No.		Set P
		CS) Examination Nov/Dec-2018
		nformatics DINFORMATICS
Time: 2	¹ / ₂ Hours	Max. Marks: 70
-	tions: All questions are compulsor	
	Iultiple choice Questions:-	14
1		ay gene expression database studying in
	bioinformatics. a) GEO	b) MMDB
	c) DDBJ	d) EMBL
2) is a graphical con view of information of phylogentic	trol element that presents a hierarchical
	a) Diagonal view	b) Alignment view
	c) Tree view	d) Domain view
3) The delta rule is often utilized by neural networks.	the most common class of ANNs called
	a) classificationc) unidirectional	b) backpropagationald) feed forward
2	,	tiple alignment program available either as
	a stand-alone or on-line program	
	a) exhaustivec) progressive	b) block basedd) iterative
5	, , <u> </u>	dy the organismal variation in phenotype as
	it changes during its life span.	
	a) Phenomicsc) Transcriptomics	b) Metabolomicsd) Genomics
6	, ,	sequence families not covered in Pfam-A.
	a) Prosite	
7	c) pfam b) Markov model descr	ibes the probability of the current state being
I	determined by the previous state	
	a) First-order c) Third-order	b) Second-order d) Zero-order
ε	,	database and text data along with sequence
	data retrieval system developed l	by NCBI.
	a) Dbget c) Entrez	b) dblink d) Srs
ç	,	s multiple alignments derived from the most
	conserved, ungapped regions of	
	a) Emotif c) Smart	b) Blocks d) Cath
1		refers to the percentage of
	matches of the same amino acid a) sequence identity	residues between two aligned sequences. b) sequence homology
	c) sequence similarity	d) sequence non homology

Set P

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d) sequence non homology

c) sequence similarity

Seat

	 11) alignment does not assume that the two sequences in question have similarity over the entire length. 	
	a) Local b) Global c) Heuristic d) Clustal	
	 12) The is a global network of computer networks that links government, academic and business information. a) FTP b) Gmail c) Telenet d) Internet 	
	 13) The scoring systems is called a which is derived from statistical analysis of residue substitution data from sets of reliable alignments of highly related sequences. a) substitution matrix 	
	a) substitution matrixb) sequence alignmentc) sequence identityd) sequence homology	
	14) The term sequence which is the similar at same position of same amino acid along with similar in physiochemical properties such as size, charge, and hydrophobicity.	
	a) identity b) similarity c) homology d) xenology	
Q.2	 A) Answer any four of the following What is interactomics? What is guide tree in phylogentics analysis? What is sequence identity in string matching algorithm? 	08
	4) What is local alignment in sequence analysis?5) What is BanKit in Genbank nucleotide sequence database?	
	 B) write notes on (Any Two) 1) Write the steps of system biology in study of associated disciplines. 2) Explain the PAM scoring matrix in dynamic programming algorithm. 3) Explain the Entrez search engine in data mining. 	06
Q.3	 A) Answer the following (Any Two) 1) Explain ANN in bioinformatics for analysis the biological data. 2) Explain the BLAST tool at NCBI in details. 3) Explain the Protein Information resources sequence database. 	08
	 B) Answer the following:- (Any One) 1) Explain multiple sequence alignment and its type in details. 2) Write in details about the application of neural network in bioinformatics. 	06
Q.4	 A) Answer the following (Any Two) 1) Explain the Phylip software Package in details. 2) Explain protein secondary sequence database in details. 3) Write the Gene array and analysis of gene array in bioinformatics in details. 	10
	 B) Answer the following:- (Any One) 1) Write EMBL nucleotide sequence database in details. 2) Explain the support vector machine in machine learning technique. 	04
Q.5	 Answer the following (Any Two) 1) Write in details note on HMM and HMM in bioinformatics analysis. 2) Explain the Genbank, EMBL, MSF biological sequence file format. 3) Explain the goals and scope and application of bioinformatics. 	14

Seat No.		Set P
	M.Sc. (Semester - I) (CBCS) E Bioinform CELL BIOLOGY A	natics
Time:	2 ¹ / ₂ Hours	Max. Marks: 70
Instru	ictions: 1) All questions are compulsory. 2) Draw neat and labeled diagram 3) Figures to the right indicate full	-
Q.1	 Rewrite the following sentences by using 1) is a cell organelle known at a) Ribosome c) Peroxisome 2) is known as programmed a) Apoptosis 	as suicide bags of the cell? b) Lysosome d) Mitochondria
	a) DNA polymerase θ	d) Metastasis cation of nuclear DNA in eukaryotes. b) DNA polymerase III d) DNA polymerase δ
	 4) In eukaryotic translation process a) tRNA^{met} c) tRNA^{val} 	acts as initiator tRNA molecule. b) tRNA ^{fmet} d) tRNA ^{leu}
	 5) In mRNA processing 7-methylguanosi end of the transcript. a) 3' and 5'end c) 2'end 	ne cap is added at the b) 3' end d) 5' end
	 6) During eukaryotic DNA replication pro- autonomously replicating sequences. a) DNA polymerase θ c) DNA polymerase γ 	cess binds directly to b) DNA polymerase III d) Origin recognition complex
	 7) Non-coding DNA sequences present in as a) Introns c) Coding region 	n the eukaryotic genes is also known b) Exons d) Euchromatin
	 8) are also known as common a) Desmosomes c) Gap junctions 	nunicating junctions. b) Hemi-Desmosomes d) Tight Junctions.
	,	r molecule during signal transduction.)Enzyme)Hormones
	 10) Cdk stands for a) Cycline dependent kinases b) Caspase dependent kinases c) Cytochrome depenent kinases d) Cycline degrading kinases 	

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	 11) is responsible for acidification of stomach. a) Facilitated diffusion b) Passive transport c) Proton pump d) Simple diffusion 	
	 12) enzyme is responsible for protein folding. a) Photolyase b) Photo-reductase c) Photo-resolvase d) Protein disulfide isomerase 	Э
	 13) In operon, responsible for synthesis of inhibitory protein a) Promoter b) Enhancer c) Regulatory gene d) Operator 	1.
	14) In BER, responsible for initial recognition of the lesion.a) AP endonucleasesb) DNA epinucleaesc) DNA Glycosylasesd) DNA exonucleases	
Q.2.	 A) Answer any four of the following. 1) Enlist functions of RER. 2) Define Cyclins. 3) What are exons and introns? 4) Define operon. 5) What is RNA editing? 6) Distinguish between prokaryotic and eukaryotic cell. 	08
	 B) Answer the following questions (Any Two) 1) What are gap junctions? 2) Write note on NPCs. 3) What is facilitated diffusion? 	06
Q.3.	 A) Answer the following. (Any Two) 1) Describe ultra structure and functions of Chloroplast. 2) Describe signal transduction via G-protein coupled receptors. 3) Explain process of translation in prokaryotes. 	08
	 B) Answer the following:- (Any One) 1) Describe different types of mutations with suitable examples. 2) Explain process of mitosis with neat labeled diagrams. 	06
Q.4.	 A) Answer the following. (Any Two) 1) Describe ultra structure of typical plan cell. 2) Describe process of apoptosis. 3) Explain law of independent inheritance with suitable example. 	10
	 B) Answer the following:- (Any One) 1) Describe process of DNA replication in bacteria. 2) Explain regulation of gene expression in lactose operon. 	04
Q.5.	 Answer the following. (Any Two) a) Describe mechanisms of D-loop model of DNA replication. b) Explain types of active transport with suitable examples. 	14

c) Explain protein trafficking in mitochondria and endoplasmic reticulum.

Seat]			
Seat No.				Set P
	M.Sc. (Semes	ster – I) (CBCS) Exam Bioinformatic		8
	INTROD	UCTION TO HTML &	-	
Time: 2	21/2 Hours		Ma	x. Marks: 70
Instruc	ctions: 1) All question 2) Figures to t	ns are compulsory. the right indicate full marks	5.	
	Complete the senter alternatives:-	nces by selecting correc	t answer fm the given	14
	 The most frequen a) spread c) skewness 	,	lata set is called mode median	
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Denis Ritchie
Mendel</td><td></td></tr><tr><td></td><td>7) A tag
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b) <body bgcolor</td><td>="yellow">
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a) An Icon
c) A Browser</td><td>b)</td><td>A File Manager
The Internet</td><td></td></tr><tr><td></td><td>10) Mean of a set of v
a) all values
c) First and last v</td><td></td><td>fifty percent values max and min values</td><td></td></tr></tbody></table></title></html>			

SLR-VC-49

		is the stan perating system.	idard tex	kt editor t	that com	nes with tl	he Micro	soft wind	lows
	a)	WordPad PPT			,	Excel Notepad			
	a)	n biostatics, group o block group	f individ	uals take	b) p	udy is cal population lock			
		he characteristics o called	•	ty that m	ay vary	from one	individu	al to ano	ther
	a)	static group dynamism			,	dynamic (/ariable	group		
	a)	ariables whose valu Qualitative variabl Absolute variables	e	not be ex	b) (l numeric Quantitati Continuo	ve varial	bles	<u> </u> .
Q.2.	1) 2) 3) 4)	Note: The second	ing of M ts in htm ts.	ark-up la I.					08
	1) 2)	rite notes on (Any MATLAB Random variable Formatting tags.	Two)						06
Q.3.	1) 2)	Explain in detail a Describe all attrib Write note on test	ccount of ta	of History able in d		1L.			08
	1)	Note: The following Write and explain Design a simple w	applicat	ions of b			y using h	ntml tags.	06
Q.4.	(1) 2)	Explain all body ta Write merits and o Design a frame te	ags in de demerits	etails also of Media	an.	mall prog	gram on i	it.	10
	1)	nswer the following Explain form elem Write a note on Al	ents in o						04
Q.5.	a) D	ver the following. (esign simple registr nd the mode from th	ation for	m using		tags.			14
		Marks	0-10	10-20	20-30	30-40	40-50	50-60	
		No. of students	3	8	15	20	10	4	

Cast		
Seat No.		Set P
	M.Sc. (Semester - I) (CBCS) Ex	
ΙΝΤΙ	Bioinform RODUCTION TO PROGRAMMING I	
	THROUGH C	
	2 ¹ / ₂ Hours	Max. Marks: 70
Instruc	ctions: 1) All questions are compulsory. 2) Figures to the right indicate full n	marks.
;	Complete the sentences by selecting co alternatives:- 1) OOPs stand for a) Oracle Oriented Programming	
	<pre>c) Operand oriented Programming 2) The output of the code below is #include<stdio.h> void main() { int x=5; if (x<1) printf("hello"); if(x == 5) printf("hi"); else printf("no"); } </stdio.h></pre>	d) Open Oriented Project
:	 a) hi c) no 3) The is the output of this C co #include<stdio.h></stdio.h> void main() { int i = 2; do { printf("Hi"); } while (i<2) } a) Compile time error 	b) hello d) hihello ode. b) Hi Hi
	 a) Completime error c) Hi 4) is the output of the following #include<stdio.h> main() { const int a = 5; a++; printf("%d", a); } a) 5 c) Runtime error</stdio.h> 	d) Varies

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			JLK-VC-	50
	5) is a set of instructions to do o		•	
	a) Programc) Calculation		Algorithm Debug	
	6) is the keyword used to mak	,	0	
	a) class		struct	
	c) int	d)	float	
	7) is valid C expression.			
	a) int my_num = 100,000;c) int my num = 1000;	b)	int my_num = 10000;	
	 8) Operating System is almost c a) Linux 		d in C. UNIX	
	c) Microsoft		Mac OS	
	 9) Every C Statement must end with a 	,		
	a) Čolon (:)		 Comma (,)	
	c) Slash(/)	d)	Semicolon(;)	
	10) C++ is an OOP language developed by			
	a) Dennis Ritchiec) Bjarne Stroustrup		Larry Ellison E. Khann	
	, ,	,		
	 is the process of wrapping up of (called class). 	Jata	and functions into a single unit	
	a) Encapsulation	b)	Inheritance	
	c) Handling	d)	Polymorphism	
	12) is a scope resolution symbol.			
	a) ; c) ""	b) d)		
		,		
	 13) plays an important role in initial a) Pointer 		Constructor	
	c) Destructor	,	Operator	
	14) C++ uses a unique keyword called this	to r	epresent an object that invokes a	
	function.	F)		
	a) Member c) Friend		Virtual Private	
Q.2.	A) Answer any four of the following.	۵,		08
Q.2.	1) Define Pointer.			00
	2) What is mean by Polymorphism?			
	3) Define Identifier.4) What is different ways to create a pr	oar	222	
	5) Write types of loops with example.	Ugi	2111 :	
	B) Write notes on. (Any Two)			06
	1) Write a note on Programming Langu	-		
	 Explain in detail decision making sta Explain data to as in Oct. 	atem	ents in C.	
~ ~	3) Explain data types in C++.			•••
Q.3.	A) Answer the following. (Any Two)1) Write a note on History of C.			08
	2) Explain in detail Array in C, also exp	lain	its types.	
	3) Write small program on addition of the			
	B) Answer the following:- (Any One)			06
	 Write a note on Operator in C++. Write a short note on Deinters with a 			
	2) Write a short note on Pointers with e	exar	ipie.	

Q.4.	 A) Answer the following. (Any Two) 1) Explain in detail Overloading? 2) What is Inheritance? Explain the types of Inheritance. 3) Conditional Statements with example in C 	10
	 B) Answer the following:- (Any One) 1) Explain in detail Polymorphism. 2) Briefly explain the structure of C Program. 	04
Q.5.	 Answer the following. (Any Two) a) Write a note on File Handling in C programming. b) Write C++ program on String function. c) Explain Dynamic memory Allocation in C. 	14

M.Sc. (Semester - II) (CBCS) Examination Nov/Dec-2018 Bioinformatics

ADVANCED BIOINFORMATICS

Time: 21/2 Hours

Seat

No.

Instructions: 1) Part-1, Question 1 is compulsory

- 2) Attempt any four questions from Part-II
- 3) Figures to the right indicate full marks.
- Answer to the Part-I and Part-II are to be written in same answer Booklet only.

Section – I

- Q.1 A) Rewrite the sentence after choosing the correct answer from the 07 given alternatives:
 - 1) _____ is clustering method for the creation of Phylogenetic trees created by Naruya Saitou and Nei in 1987.
 - a) MP b) ML c) UPGMA d) N-J
 - 2) The ______ based method makes predictions based on significant matches of the query sequence with sequences of known genes.
 - a) Ab initio b) Homology
 - c) HMM d) Neural network
 - 3) _____ is the replacement of a single amino acid in the primary structure of a protein with another single amino acid, which is accepted by the processes of natural selection.
 - a) PAMb) BLOSUMc) Pairwised) Multiple
 - 4) _____ plant database resource for integrative and comparative plant genome research.
 - a) Pathogen

c) MIPS

- b) PHI BASE
- d) All of these
- 5) ORF stands for _____
 - a) Open reading frequency
 c) Open reading Frame
- b) Open random framed) None of these
- 6) _____ is a web-based program that combines neural network with multiple sequence alignment.
 - a) PSI PRED b) PHD
 - c) Protparam d) pfam
- 7) _____ is a technique used by molecular biologists to produce a snapshot of the messenger RNA population.
 - a) OMIM b) SNP
 - c) SIFT d) SAGE

Max. Marks: 70

Set F

Definitions B)

- Sequence identity
 Chip
 Coil

- 4) Suffix Tree
- 5) Protparam
 6) Rooted Tree
- 7) Profile

Section – II

	Answer any four of the following.	
Q2	Explain identification of SNP in disease and SNP database with its applications.	14
Q3	Explain in detail UPGMA, NJ, Maximum Parsimony phylogenetic method in details.	14
Q4	Explain the gene prediction in prokaryotic and eukaryotic in details and add a note on plant and animal database.	14
Q5	 Answer any two of the following. a) Write a note on ExPasy server. b) Explain KEGG database in detail. c) Write a note on DNA microarray and its applications. 	14
Q6	 Write short notes on. (Any two) a) Smith-waterman algorithm b) OMIM database c) PSI and Mega Blast 	14

Seat No.		ç
	M.Sc. (Semes	ter - II) (CBCS) Examination Nov/Dec-2018
		Bioinformatics

MICROBIOLOGY AND BIOTECHNOLOGY

Time: 2¹/₂ Hours

Instructions: 1) Part-1, Question 1 is compulsory.

- 2) Attempt any four questions from Part-II.
- 3) Figures to the right indicate full marks.
- 4) Answer to the Part-I and Part-II are to be written in same answer Booklet only.

Section – I

Q.1 Rewrite the sentence after choosing the correct answer from the A) 07 given alternatives: The concept of totipotency was proposed by _____. a) Haberlandt b) Skoog c) Cocking d) Miller The genome size of mycoplasma is ____ b) 1.7 - 2.3mbps a) 1.5 - 2.5mbps c) 2.0 - 2.2mbps d) 0.5 - 1.3 mbps 3) _ _____ Hormone used to induce shoots in plant. b) Cytokines a) Gibberellins c) Both a & b d) None 4) Malachite green is used in ______ structural staining. a) Cell wall b) Capsule d) Falgella c) Endospore 5) LacZ gene is present in _____ plasmid. a) pUC19 b) pBR322 c) pUC18 d) Both a & c Serum media is used in _____ a) Plant cell culture b) Animal cell culture c) Bacterial d) None Molecular taxonomy is based on ____ b) 16S rRNA a) 28S rRNA c) 30S rRNA d) None of these 07 **Definitions:** B) 1) DNA vaccine 2) HindIII 3) Bacteriophage λ 4) Nucleoid 5) ICSB 6) Clustal W 7) Lytic cycle

Set

Max. Marks: 70



Section – II

Answer any four of the following.

Q2	Explain general characteristics and classification of plant viruses.	14
Q3	Explain the different artificial media used in animal cell culture media.	14
Q4	Explain the technique of Sanger's method of DNA sequencing	14
Q5	 Answer any two of the following. a) Explain the different staining techniques. b) Molecular genetic analysis of human diseases. c) Write a note on bacterial Transformation. 	14
Q6	 Write short notes any two of the following. a) GMOs advantages and disadvantages. b) Gene therapy c) Phylogenetic tree 	14

M.S	Sc. (Semester - II) (CBCS) E Bioinform	
	BASIC BIOCHEMISTRY	AND IMMUNOLOGY
Time: 2½ Hou	rs	Max. M
	 Part-I, Question 1 is compulsor Attempt any four questions fror Figures to the right indicate full Answer to the Part-I and Part-II only. 	n Part-II
	Part –	I
ģiv	ewrite the sentence after choosi ven alternatives: Standard free energy is denoted a) ΔE° c) ΔH°	by $\frac{1}{\Delta G^{o}}$. d) ΔF^{o}
2)	The name protein was suggested a) Kuhne c) Watson	d by b) Leninger d) Brezelius
3)	Enzymes belong to the class of _ a) Fibrous c) Membrane	proteins. b) Globular d) None
4)	is a storage polysacc	charide.

No.

Seat

-2018

Max. Marks: 70

07

answer Booklet

b) Cellulose a) Starch c) Pectin d) Chitin

5) _____ is an example of agranulocyte.

a) B cell b) Basophil d) Eosinophil c) Neutrophil

6) The agent which induces the immune response is _____.

- a) Allogen b) Autogen c) Antigen d) Antibody Interleukins are produced by _____ b) WBCs
 - a) RBCs d) None of these c) Muscle cells

Definitions B)

- 1) Enzyme
- 2) Thermodynamics
- 3) Cholesterol
- 4) Vitamin
- 5) Neutrophil
- 6) Autoimmunity
- 7) Monoclonal antibody

07





Answer any four of the following

Q.2	Explain the structural classification of proteins.	14
Q.3	Write a detailed note on innate immunity.	14
Q.4	Explain different types of antigen antibody interactions.	14
Q.5	 Answer any two of the following a) Write a note on functions of vitamins. b) Add a note on classifications of nucleic acids. c) Define enzyme. Write a note on their classification. 	14
Q.6	 Write short notes on any two a) Protein folding b) Cytokines c) Hybridoma technology 	14

		M.) Examination Nov/Dec-2018 rmatics
		IN	DUSTRIAL AND ENVIRON	MENTAL BIOTECHNOLOGY
Time	: 2½	Ηοι	Irs	Max. I
Instr	uctio	ons:	 Part-I, Question 1 is compute Attempt any four questions f Figures to the right indicate f Answer to the Part-I and Partonly. 	rom Part-II
			Par	t – I
Q.1	A)	giv	write the sentence after choo ven alternatives: Out of following is fo a) PAH c) Textile dyes	sing the correct answer from the ound to be most carcinogenic. b) Heavy metals d) Air Pollutants
		2)	Phenyl acetic acid acts as a prea) Penicillin Vc) Penicillin M	ecursor for the production of b) Penicillin G d) Cyclosporine
		3)	In Bioreactors ar a) Spargers c) Baffles	e used to prevent vortex formation. b) Impellers d) Both B & C
		4)	Amylase is a starch hydrolyzing a) <i>A. Oryzae</i> c) <i>B. Licheniformis</i>	g enzyme can be obtained by using b) <i>S. Cerevisiae</i> d) Both A & C
		5)	Which of the following techniquea) Slant Culturec) Iyophilized form technique	ies are used to maintenances of cultb) Glycerol Storage techniqued) All of these
		(\mathbf{C})	Computers are used in Disress	tor for

Seat No.

LOGY

e answer Booklet

- by using
 - es of cultures echnique

6) Computers are used in Bioreactor for

- a) Data acquisition c) A+B
- b) Data analysis d) Either A or B

d) All of these

- 7) Which of the following are Antifoaming Agent? b) Esters a) Alcohol
- c) Silicones

B) Definitions

- 1) Microbial Nutrition
- 2) Microbial Enzyme
- 3) SCP
- 4) Bio indicator
- 5) Bioremediation
- 6) Environmental Policy
- 7) Environmental impact assessment

07



SLR-VC-55

Max. Marks: 70

07

SLR- VC-55

	Answer any four of the following	
Q2	Describe about the microbial cell cultivation system.	14
Q3	Discuss in brief on Kinetics of Microbial growth.	14
Q4	Write in details of solid waste management.	14
Q5	 Answer any two of the following a) Biotechnology for clean environmental b) Heavy metal resistance in microbes c) Streptomycin production 	14
Q6	 Answer any two of the following a) Air and media sterilization b) Measurement and control of bioprocess parameters 	14

Part – II

c) Degradation of dyes

Seat No.		Set P			
	M.Sc. (Semester - III) (CBCS) Examination Nov/Dec-2018 Bioinformatics BIOLOGICAL DATABASE MANAGEMENT SYSTEM				
Time: 2	2½ Hours	Max. Marks: 70			
Instru	ctions: 1) All questions are compulsory 2) Figures to the right indicate full	l marks.			
Q.1	Complete the sentences by selecting c alternatives:- 1) In the relational modes, cardinally is te a) Number of tuples c) Number of tables	_			
	2) Cartesian product in relational algebraa) A Unary operatorc) A Ternary operator	a is b) A Binary operator d) Not defined			
	 3) DML is provided for a) Description of logical structure of d b) Addition of new structures in the da c) Manipulation & processing of data d) Definition of physical structure of d 	database system abase			
	4) In a relational model, relations are terra) Tuplesc) Tables	rmed as b) Attributes d) Rows			
	 5) An entity set that does not have sufficient is a a) Strong entity set c) Simple entity set 	b) Weak entity set			
	 6) In a Hierarchical model records are or a) Graph c) Links 	 d) Primary entity set brganized as b) List d) Tree 			
	7) Related fields in a database are groupea) Data filec) Menu	bed to form a b) Data record d) Bank			
	8) In an E-R diagram an entity set is represented a) Rectanglec) Diamond box	resent by a b) Ellipse d) Circle			
	 9) A table joined with itself is called a) Join c) Outer join 	b) Self join d) Equi join			
	10) Which of the following is used to declara) %ROWTYPEc) %CHAR	lare a record? b) %TYPE d) %DATE			

SLR-VC-56 Г

	 11) is a particular property, which describes the entity. a) Table b) Relation c) CHAR d) Attribute 	
	 12) A with respect to DBMS relates to user commands that are used to interact with a data base. a) Command b) Query c) Function d) Database 	
	13) You communicate with a RDBMS using a) Java b) Perl c) C d) SQL	
	14) While working at IBM, created the relational database model.a) Dennis Ritchieb) Larry Ellisonc) E.F. Coddd) E. Kanh	
Q.2.	 A) Answer of the following (Any Four) 1) What is mean by Sub query? 2) Name and describe three types of binary relationships. 3) Define Aggregation. 4) Write components of DBMS. 5) What is Relationship? 	08
	 B) Write notes on (Any Two) 1) Write a note on Relational Algebra. 2) Explain in detail Functional Dependency. 3) Write a short note on RDBMS KERNEL. 	06
Q.3.	 A) Answer the following (Any Two) 1) Write a note on 'Database Trigger'. 2) Describe in detail SQL and its Statements. 3) Explain in detail data types in PLSQL. 	08
	 B) Answer the following (Any One) 1) Create table using DCL commands. 2) Write a note on three schema architecture in DBMS. 	06
Q.4.	 A) Answer the following (Any Two) 1) What are partial, alternate, artificial, compound and natural key? 2) Write a simple program on DDL statements. 3) Describe Data mining tools in detail. 	10
	 B) Answer the following (Any One) 1) Write a short note on OLAP. 2) Eveloping in detail demain constraints 	04
Q.5.	 2) Explain in detail domain constraints. Answer the following (Any Two) a) Give brief account on features of DBMS. b) Write a note on Data Normalization. c) Explain in detail account on Data Model 	14

c) Explain in detail account on Data Model.

Seat No.

M.Sc. (Semester - III) (CBCS) Examination Nov/Dec-2018 **Bioinformatics** ADVANCED BIOPHYSICAL TECHNIQUES

Time: 2¹/₂ Hours

Instructions: 1) All questions are compulsory.

2) Figures to the right indicate full marks.

Q.1 Multiple choice Questions:-

- of the intermolecular bonding interactions below are 1)
 - possible for a secondary amide.
 - a) Hydrogen bonding only
 - b) Vander Waals Interactions Only
 - c) Ionic Bonding Only
 - d) Both hydrogen Bonding and ionic Bonding
- of the following statements is true regarding a secondary amide. 2)
 - a) It can only participate in hydrogen bonding as a hydrogen bond donor
 - b) It can only participate in hydrogen bonding as a hydrogen bond acceptor
 - c) It can participate in hydrogen bonding both as a hydrogen bond donor and a hydrogen bond acceptor
 - d) It cannot participate in hydrogen bonding at all
- 3) ______ of the following major aims in drug design is not related to the pharmacodynamics of a drug.
 - a) The reduction of side effects
 - b) The maximization of activity
 - c) The reduction of toxicity
 - d) The maximization of oral bioavailability
- _____ of the following is a noble gas configuration. a) $1s^2 2s^2$ 4)
 - b) 1s² 2s²2p⁶
 d) 1s² 2s²2p⁶3s²3p⁶3d¹⁰ c) $1s^2 2s^2 2p^6 3s^2$
- 5) There are _____ types of molecular orbitals.
 - a) 1 b) 2
 - c) 3 d) 4
- 6) ____ _____ of the following is non-polar bond.
 - a) C-H b) C=O d) C-C c) N-H

7) ____ _____ of the following bonds is most strongly polarized.

- a) C-H b) C=O
- d) C-N c) N-H

8) ______ of the following molecules will not display an infrared spectrum.

a) CO_2 b) N₂ c) Benzene d) HCCH

One of the following nuclei has a magnetic moment (so that an 9) NMR experiment can be performed).

b) ¹⁶O a) ¹²C c) ${}^{14}N$ d) ${}^{32}S$

14

Max. Marks: 70

SLR-VC-57



- 10) In a proton NMR experiment with a frequency (v) of 60 MHz, the magnetic field B require for resonance is 1.4 T. calculate the magnetic field required for resonance of the proton in a spectrometer with a frequency of 500 MHz.
 - a) 2.8 T

- b) 11.7 T
- c) 0.7 T d) 14.0 T
- 11) In a time-of-flight mass spectrometer, the velocity v of an accelerated ion is related to its mass by of the following.
 - a) Proportion to m (its Mass)
 - b) Inversely proportional to its mass
 - c) Proportional to the square root of its mass
 - d) Inversely proportional to the square root of its mass
- 12) For the molecule CBr4, the number of peaks which comprise the cluster for the molecular ion will be of the following.
 - a) 1 b) 4
 - d) 3 c) 5

13) A device which converts electrical energy in the form of a current into optical energy is called as b) Optical coupler

- a) Optical source
- c) Optical isolator d) Circulator

_____ of the following pairs of molecules exhibits both a pure 14) Identify rotational spectrum and a rotational Raman spectrum.

- b) CO₂ and N₂O a) O_2 and H_2O c) CO and CH_4 d) NO and DCCH
- Q.2. A) Answer any four of the following. 80 1) Valency 2) Hydrogen bond 3) Spectroscopy 4) Circular Dichroism 5) Cuvette B) Write notes on. (Any Two) 06 1) X-radiation 2) Atomic Force microscopy 3) Electron density Q.3. A) Answer the following. (Any Two) 80 1) Non-bonding interactions 2) Instrumentation of UV-Visible spectrophotometer 3) Difference between of CD and ORD. B) Answer the following:- (Any One) 06 1) Write the principle of MALDI TOF. 2) Brief out Electromagnetic spectrum. A) Answer the following. (Any Two) 10 Q.4.
 - 1) Explain the X-Ray Diffraction.
 - 2) Explain principle of Scanning Electron Microscopy.
 - Add a note on Fluorescent microscopy.

SLR-VC-57

	 B) Answer the following:- (Any One) 1) Define covalent bond. Add a note on its types. 2) Write the working of FTIR. 	04
Q.5.	 Answer the following (Any Two) a) Briefly explain the principles and theory and instrumentation of NMR b) Discuss the types of lasers with examples. c) Application of X-rays in medicine 	14

Seat No.		Set	Ρ
	M.Sc. (Semester - III) (CBCS) Ex Bioinforma OMPUTATIONAL STRUCTURE BIO	itics	3
	2 ¹ / ₂ Hours	Max. Mark	
Instruc	ctions: 1) All questions are compulsory 2) Figures to the right indicate full m	arks.	
Q.1	Multiple choice Questions:- 1) is web based service for ar of NMR structure. a) Vivaldi c) Nmrcore	nalysis, visualization and validation b) Olderado d) Nmrclust	14
	 2) Metabolism of an oral drug via the liver is a) First pass metabolism c) Last pass metabolism 	s called b) Second pass metabolism	
	 3) The secondary structure is stabilized by & a) i+5 c) i+4 	hydrogen bonds between residue I b) i+7 d) i+2	
	 4) Polar surface area of drug should be a) <120 A^o c) <150 A^o 	b) <140 A ^o d) <200 A ^o	
	 5) is database of protein models ge a) PDB c) Modbase 	enerated by homology programme. b) CSA d) 3D crunch	
	 6) Drugs are approved by a) FAD c) FDA 	b) FDD d) FAAD	
	 7) is a method of protein mod proteins that do not have homologous proteins that do not have homologous protein threading c) Protein threading 		
	 8) is the major drug metabolizi a) CYP2C19 c) CYP2D6 	ng enzymes. b) CYP2A d) CYP450	
	 9) The loop refers to a single adjacent base paired segments to form l a) Buldge c) Multibranch 		
	10) Pharmacologically inactive compoundsa) Prodrugc) Postdrug	are called b) Predrug d) Biodrug	

SLR-VC-58

	11) SSAP automatic structural alignment p database.	rogramme classifies the protein in	
	a) SCOP c) PDBsum	b) CATH d) PDBeFold	
	 12) A single transmembrane <i>α</i>-helix is calle a) Peripheral membrane protein b) Polytopic transmembrane <i>α</i>-helical c) Bitopic membrane protein d) Polytopic transmembrane <i>β</i>-sheet p 		
	13) A pattern of DNA sequence that is simi called	-	
	a) Motif c) Tunnels	b) Domain d) Pores	
	14) motif is the most commonly u DNA recognition.		
	a) Beta-Helix-Beta c) Zinc Finger	b) Helix-Beta-Helixd) Ca Finger	
Q.2.	 A) Answer any four of the following. 1) What is STRING DATABASE? 2) What is role of wwPDB database in 3) What is nuclplot in pdbsumdatabase 4) What is GRID? 5) What is interior loop in RNA structure 	protein data bank database?	08
	 B) Write notes on. (Any Two) 1) Explain the biological pathway datas 2) Describe the consequences of drug 3) Explain the development of lead libr 	target mutation.	06
Q.3.	 A) Answer the following. (Any Two) 1) Explain PDBe Fold database in deta 2) Explain the Lipnski rule of five in det 3) Explain the architecture of protein st softwares. 	ail.	08
	 B) Answer the following:- (Any One) 1) Explain the combinatorial chemistry 2) Explain the threading method for presstructures. 		06
Q.4.	 A) Answer the following. (Any Two) 1) Write in details about the Protein-DN 2) Write in detail about pharmacodynam 3) Explain protein structure classification 	mics.	10
	 B) Answer the following:- (Any One) 1) Explain the structure based drug des 2) Explain the Chou-fasman secondary 		04
Q.5.	 Answer the following. (Any Two) a) Write in details about the protein folding b) Explain the different methods of drug ta c) Explain the RCSB PDB database and p 	rget identification.	14

No.		
	M.Sc. (Semester -	· IV) (New) (CBCS) Examination Nov/
	•	Bioinformatics
	BIOLOG	IAL SIMULATIONS AND MODELING

Time: 2¹/₂ Hours

Q.1

A)

Seat

Instructions: 1) Part-1, Question 1 is compulsory

given alternatives:

- 2) Attempt any four questions from Part-II
- 3) Figures to the right indicate full marks.
- 4) Answer to the Part-I and Part-II are to be written in same answer Booklet only.

Section – I Rewrite the sentence after choosing the correct answer from the

1) Python was developed by _____ a) Stave Jobs b) Dennis Ritchi c) Guido van Rossum d) None _____ is a named piece of memory that can store a value. 2) ___ b) Array a) Constant d) Variable c) Static 3) IDLE stands for _____ a) Integrated Development Environment b) Invented Development Environment c) Independent Development Environment d) None of these _____ is a sequence of text characters in a program. 4) b) String a) Code c) Program d) Data types 5) Simulation is mimicking of _____ a) Virtual event b) Real event c) Both a & b d) None Simulation finds its application in _____ b) Chemistry a) Physics c) Biology d) All Energy is a parameter of _____ b) MC a) MD c) Both a & b d) None B) **Definitions:** 07 1) Class 2) Generators 3) Methods 4) Dictionary

on Nov/Dec-2018

Set |

Max. Marks: 70

Ρ

07

- 5) Simulation
- 6) Bacterial model
- 7) Force Field

SLR-VC-60

Section – II

Q.2	Write a note on principles and applications of simulations.	14
Q.3	Write features of python? Explain features in detail.	14
Q.4	Write a note on molecular mechanics with reference to bio-molecules.	14
Q.5	 Answer any two of the following. a) What is list? Explain with example. b) Explain python Tuples with syntax and example. c) Add a note on biological models of simulations. 	14
Q.6	 Write short notes on. (Any two) a) Characteristics of Python. b) Energy minimization in simulations. 	14

c) Examples of molecular dynamics

07

c. (Semester - IV) (New) (CBCS) E Bioinformati CLINICAL BIOINFO	cs
Hours	Max. Marks: 70
2) Attempt any four questions from Pa3) Figures to the right indicate full ma	
Part – I	
Rewrite the sentence after choosing to given alternatives:	the correct answer from the 07
1) AB SOLID 3 System generates over	gigabases per
turn. a) 30 k	b) 20
	Bioinformati CLINICAL BIOINFO Hours ns: 1) Part-1, Question 1 is compulsory. 2) Attempt any four questions from Pa 3) Figures to the right indicate full ma 4) Answer to the Part-I and Part-II are only. Part – I Rewrite the sentence after choosing to given alternatives:

- c) 15 d) 50
- _____ locates the pair of genes on the chromosomes. 2) _____
 - a) Map Viewer b) Linkage d) All of these c) Genetic Mapping
- 3) _____ disease is progressive loss of function of neurons including death of neuron.
 - a) Alzeimer's b) Huntigton's
 - d) All of these c) Parkinsons

4) A wide variety of microarray analysis tool written in _____.

- b) C++ a) C c) Python d) R
- _____ jointly sponsored by American College of Surgeons & 5) _ American Society.
 - a) NCDB b) MHDC
 - c) NHRD d) PHIBASE
- _____ is the intermediates and products of metabolism. 6) _
 - a) Metabolites b) Genomics d) None
 - c) Proteomics
- _____ to determine the amount to paid to provider. 7) b) Payer

d) Hospital

- a) Provider
- c) Patient
- Definitions. B)
 - 1) Linkage
 - 2) COPD
 - 3) Clinical Data Management
 - 4) Adverse Drug Reaction
 - 5) ChIP
 - 6) Transcriptomics
 - 7) System Dynamics

Set

Ρ

07

Seat No.

SLR-VC-61

Part – II

Answer any four of the following

Q.2	Explain Host Pathogen interaction in detail and add a note on host pathogen interaction database.	14
Q.3	Give a brief account on pharmacogenomics and their application in details.	14
Q.4	Write in detail Next Generation Sequencing QC tools and add a note on basic NGS Chemistry.	14
Q.5	 Answer any two of the following. a) Explain pathology informatics in details. b) Explain De Novo genome sequencing c) Explain goals and future challenges in Human Genome Project. 	14
Q.6	 Write short notes any two of the following. a) International Classification disease b) NGS Platforms c) Ensembl Database 	14

No. M.Sc. (Semester - IV) (New) (CBCS) Examination Nov/Dec-2018 **Bioinformatics**

RESEARCH METHODOLOGY AND IPR IN BIOINFORMATICS

Time: 2¹/₂ Hours

Instructions: 1) Part-1, Question 1 is compulsory

- 2) Attempt any four questions from Part-II
- 3) Figures to the right indicate full marks.
- 4) Answer to the Part-I and Part-II are to be written in same answer Booklet only.

Section – I

Q.1 Rewrite the sentence after choosing the correct answer from the 07 A) given alternatives: 1) ______ of the following is a major method of data collection. a) Questionnaires b) Secondary data d) All of these c) Interviews 2) It is in this section that you fully interpret & evaluate your results a) Introduction b) Method c) Results d) Discussion A literature review requires _____ a) Planning b) Clear writing c) Good writing d) All of these 4) When citation includes more than _____ authors, only the surname of the author is cited followed by et al. a) 2 b) 4 c) 5 d) 6 5) The term 'Intellectual Property Rights' covers a) Copyright b) Patent c) Trade dress d) All of the above World Intellectual Property Organization was established in _____. a) 14 March, 1959 b) 14 July, 1967 c) 14 August, 1965 d) 14 October, 1960 7) _____ is a preferred sampling method for the population with finite size. b) Cluster sampling a) Area sampling c) Purposive sampling d) Systematic sampling Definitions. 07 B) 1) Scientific journal 2) Research report 3) ANOVA 4) Hypothesis 5) Trade secrets 6) Impact factor 7) Fundamental research

SLR-VC-62

Max. Marks: 70



Section – II

Answer any four of the following.

Q2	Explain in detail the title and abstract guidelines for preparation manuscript.	14
Q3	What is Sampling? Explain in detail Types of Sampling.	14
Q4	What is research methodology? Explain in detail steps in research.	14
Q5	 Answer any two of the following. a) Write a note patenting of biological materials. b) Write a note computer and internet application in research. c) Write a note on Review of Literature. 	14
Q6	 Answer any two of the following. a) Intellectual property b) Sampling and non sampling error c) Plant variety protection in India 	14

No. M.Sc. (Semester - IV) (New) (CBCS) Examination Nov/Dec-2018 **Bioinformatics**

EMERGING AREAS OF BIOINFORMATICS

Time: 2¹/₂ Hours

Seat

Instructions: 1) Part-1, Question 1 is compulsory

- 2) Attempt any four questions from Part-II
- 3) Figures to the right indicate full marks.
- 4) Answer to the Part-I and Part-II are to be written in same answer Booklet only.

Part – I

Q.1 Rewrite the sentence after choosing the correct answer from the 07 A) given alternatives: 1) _____ is a multilateral treaty to protect endangered plants and animals. a) SITES b) ISBN d) CITES c) ICZN The most formats was created by _ Information Systems. b) SDF a) MDL c) US d) Canada medicine is a medical procedure that separates patients into different groups. a) genetic b) Natural c) Preservative d) Personalized 4) MHCpred 2.0 an updated quantitative _____ epitope prediction server. a) B-Cell b) Cyctotoxic d) All c) T-Cell 5) The _____ is a complex system of the human body and understanding it is one of the most challenging topics in biology. a) Blood b) immune c) Respiratory d) Reproductive 6) _____ coined the term the nanotechnology. a) Richard Fymann b) Nario Taniguchi c) Gopal Chandra d) Robert Koch _____ is a database of chemicals which is owned by the Royal 7) Society of Chemistry. a) ChemSpider b) Pubchem c) Zinc d) ChEBI



Set

Max. Marks: 70

B) Definitions

- 1) QSPR
 2) 2D compounds
 3) Polyphen
 4) Nano

- 5) Disease
- 6) CML
- 7) TDWG

Part – II

Answer any four of the following

Q2	Explain different chemical file format and add a note on chemical database.	14
Q3	Give a detailed account on Biodiversity informatics and explain the molecular data types in phylogenic study.	14
Q4	Explain in brief about the immunoinformatics and add a note on reverse vaccinology.	14
Q5	 Answer any two of the following a) Write a note on personalized medicine in details. b) Describe physical method for synthesis of nanomaterials. c) Write a note on MHC prediction in detail. 	14
Q6	 Answer any two of the following a) FTIR and its application b) SNP database c) Species 2000 	14

No.		
	M.Sc. (Semester	- IV) (Old) (CBCS) Examination Nov/De
	·	Bioinformatics
	BIOLOG	IAL SIMULATIONS AND MODELING

Time: 2¹/₂ Hours

Instructions: 1) Part-1, Question 1 is compulsory

- 2) Attempt any four questions from Part-II
- 3) Figures to the right indicate full marks.
- 4) Answer to the Part-I and Part-II are to be written in same answer Booklet only.

Section – I Rewrite the sentence after choosing the correct answer from the

given alternatives: 1) Python was developed by _____ a) Stave Jobs b) Dennis Ritchi d) None c) Guido van Rossum _____ is a named piece of memory that can store a value. 2) _ a) Constant b) Array d) Variable c) Static 3) IDLE stands for ____ a) Integrated Development Environment b) Invented Development Environment c) Independent Development Environment d) None of these _____ is a sequence of text characters in a program. 4) a) Code b) String d) Data types c) Program Simulation is mimicking of _____ a) Virtual event b) Real event c) Both a & b d) None Simulation finds its application in ____ b) Chemistry a) Physics d) All c) Biology Energy is a parameter of _____ b) MC a) MD c) Both a & b d) None B) **Definitions:** 07 1) Class 2) Generators 3) Methods 4) Dictionary 5) Simulation 6) Bacterial model 7) Force Field



Max. Marks: 70

07

Q.1

A)

SLR- VC-65

Section – II

Q.2	Write a note on principles and applications of simulations.	14
Q.3	Write features of python? Explain features in detail.	14
Q.4	Write a note on molecular mechanics with reference to bio-molecules.	14
Q.5	 Answer any two of the following. a) What is list? Explain with example. b) Explain python Tuples with syntax and example. c) Add a note on biological models of simulations. 	14
Q.6	 Write short notes on. (Any two) a) Characteristics of Python. b) Energy minimization in simulations. 	14

c) Examples of molecular dynamics

No. **Bioinformatics** Time: 2¹/₂ Hours **Instructions:** 1) Part-1, Question 1 is compulsory. 2) Attempt any four questions from Part-II. 3) Figures to the right indicate full marks. 4) Answer to the Part-I and Part-II are to be written in same answer Booklet only.

Part – I

Rewrite the sentence after choosing the correct answer from the Q.1 A) 07 given alternatives: 1) AB SOLID 3 System generates over _____ gigabases per turn. a) 30 b) 20 c) 15 d) 50 2) _____ _ locates the pair of genes on the chromosomes. a) Map Viewer b) Linkage c) Genetic Mapping d) All of these disease is progressive loss of function of neurons including death of neuron. a) Alzeimer's b) Huntigton's d) All of these c) Parkinsons A wide variety of microarray analysis tool written in ______. b) C++ a) C c) Python d) R jointly sponsored by American College of Surgeons & 5) _ American Society. a) NCDB b) MHDC c) NHRD d) PHIBASE _____ is the intermediates and products of metabolism. 6) _ b) Genomics a) Metabolites c) Proteomics d) None ____ to determine the amount to paid to provider. 7) a) Provider b) Payer d) Hospital c) Patient **Definitions.** 07 B) 1) Linkage 2) COPD 3) Clinical Data Management 4) Adverse Drug Reaction 5) ChIP 6) Transcriptomics 7) System Dynamics

M.Sc. (Semester - IV) (Old) (CBCS) Examination Nov/Dec-2018 CLINICAL BIOINFORMATICS

Seat

SLR-VC-66

Ρ Set

Max. Marks: 70

Part – II

Answer any four of the following

Q.2	Explain Host Pathogen interaction in detail and add a note on host pathogen interaction database.	14
Q.3	Give a brief account on pharmacogenomics and their application in details.	14
Q.4	Write in detail Next Generation Sequencing QC tools and add a note on basic NGS Chemistry.	14
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Q.6	 Write short notes any two of the following. a) International Classification disease b) NGS Platforms c) Ensembl Database 	14

M.Sc. (Semester - IV) (Old) (CBCS) Examination Nov/Dec-2018 **Bioinformatics**

EMERGING AREAS OF BIOINFORMATICS

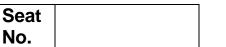
Time: 2¹/₂ Hours

Instructions: 1) Part-1, Question 1 is compulsory

- 2) Attempt any four questions from Part-II
- 3) Figures to the right indicate full marks.
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Part – I

Q.1 Rewrite the sentence after choosing the correct answer from the 07 A) given alternatives: 1) _____ is a multilateral treaty to protect endangered plants and animals. a) SITES b) ISBN d) CITES c) ICZN The most formats was created by _ Information Systems. b) SDF a) MDL c) US d) Canada medicine is a medical procedure that separates patients into different groups. a) genetic b) Natural c) Preservative d) Personalized 4) MHCpred 2.0 an updated quantitative _____ epitope prediction server. a) B-Cell b) Cyctotoxic d) All c) T-Cell 5) The _____ is a complex system of the human body and understanding it is one of the most challenging topics in biology. a) Blood b) immune c) Respiratory d) Reproductive 6) _____ coined the term the nanotechnology. a) Richard Fymann b) Nario Taniguchi c) Gopal Chandra d) Robert Koch _____ is a database of chemicals which is owned by the Royal 7) Society of Chemistry. a) ChemSpider b) Pubchem c) Zinc d) ChEBI





Set

Max. Marks: 70



B) Definitions

- 1) QSPR
 2) 2D compounds
 3) Polyphen
 4) Nano

- 5) Disease
- 6) CML
- 7) TDWG

Part – II

Answer any four of the following

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Q5	 Answer any two of the following a) Write a note on personalized medicine in details. b) Describe physical method for synthesis of nanomaterials. c) Write a note on MHC prediction in detail. 	14
Q6	 Answer any two of the following a) FTIR and its application b) SNP database c) Species 2000 	14