BBLEBAL JE BLABE HALL BEDIG BEM BREI.		
Reg. No. :	SECTION - B	
Name :	(Bioinformatics)	
	B.Sc. Degree (CCSS-Regular/Sup Examination, May 2014 CORE COURSE IN ZOOLOGY Control Informatics and Bioin	
Time 2 Hours		Max. Weightage: 25
Time: 3 Hours	SECTION - A	
	(General Informatics)	
	agendanta connuese noto	
I. Answer any one (V	Weight 4):	upper no manero action
1) Give an accoun	nt on the pros and cons of IT in modern	life.
2) Write an essay	on guidelines for the proper use of com	puters and internet. (1×4=4)
II. Answer any two (\	Weight 2 each):	
3) Comment on virt	tual reality	
4) Comment on fre	o dominaro de la	
5) Give the standar	rd specifications of a modern personal o	computer.
6) Account on diffe	rent antivirus package used in compute	ers. (2×2=4)
III. Answer any two (Weight 1 each):	
7) What is applica		
8) What is INFLIB	BNET?	
0) What is antiviru	118 2	
10) What is an inte	rnet browser?	
11) What is digital		(2×1=2)
IV. Answer the follow		URBUAL D
	for an application software is	
	_ is an antivirus package is an operating system.	
	- for	(1×1=1)
(S=1×Su) DOS stands	\$101	



SECTION-B

(Bioinformatics)

V. Ar	nswer any one (Weight	Semester B.Sc. Degrae (CCSS-Reg.: (
13)	Give an account on Mic	ve an account on Micro arrays. Describe the methods involved in it.		
14)	Comment on the history	, development and scope of bioinformatics.	(1×4=4)	
VI. An	swer any two (Weight 2	each):		
15)	Comment on pair wise s			
16)	Differentiate between G	enomics and Proteomics.		
17)	ccount on Nucleotide sequence databases.			
18)	Comment on Metabolor	CATTRIVIAN SHOVEN NO	(2×2=4)	
VII. An	swer any four (Weight	each):		
	What are database sea			
20)	Mention the scope of bioinformatics			
	What is EMBL?			
22)	What are primary datab			
200	Give any two application			
/III. Ans	wer the following: (Wei			
	a) Block is a			
	b) NCBI stands for	Cachillas at tac		
	c)is an e	xample for metabolite database.		
	d) Genbank is a	v yearlichting v		
27)	Match the following:			
	a) Prosite	Search Engine		
	b) EcoCyte	Secondary database		
	c) BLAST	Metabolite database		
	d) NCBI	Sequence similarity search MIPS	(2×1=2)	