M. SC. BIOINFORMATICS SEM.-III (2013 COURSE) (CHOICE BASED CREDIT SYSTEMS) : SUMMER - 2018

SUBJECT : CHEMINFORMATICS & DRUG DESIGN

Day: Date:	Saturday 07/04/2018		S-2018-1131		TIME:		
N.B :	ea 2) Fig	 Q. 1 and Q. 5 are Compulsory. Out of the remaining questions, attempt any 2 from each sections. Figures to the right indicate Full marks. Answers to both the sections should be written in Separate answer books. 					
Q. 1	a) c) e)	Section - I Define: Hosoya index b) Adjacency matrices E-state d) Tanimoto coefficient Lipinski's rule					
Q. 2	a) b) c)	What is the full form of SMILES and SDF in molecular file format? How can you search molecular patterns using SMARTS? Explain the role of cheminformatics in pharmaceutical research. OR Explain descriptors with example of 1D, 2D and 3D descriptors.					
Q. 3	a) b)	Explain molecular graph in brief. What is the lead molecule? How molecular databases are crucial to find lead molecules? OR Explain Partial Least Square method.					
Q. 4	c) a) b) c) c)	Explain – "Role of similarity matrix in molecular similarity search". Define molecular refractivity. Write short note on electronic descriptors. Discuss importance of ADMET profile being a drug molecule. OR Explain – "Role of small molecular databases to identify potential lead molecules".				(04) (02) (04) (04)	
Q. 5	a) c) e)	Define: Pharmacophore kees	eys	Second b) d)	tion - II Lead optimization Test set	(10)	
Q. 6	a) b) b) c)	Define receptor. Write short note on ligand-based drug design. OR Discuss the role of drug design in pharmaceutical industry. How bioinformatics merged to cheminformatics to develop lead molecules?				(02) (04) (04)	
Q.7	a) b) b) c)	Define QSPR. Write short note on 3D QSAR. OR Write short note on applicability domain of QSAR. Write short note on quantum chemical descriptors.					
Q.8	a) b) c)	Define scoring function in molecular docking. Define virtual screening. Explain role of virtual screening for identification of lead molecules. Explain receptor-based pharmacophore model. OR					
	c)	Differentiate between ligand-based and receptor-based drug designing.					

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