

JAYPEE UNIVERSITY OF INFORMATION TECHNOLOGY, WAKNAGHAT

TEST -3 EXAMINATION- 2025

B.Tech-VII Semester (BT)

COURSE CODE (CREDITS): 18B1WBI731(3)

MAX. MARKS: 25

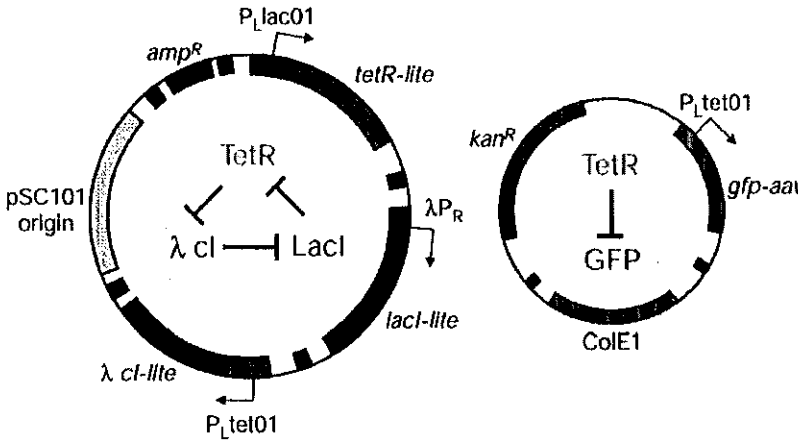
COURSE NAME: Computational Systems Biology

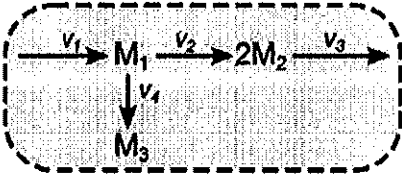
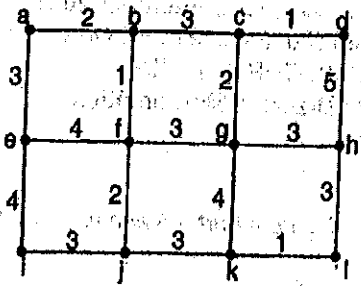
COURSE INSTRUCTORS: Dr. Raj Kumar

MAX. TIME: 2 Hours

Note: (a) All questions are compulsory.

(b) The candidate is allowed to make Suitable numeric assumptions wherever required for solving problems

| Q.No | Question | CO | Marks | | | | | | | | | | | | |
|--------------|--|--------|--------|-----|--------------|--------|---|-------------|-------|---|--------------|------|----|---|---|
| Q1 | Give a brief account of the various types of protein–protein interactions, and enumerate the methods used for identifying PPIs. | 5 | 3 | | | | | | | | | | | | |
| Q2 | The BioCyc database collection is a set of several pathway/genome databases (PGDBs). Provide a short summary of tasks involved in curating BioCyc databases? | 2 | 3 | | | | | | | | | | | | |
| Q3 | Explain the working of the given synthetic gene circuit:  | 3 | 5 | | | | | | | | | | | | |
| Q4 | Create a possible XML output for the following Vet. Data: <table border="1" data-bbox="309 1702 647 1859"><thead><tr><th>Person</th><th>Animal</th><th>Age</th></tr></thead><tbody><tr><td>Hafsah Downs</td><td>Cashew</td><td>2</td></tr><tr><td>Carrie Pope</td><td>Chase</td><td>1</td></tr><tr><td>Jim Chandler</td><td>Otis</td><td>20</td></tr></tbody></table> | Person | Animal | Age | Hafsah Downs | Cashew | 2 | Carrie Pope | Chase | 1 | Jim Chandler | Otis | 20 | 4 | 5 |
| Person | Animal | Age | | | | | | | | | | | | | |
| Hafsah Downs | Cashew | 2 | | | | | | | | | | | | | |
| Carrie Pope | Chase | 1 | | | | | | | | | | | | | |
| Jim Chandler | Otis | 20 | | | | | | | | | | | | | |

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|----|---|---|-----------------------|
| Q5 | <p>Given a simple metabolic network with four reactions and three metabolites, construct the stoichiometric matrix and set up the FBA optimization problem (objective function + constraints).</p> <p style="text-align: center;">Metabolic network</p>  | | 5 |
| Q6 | <p>Use Kruskal's Algorithm to find a minimum spanning tree in the weighted graph given below:</p>  | 5 | 4 |
| Q7 | <p>Systems biology is a scientific approach that studies complex biological systems. Explain the following in the context of systems biology.</p> <ol style="list-style-type: none"> Reductionist Vs. systems approach Emergent properties Model organisms Kinetic Modelling Systems biology and synthetic biology | 1 | 2×5 $=10$ |