## **UPGMA Exercise Solution**

	Α	В	С	D
Α	0	3	10	10
В	3	0	2	12
С	10	2	0	4
D	10	12	4	0

- 1. Run the UPGMA Algorithm on distance matrix D, using centroid based (average) linkage when computing inter-cluster distances, showing intermediate steps. Parts (a) and (c) are worth 5 points each. Parts (b), (d), and (e) are worth 10 points each.
  - a. Which pair of nodes is Merged first? B and C
  - **b.** What is the resulting inter-cluster distance matrix?

	ВС	Α	D
ВС	0	6.5	8
Α	6.5	0	10
D	8	10	0

- c. Which pair of nodes or clusters is Merged next? (BC) and A
- **d.** What is the resulting inter-cluster distance matrix?

	ABC	D
ABC	0	26/3 = 8.66667
D	26/3 = 8.66667	0

e. What is the final phylogenetic tree based on the above UPGMA algorithm run?

(((B C) A) D)