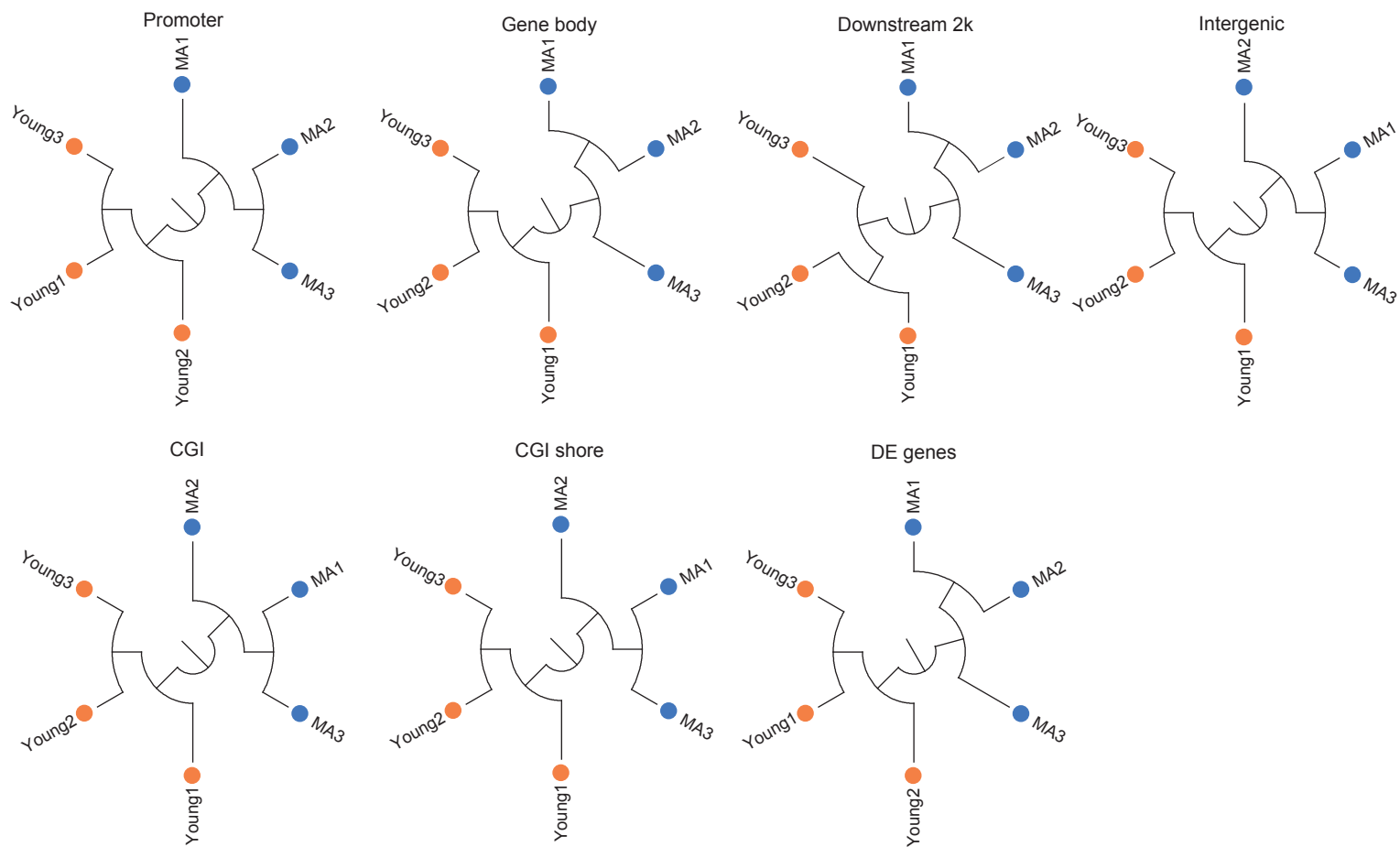




Additional file 8: Hierarchical clustering of samples using DMRs in various genomic elements and differentially expressed (DE) mRNAs. Clustering was performed using MultiExperiment Viewer software. The distance metric applied for clustering was Pearson correlation across samples.