence of bases in DNA or RNA into codons is
	called(Reading frame).
73.	Beginning with the start codon and which can be translated into a protein is known as an
	(Open Reading Frame(ORF)).
74.	mRNA also have some additional sequence that are not translated and are referred to as
	(Untranslated Region(UTR)).
75.	This ribosome binding site is called the(Shine- Dalgarno sequence or S-D sequence).
76.	Steps requires the correct transfer RNA, another GTP and proteins called(Elongation factors).
77.	E.coli requires three enzymes,and(Permease, β-galactosidase and
	Transacetylase).
78.	and(Jacob and Monod) proposed the classical model of lac operon to explain gene
	expression and regulation in E.coli.
79.	The international human genome project was launched in the year(1990).
	Human genome is said to have approximately(3×10 ⁹ bp).
	HGP was closely associated with the rapid development of a new area in biology called
	(Bioinformatics0.
82.	Identify all the genes approximately(3000) in human DNA.
83.	Bacteria and yeast are two commonly used hosts and these vectors are called asand-
	(BAC-Bacterial Artificial Chromosomes and YAC –Yeast Artificial Chromosome).
84.	Some repetitive DNA sequence called(Microsatellites).
85.	The latest method of sequencing even longer fragments is by a method called(Shotgu Sequence).
86.	An average gene consists of 3000 bases, the largest known human gene being (Dystrophin) with 2.4 million bases.
87.	Chromosome 1 has(2968) genes whereas Chromosome'Y' has(231) gene.
	(Pharmacogenomics) is the study of how gene affect a person's response to drugs.
	(Pharmacology) is the science of drugs.
	(Genomics) is the study of genes and their functions.
	The DNA fingerprint technique was first developed by(Alec Jeffreys)in 1985.
92.	There are(23) pairs of human chromosomes with 1.5 million pairs of genes.
93.	These nucleotide sequence are called as(Variable Number Tandem Repeats(VNTR)).
94.	DNA fingerprinting involves identifying difference in some specific regions in DNA sequence
	called(Repetitive DNA).
95.	The bulk DNA forms a major peak and the other small peaks are referred to as
	(Satellite DNA).DNA is classified into many sub categories such as(micro –
	satellite, mini-satellite),etc.

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