## Department of Biochemical Engineering and Biotechnology

## BEL714 Protein Science and Engineering

Minor-I (1st semester 2016-17)

	Controduct 2010-171			
Max Marks:25	Max Marks:25		Time:1 hour	
Q.1 Write down the structu	are of the following peptide:	0,00	our px_C	
Trp-Pro-Asp-Lys	7.5	, c	(2)	
V 101 tile same! Show schemat	nantly made up of beta structures. I ically the arrangement of the hydrodipole moment. How does it arise	ogen honds in allu-pa	could	
<ul> <li>Q.3 Explain:         <ul> <li>How globin fold has been preserved in spite of low sequence identities.</li> <li>How new enzyme activities have evolved (are new enzymes formed from random sequences generated by recombination or do these arise from a preexisting set of enzyme activities)</li> </ul> </li> </ul>				
Is there any experiment	al evidence in literature that supp	ort your answer?	(4)	
proteins that show this pack	t down for helix-helix packing and ing. Would you expect this packi r answer with concrete examples.	ng to be affected by	nples of amino (5)	
0.5 Examine the protein str	ructure on the next page (Fig.1) at where the active site may be loca	nd represent it in the	form of (3)	
Q.6 Explain giving suitable	e examples:			
Domain shuffling and second database and its			(3) (3)	