Figure 4.10a. Clustering of significantly differential expressed orthologous genes using k-means clustering method according to the gene expression profiles in Bud, Flower and Peg.

Figure 4.10b. Clustering of significantly differential expressed orthologous genes using k-means clustering method according to the gene expression profiles in S1, S2 and S3.
Figure 4.10c. Clustering of significantly differential expressed orthologous genes using k-means clustering method according to the gene expression profiles in UL1, UL2 and UL3.

Figure 4.10d. Clustering of significantly differential expressed orthologous genes using k-means clustering method according to the gene expression profiles in LL1, LL2 and LL3.