

JAYPEE UNIVERSITY OF INFORMATION TECHNOLOGY, WAKNAGHAT
TEST -1 EXAMINATION

September 2018

B.Tech. (Bioinformatics): Vth Semester

Course Code: 15B11BI512

MAX. MARKS: 15

Course Name: Computational Genomics

Course Credits: 04

MAX. TIME: 1 Hr

Note: All questions are compulsory. Carrying of mobile phone during examinations will be treated as case of unfair means.

Q.1. Explain the following terms with a suitable example:

(1×5=5)

- (i) PSI BLAST (ii) HSPs in BLAST (iii) NGS technologies and applications
- (iv) Offset vector in FASTA (v) Personalized genomics for Drug Discovery [CO-1, 2]

Q.2. Discuss all the challenges and perspectives of computational genomics towards life. (2)

[CO-1-3]

Q.3. What is fragment assembly? Explain its complications with an example of each. (3) [CO-2]

Q.4. Assemble following fragments to generate a candidate genome. Keep in mind that orientation is unknown to the assembler. Check for length and overlaps in the final output.

(3) [CO-2]

- ACGT
- CGCTGC
- CTAGCA
- CGTGC

Q.5. Provide a comparative analysis of three genomics types with their respective applications.

(2) [CO-1]