

HDS-003-006509

Seat No.

B. Sc. (Bioinformatics) (Sem. V) (CBCS) Examination November / December - 2017

BI - 505 A: Molecular Modelling & Drug Designing

Faculty Code: 003

Subject Code: 006509

Time : $2\frac{1}{2}$ Hours] [Total Marks : 70

Instructions: (1) All questions are compulsory.

(2) The right-side figure indicates total marks of the question.

PART - A

1	The average angle between the orbitals of the central atom containing the bonding electron pairs in the molecule is known as between the atoms.
2	Non-covalent interactions can be classified into different categories, such as involve the attraction of ions or molecules with full permanent charges of opposite signs.
3	π -effects can be broken down into numerous categories, including u
4	Full form of SMD?
5	What is the abbreviation of SUMO protein?
6	The function performed by SUMO proteins is known as?
7	Why the product is formed unstable in the active site?
8	What is the aim of virtual screening?
9	ADMET
10	Covalent Bonds are and with respect to drug binding receptor.

11

What type of interactions in hydrogen bonding provides

extension of attraction of opposite charges?

12	is a group of diseases involving abnormal cell growth with the Potential to invade or spread to other parts of the body.						
13	Wha	What are oncogenes?					
14	Defi	Define malignancy.					
15	Defi	Define infectious disease.					
16	and have been used to generate a model of the FDA-approved inhibitor Sustiva bound to HIVRT.						
17	How	we	get semisynthetic drugs?				
18	Which theory of the activated complex by enzymes was first proposed by Pauling?						
19	Defi	ne si	uicide inhibition.				
20	Data	Data mining approaches used for target identification					
			PART – B				
1	(A)	Exp	lain Any Three :	6			
		(1)	Bond length				
		(2)	Define Valence. bond method.				
		(3)	Application of pharmacophore				
		(4)	What is cancer? Mention types of cancer.				
		(5)	What are proto onco genes?				
		(6)	Which are the steps to increase accuracy modeling the solvent effect during scoring?				
	9						
		(1)	Use of molecular mechanism				
		(2)	What are the Empirical potentials?				
		(3)	Virtual Screening				
		(4)	Describe mechanisms of anti-cancer drugs.				
		(5)	What are the steps involved in post screen analysis focusing on protein-ligand interaction. Structure based drug design?	_			
		(6)	Steps involved in the concept of Enzyme inhib	ition.			
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	(1)	Non-covalent interactions	
	(2)	Explain molecular dynamics example of applications?	
	(3)	Explain in detail about ab initio method.	
	(4)	Ligand based drug design	
	(5)	Describe mechanisms of antibiotics and antibiotic resistance.	
2 (A	a) Exp	olain Any Three :	6
	(1)	Bond Angle	
	(2)	Define Canonical ensemble (NVT).	
	(3)	Features of pharmacophore	
	(4)	What is p53 and what does it do?	
	(5)	How drug design is useful in drug discovery?	
	(6)	Features of enzyme structures and reaction pathways	
(E	B) Exp	olain Any Three :	9
	(1)	Explain molecular dynamics algorithms.	
	(2)	MMFF	
	(3)	Explain drug-induced autophagy	
	(4)	Challenges of Structure based drug design.	
	(5)	Steps involved in evaluating a structure for Structure based drug design.	
	(6)	Active site identification	
(0	C) Atte	empt Any Two :	10
	(1)	Write a note on Monte Carlo Simulation for conformational analysis	
	(2)	Drug-receptor interactions	
	(3)	Explain diseases related to cancer	
	(4)	How proto oncogene converts to an oncogene?	
	(5)	Describe infectious disease and its mechanisms in detail.	
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(C) Attempt Any **Two**: