| Name: | Reg #: | Section: |
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## **National University of Computer and Emerging Sciences, Lahore Campus**

Course: Bioinformatics Course Code: CS4054

Section: BCS-8A **Program: BS(Computer Science) Semester: Spring 2025** 

**Duration: 30 Minutes Total Marks: 10** Paper Date: 24-April-2025

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Exam: Quiz 3

Q1 Consider the following gene expression distance matrix from a microarray experiment for 5 genes. A biologist is trying to find out whether these 5 genes can be separated into clusters based on their behavior in the experimental conditions.

|                   | Gene <sub>1</sub> | Gene <sub>2</sub> | Gene <sub>3</sub> | Gene <sub>4</sub> | Gene₅ |
|-------------------|-------------------|-------------------|-------------------|-------------------|-------|
| Gene <sub>1</sub> | 1.00              | 0.90              | 0.10              | 0.65              | 0.20  |
| Gene <sub>2</sub> | 0.90              | 1.00              | 0.70              | 0.60              | 0.50  |
| Gene <sub>3</sub> | 0.10              | 0.70              | 1.00              | 0.40              | 0.30  |
| Gene <sub>4</sub> | 0.65              | 0.60              | 0.40              | 1.00              | 0.80  |
| Gene <sub>5</sub> | 0.20              | 0.50              | 0.30              | 0.80              | 1.00  |

Use average link to update your distance matrix. Indicate the data points belonging to each cluster. Create a dendrogram to visualize the results. If the biologist wants to group the genes into two clusters, identify the genes belonging to each cluster. [10 marks]