Dr. Tirth Ray Sigh

(CO1-4) [2\*2=4]

## JAYPEE UNIVERSITY OF INFORMATION TECHNOLOGY, WAKNAGHAT TEST -2 EXAMINATIONS April 2019

## B.Tech VIII Semester (BT and BI branches)

Course Code: 15B1WBI834		MAX. MARKS: 25	A STATE
Course Name: Computational Molecular Evolu	tion	MAX. TIME: 1.5 hrs	
Course Credits: 03			
			*****
Note: All questions are compulsory. Carrying of n	nobile phone during ex	aminations will be treat	ed as
case of unfair means.			
Q1. Define Kimura's two parameter model for nu	cleotide substitutions for	or studying evolution alo	ong wit
ts derivation.		(CO3)	[5]
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Q2. What is codon usage bias in protein coding ge	enes? Define Codon Ac	daptive Index (CAI) for	the
codon usage.		(CO2,	3) [4]
		(001	4) [0]
Q3. Explain the evolution of human in your scient	tific opinion.	(CO1-	4) [2]
Q4. Define introns evolutionary theories and their	cignificance	(CO2,	3) [3]
2 Domic mirons evolutionary incomes and then	Significance.	(002,	ردا (۲
Q5. What is gene duplication? How gene duplicat	ion is associated with f	functional aspects of gen	e? Hov
you date gene duplication in any species?		(CO2-	
, and the same of		(002	., г.,
Q.6. Define following terms.			
(a) Genetic drift (b) Selection	(c) Gene loss	(CO1-4) [1	*3=3]
0.7 Differentiate between the following:			
1) / Lutterenfiete between the following:			

(b) Orthologs and Paralogs

(a) Homology and Homoplasy