Reg. No. :	M 8586
Name:	V. Answereny one i
IV Semester B.Sc. Degree (CCSS-Reg./Supple./Imp Core Course in Zoology 4B04 ZLG : GENERAL INFORMATICS AND	/
Time: 3 Hours	Total Weight: 25
Ingles of the development in segal	
SECTION -A	
(General Informatics)	
Answer any one: Justify internet as a knowledge repository with specific and the s	(Weight 4) ecial reference to academic
networking. 2) Give an account on operating systems and major ap	oplication softwares. (1×4=4)
II. Answer any two:	(Weight 2 each)
3) Explain the special devices and softwares used 4) Comment on Artificial intelligence. 2. A special and othical issues of the second secon	
	(2×2=4)
III. Answer any two:	(Weight 1 each)
9) What is an operating system? 10) What is virtual reality? 11) What is cyber crime?	(2×1=2)
IV. Answer the following: 12) a) INFLIBNET stands for	(Weight 1)

b) VIRUS stands for _____

d) HTML stands for _____

c) An example for a system software is _____

(W=1)

P.T.O.



SECTION-B

(Bioinformatics)

٧.	Ans	wei	rany one :	739/		(Weight 4)	
	10.0	PR	INTS.	atabases with special re			
	14)	 Describe various database search engines in bioinformatics. Add a note on their merits. (1×4=					
VI.	Answer any two :				(Weight 2 each)		
	15)	5) What is Bioinformatics ? Give a short note on its development in recent years.					
	16)	Co	mment on Multiple	sequence alignment.			
	17)	Giv	e the applications	of Proteomics.			
	18)	De	scribe the structure	of SWISS-PORT entries.		(2×2=4)	
VII.	Ans	wei	rany four:			(Weight 1 each)	
F=A: rloa	19)	Giv	ve any two applicati	on of Metabolomics.			
			at are Micro arrays				
	21)	21) Mention any two ethical issues involved in Bioinformatics.					
	M-	2) What is DNA sequencing?					
	23)	23) What is Genbank?					
	24)	4) What are secondary databases ?					
			ention the scope of I			(4×1=4)	
VIII.	Ans	wei	r the following:			(Weight 1 each)	
	26)		FASTA stands for				
	·	b)		or primary database.			
		c)		otide sequence database	gallgraga na		
			DDBJ stands for _				
	27)		tch the following:				
	IBY		SWISS-PROT	Nucleotide database			
			KEGG	Protein database			
		c)	STAG	Metabolite database			
		100	EMBL	Search Engine			
				Blocks		(2×1=2)	