CET (PG)-2015

Sr. No. :

202083

Ouestion Booklet Series: A

Ans	wer Sheet.		
Roll No.	In Figures	In Words	
OMP Ans	wer Sheet Serial No.		

Subject: M.Sc. (Two Year Course)-(Bioinformatics)/(System Biology)

Time: 90 minutes Number of Questions: 75 Maximum Marks: 75

DO NOT OPEN THE SEAL ON THE BOOKLET UNTIL ASKED TO DO SO

INSTRUCTIONS

- Write your Roll No. on the Question Booklet and also on the OMR Answer Sheet in the space provided and nowhere else.
- Enter the Subject and Series Code of Question Booklet on the OMR Answer Sheet. Darken the corresponding bubbles with Black Ball Point / Black Gel pen.
- 3. Do not make any identification mark on the Answer Sheet or Question Booklet.
- 4. To open the Question Booklet remove the paper seal gently when asked to do so.
- Please check that this Question Booklet contains 75 questions. In case of any discrepancy, inform the Assistant Superintendent within 10 minutes of the start of test.
- Each question has four alternative answers (A, B, C, D) of which only one is correct. For each question, darken only one bubble (A or B or C or D), whichever you think is the correct answer, on the Answer Sheet with Black Ball Point / Black Gel pen.
- If you do not want to answer a question, leave all the bubbles corresponding to that question blank in the Answer Sheet. No marks will be deducted in such cases.
- Darken the bubbles in the OMR Answer Sheet according to the Serial No. of the questions given in the Ouestion Booklet.
- Negative marking will be adopted for evaluation i.e., 1/4th of the marks of the question will be deducted for each wrong answer. A wrong answer means incorrect answer or wrong filling of bubble.
- For calculations, use of simple log tables is permitted. Borrowing of log tables and any other material is not allowed.
- 11. For rough work only the sheets marked "Rough Work" at the end of the Question Booklet be used.
- 12. The Answer Sheet is designed for computer evaluation. Therefore, if you do not follow the instructions given on the Answer Sheet, it may make evaluation by the computer difficult. Any resultant loss to the candidate on the above account, i.e., not following the instructions completely, shall be of the candidate only.
- 13. After the test, hand over the Question Booklet and the Answer Sheet to the Assistant Superintendent on duty.
- 14. In no case the Answer Sheet, the Question Booklet, or its part or any material copied/noted from this Booklet is to be taken out of the examination hall. Any candidate found doing so, would be expelled from the examination.
- 15. A candidate who creates disturbance of any kind or changes his/her seat or is found in possession of any paper possibly of any assistance or found giving or receiving assistance or found using any other unfair means during the examination will be expelled from the examination by the Centre Superintendent/Observer whose decision shall be final.
- Telecommunication equipment such as pager, cellular phone, wireless, scanner, etc., is not permitted inside the examination hall. Use of calculator is not allowed.

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1.	The BLOSUM matrices are used for scoring	substitu	itions in :	
	(A) Amino acid sequences	(B)	Carbohydrate sequences	
	(C) Lipid sequences	(D)	Amino glycoside sequences	
2.	Which of the following steps is blocked by Ac	ctinomy	cin D in prokaryotes and Eukaryotes?	
	(A) Inhibiting initiation step of replication	(B)	Inhibiting terminating step of replication	
	(C) Inhibiting elongation step of transcription	(D)	Inhibiting elongation step in transcription	
3.	ASCII acronym in computers stands for:			
	(A) American Subscriber Code for Information	Interchar	nge	
	(B) American Service Code for Information Inte	erchange		
	(C) American Standard Code for Information In	nterchang		
	(D) American Standard Code for Interventional	Intercha		
4.	While scripting a programme in Perl langua	ge a stat	ement ends in :	-
	(A) Colon	(B)	Semicolon	
	(C) No punctuation marks needed	(D)	Numeric characters	
5.	The overall process of anaerobic glycolysis i	in muscl	e yields :	
	(A) Thirty six ATPs	(B)	Six ATPs	
	(C) Four ATPs	(D)	Two ATPs	
6.	All of the below mentioned radioactive isoto	pes emi	t β-type of radiations except:	
	(A) ³ H	(B)	(4C)	
	(C) 125I	(D)	³² P	
7.	Homologous proteins from different species	that po	ssess the same function are called :	
	(A) Orthologs	(B)	Paralogs	
	(C) Metalogs	(D)	Analogs	
8.	All of these are true for relational database	except:	the second property of the	
	(A) It follows E.F. Codd's rules			
	(B) It is a series of mathematical steps not neces	ssarily in	any logical order	
	(C) The rows correspond to record, the column	is corresp	oond to attributes	
	(D) These used structured query language for da	ata defini	tion	
9.	The RNA molecule can be present in all of the	he follow	ving forms except :	
	(A) Ribozymes	(B)	snoRNA	
	(C) miRNA	(D)	Ribonuclease	
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10.	While making a gel for electrophoresis, the polymerization reaction of acrylamide to form polyacrylamidis:
	(A) Initiated by bisacrylamide and catalyzed by persulphate
	(B) Initiated by persulphate radicals and catalyzed by TEMED
	(C) Initiated by Tris and catalyzed by persulphate radicals
	(D) Initiated by TEMED and catalyzed by bisacrylamide
11.	Which of the following statements is not true for mucopolysaccharides?
11.	(A) Polysaccharides with viscous jelly like consistency
	(B) Present in connective tissue
	(C) Contain O-glycolipid linkages
	(D) Contain O-glycosidic linkages
12	Lysosomes contain which of the following enzymes?
1.20	(A) Kinases (B) Lyases
	(C) Ligases (D) Hydrolases
13	All of the following statements are true for plasmids except :
1.3.	(A) These are self replicating
	(B) These are extra-chromosomal DNA molecules in bacteria
	(C) They usually confer a specialized characteristic to bacteria
	(D) They are always found in association with plasma membrane
14	One complete round of citric acid cycle yields :
14.	(A) Two molecules of CO ₂ , one NADH, three FADH ₂ , one high energy compound GTP
	(B) Two molecules of CO ₂ , three NADH, one FADH ₂ , one high energy compound GTP
	(C) Two molecules of CO ₂ , three NADH, one FADH, two high energy compound GTP
	(D) Two molecules of CO ₂ , one NADH, one FADH ₂ , two high energy compound GTP
15	Watern and Crick model is composed of:
15	
	See the second of the second o
	(B) Two antiparallel complementary strands (C) Two strands in parallel noncomplimentary alpha helix conformation
	n to a second to a
16	ab absent fragment must be initiated by action of which of
	(A) Helicase (B) DNA gyrase
	(C) DNA polymerasel (D) Primase
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17.	Asa	part of the electromagnetic spectrum, which	ch of	the following has the highest energy?
	(A)	Infrared	(B)	Microwave
	(C)	X-ray	(D)	Ultraviolet
18.	Whi	ich of the following could be considered as t	he be	st method for ascertaining paternity?
	(A)	DNA quantification	(B)	DNA fingerprinting
	(C)	RNA quantification	(D)	Protein fingerprinting
19.	The	First draft of human genome sequence was	rele	ased in :
	(A)	1995	(B)	2000
	(C)	2001	(D)	2003
20.	The	first biological sequence database was a:		
	(A)	Nucleotide sequence database		
	(B)	Protein database		
	(C)	Database containing nucleotide as well as prote	ins	
	(D)	Database containing nucleotides, proteins and o	other	biomacromolecules
21.	Mot	tif in a protein structure is :		
	(A)	Primary structure	(B)	Secondary structure
	(C)	Super secondary structure	(D)	Tertiary structure
22.	The	proteins functioning as gene expression re	gulat	tors do so by :
	(A)	Influencing the DNA polymerase catalysis	(B)	Activation or repression of RNA polymerase activity
	(C)	Binding to mRNA to synthesize it or destroy it	(D)	Binding to rRNA to synthesize it or destroy it
23.	The	first algorithm for comparison of protein a	nd D	NA sequences was published in 1970 by :
	(A)	Needleman and Wunsch	(B)	Smith and Waterman
	(C)	Altschul et al	(D)	Pearson and Lipman
24.	The	nucleotide sequence data retrieved from a	data	base can be in all the following formats, except:
	(A)	ASN.1	(B)	FASTA
	(C)	GenBank	(D)	PDB
25.	The	data presentation in a PDB file has :		
	(A)	Each line 60 columns wide and is terminated b	yanı	mber
	(B)	Each line 80 columns wide and is terminated b	yanı	imber
	(C)	Each line 60 columns wide and is terminated b	y an e	end-of-line indicator
	(D)	Each line 80 columns wide and is terminated b	y an c	end-of-line indicator
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26.	The selection criteria of the PAM matrix con	struction	algorithm variants will be gove	rned accord
	to:		and securopes	
	(A) PAM1 and PAM100 are both suitable for c			
	(B) PAM1 and PAM100 are both suitable for d	istantly re	lated sequences	
	(C) PAM1 and PAM100 are suitable for distant	ly and clo	sely related sequences, respectively	i a .
	(D) PAM1 and PAM100 are suitable for closely	and dist	antly related sequences, respectively	
27.	Which of the following programming lang systems?	guages w	as used initially for scripting U	NIX operat
	(A) Fortran	(B)	Java	
	(C) Assembler	(D)	Perl	
28.	Which of the following processes can be inh	bited by	rifampicin in E.Coli?	
	(A) Elongation of peptide chains	(B)	Replication of deoxyribonucleotide	sequences
	(C) Transcription of RNA sequences	(D)	Translation of RNA sequences	
29.	and the second s	are best	visible in:	
	(A) Phase contrast microscope	(B)	Flurorescence microscope	
	(C) Bright field microscope	(D)	Electron microscope	
30.	Which of the following bioinformatics progra	ams can l	oe used for predicting the transme	mbrane don
	of a receptor ?			E (A)
	(A) TMHMM	(B)	HMM	
	(C) IMM	(D)	MM	
31.	. Identify the Global alignment algorithm for	databas	e searching:	
	(A) Needleman and Wunsch	(B)	Smith and Waterman	
	(C) BLAST	(D)	Fasta	
32	. Thiamine is a:			
12	(A) Water soluble vitamin	(B)	Fat soluble vitamin	
	(C) Pyrimidine	(D)	Purine	
33	. The database InterPro is an :			
	(A) Integrated Resource of Protein sequence a	nd classif	ication	
	(B) Integrated Repository of Protein and DNA	sequenc	es	
	(C) Integrated Resource of Proteins classificate			
	(D) Integrated Repository of Protein and DNA		ation	
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34.	Ide	ntify the mismatch pair of functional group	and a	mino acid :		
	(A)	Imidazole-Histidine	(B)	Guanidino-Arginine		
	(C)	Imino-Proline	(D)	Indole-Phenylalanine		
35.	Wh	at is the molar concentration of HOH speci	ies in p	oure water ?		
	(A)	50.0 M	(B)	50.5 M		
	(C)	55.0 M	(D)	55.5 M		
36.	Wh	ich of the following cannot be used for cultu	ring e	mbryonic organs ?		
	(A)	Plasma clots	(B)	Media modified with agar		
	(C)	Fluid media	(D)	Media modified with acryl amide		
37.	Wh	ich of the following was first published con	plete	genome sequence ?		
	(A)	H. Pylori	(B)	E.Coli		
	(C)	H. influenzae	(D)	S. cerevisiae		
38.	Wa	ter as a nonlinear, polar molecule can serve	direc	tly all of the following functions in cells, except		
	(A)	It is an effective solvent	(B)	It is a reactant molecule		
	(C)	It can generate H ⁻ or OH ions	(D)	It can generate Hydride ions		
39.	Wh	ich of the following statements is not true f	or fea	tures of Protein Data Bank?		
	(A) It is the source of primary data					
	(B)	Provides efficient collection and curation of da	ita			
	(C)	Provides crystallographic and NMR structura	data			
	(D)	Does not provide any validation server				
40.	Mul	ltiple Sequence Alignment is very often use	d for	the following except :		
	(A)	Detecting similarities between the sequences	(B)	Detecting conserved regions in the sequences		
	(C)	Inferring evolutionary trees	(D)	Predicting the quaternary structure of sequences		
41.	Hyp	perchromic shift observed in a DNA sample	refle	cts which of the following ?		
	(A)	Denaturation	(B)	Renaturation		
	(C)	Destruction	(D)	Precipitation		
42.	Ide	ntify the true statement for eukaryotic RN	A poly	merases:		
	(A)	It can synthesize all types of RNAs				
	(B)	It can synthesize selective types of RNAs depe	ending	on cell types		
	(C)	There is a unique polymerase for each type of	RNA			
	(D)	There is one enzyme with various functional do	mains			

43.	The Glimmer program for gene prediction is a:		The second of th	
	(A) UNIX program from DDBJ		WINDOWS program from DDBJ	
	(C) UNIX program from TIGR	(D)	WINDOWS program from TIGR	
44.	The GenBank sequence format flat file contains	11		
11000	(A) All the information in three sections	(B)	All the information in two sections	
	(C) All the information in one section only		All the information in variant number of secti	
45.		prog	gramming language into machine languag	e woi
	be:			
	(A) Compiler	(B)	Controller	
	(C) Coprocessor	(D)	Processor	
46.	Identify the group of databases which are exclu	isivel	y specialized databases :	
	(A) Flybase, GenBank, PubMed	(B)	Flybase, AceDB, TAIR	
	(C) Flybase, GenBank, TrEMBL	(D)	Flybase, SWISS-PROT, PubMed	
47.	The Acronym of the networking specification la	ingua	age COBRA is:	MINISTER OF
	(A) Common Object Request Broker Architecture	(B)	Common Object Relational Broker Archite	cture
	(C) Common Object Request Based Architecture	(D)	Common Object Relational Based Architec	ture
48.	THE RESERVE OF THE PARTY OF THE			
	(A) The banding patterns of stains coloring cytosol	S		
	(B) The banding patterns of stains coloring chromo		s	
	(C) The banding patterns of stains coloring cell me	mbra	nes	
	(D) The banding patterns of stains coloring centros			
49	AND THE PARTY OF T			
	(A) Write only Read Many	(B)	Write once Read Many	
	(C) Write only Read Memory	(D)	Write once Read Must	
50	. The 'ktups' in FASTA database similarity sear	ch to	ol are:	
	(A) Short stretches of identical residues with a len			
	(B) Short stretches of identical residues with a ler		=I	
	(C) Short stretches of identical residues with a len			
	(D) The k number of stretches of identical residue			
51	. A cDNA library contains :			
-2.8	C. L. La La Companion	(B	Complete transcriptome information	
	The second secon	(D	1 1 C - 1 C	
	(C) Some genomic information			
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52.	Th	e low complexity regions in protein se	quences ar	e;	
	(A)	Segments represented by N-terminal an	nino acid res	idues	
	(B)	Segments represented by C-terminal arr	ino acid res	idues	
	(C)	Segments overrespresented by a small re	number of an	nino acid residues	
	(D)	Segments represented by variant types of	of amino aci	d residues	
53.	The	protein structures from secondary to	quaternar	v are maintained primarily by :	
	(A)	Metallic forces	(B)	Coordinate forces	
	(C)	Covalent forces	(D)	Noncovalent forces	
54.	Wh	ich of the following programs cannot l	be used for	molecular structure visualization?	
	(A)	Rasmol	(B)	Molscript	
	(C)	Grasp	(D)	CONPRO	
55.	Int	he first trial of gene therapy, ADA the	rapy the pa	atient was given :	
	(A)	Lymphocyte transfected with ADA gene	by simian v	iral vector	
	(B)	Lymphocyte transfected with ADA gene	by Adeno v	iral vector	
	(C)	Lymphocyte transfected with ADA gene	by retrovira	l vector	
	(D)	Lymphocyte transfected with ADA gene	without any	vector	
56.	Identify the false statement in the following:				
	(A)	RAM is a type of volatile memory			
	(B)	RAM requires a flow of electricity to reta	ain data		
	(C)	RAM is the most common type of memo	ory found in	devices such as printers	
	(D)	RAM is a memory, modules of which ca	unnot be late	r added to make the computer faster	
57.	The	graphical analysis of Phi-Psi angles fo	or protein c	onformation is known as :	
	(A)	Rajaraman plot	(B)	Ramnujan plot	
	(C)	Ramachandran plot	(D)	Ramaswami plot	
58.	Whi	ch of the following statements is not t	rue for feat	ures of SWISS-PROT ?	
	(A)	It provides core data	(B)	It provides annotation of data	
	(C)	It provides maximal redundant data	(D)	Integration with other database collections	
59.	The	following types of repetitive sequence	es can be us	sed for DNA fingerprinting, except:	
	(A)	Variable Number Tandem repeats	(B)	Short Tandem repeats	
	(C)	Simple sequence repeats	(D)	Oligopeptide repeats	

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60.	Which of the following reactions is co		Deamination
	(A) Transamination	(B)	
	(C) Carboxylation		Decarboxylation
61.	Apoenzyme is that part of the enzyme		
	(A) Polypeptide with prosthetic group of		
	(B) Polypeptide without prosthetic grou		The second secon
	(C) Prosthetic group or cofactors only w	ithout polypepti	de
	(D) Prosthetic group only		result exception in
62.	Monod-Wyman-Changeux (MWC) e	oncerted mode	are used to describe:
	(A) Action of enzymatic reactions		
	(B) Action of regulatory molecules in mo	odulating enzyma	aticaction
	(C) Action of folding accessory proteins		
	(D) Action of regulatory molecules on f	olding accessory	proteins
63.	Shine Dalgarno Sequences are prese	nt in:	
	(A) rRNA	177	tRNA
	(C) mRNA	7.800.00	snRNA
64.	Amino acid substitution from normal hae	moglobin to mut	ant Haemoglobin S is:
	(A) 6 Glu → Lys		6 Glu → Val
	(C) 6 Glu → Ala		6 Glu → Ile
65.	Which of the following conditions pro	evail when dena	turation of double stranded DNA occur
	(A) Breakage into short double strande	d fragments	
	(B) Breakage of DNA backbone		
	(C) Breakage of bases from sugar pho-	sphate backbone	The establishment of the Establishment of
	(D) Breakage of hydrogen bonds between	een two strands i	o generate single stranded fragments
66.	Identify which of the following horn	iones is not seci	eted by pancreas:
	(A) Insulin		
	(C) Somatostain	190	Glucocorticoids
67.	 In computational biology programs commonly called: 	nes, the logistic	process of analyzing a string of symbo
	(A) Priming	(B	
	(C) Predicting	(D) Pretexting

6	58. Edman degradation is the method to determine	ne the	sequence of :
	(A) Deoxyribonucleotides	(B	
	(C) Polypeptides	(D)	
6	9. What is PERL?	- 028	
	(A) Practical Extraction and Report Learning	(B)	Practical Extraction and Report Language
	(C) Practice for Exclusive and Report Language	(D)	Practical Exclusive and Report Language
7	0. To resolve proteins on the basis of their molecused?	ular	weight which of the following techniques can be
	(A) Paper chromatography	(B)	Affinity chromatography
	(C) Adsorption chromatography	(D)	Gel filtration chromatography
71	 Identify the epimeric pair of sugars from follow 	ing:	
	(A) Glucose and Fructose	(B)	Glucose and Galactose
	(C) Fructose and Mannose	(D)	Fructose and Galactose
72	The HAT selection media used for monoclonal a	ntibo	dy production would allow:
	(A) Hybrid Cells to grow in the presence of antibio	tie	
	(B) Hybrid Cells to grow in any other media		
	(C) Hybrid Cells to grow that can use salvage path	way o	f nucleotide biosynthesis
	 (D) Hybrid Cells to grow that cannot use salvage pa 	thway	of nucleotide biosynthesis
73.	Which of the following proteins have light and h	eavy	chains in their structures ?
	(A) Myoglobin		Albumin
	(C) Immunoglobulins	(D)	Insulin
74.	payagenetic tree that are descended	from	a single common ancestor
	CO \ NA concession of a con-		Monoplasty
	(C) Monomoria		Monotomy
75.			
	(A) Unweighted Protein Group Method with Arithm	etic M	eans
	(B) Unweighted Pair Group Method with Arithmetic	Mear	ns.
	(C) Unweighted Partial Group Method with Arithme	etic M	eans
	(D) Unweighted Perfect Group Method with Arithme	etic M	cans