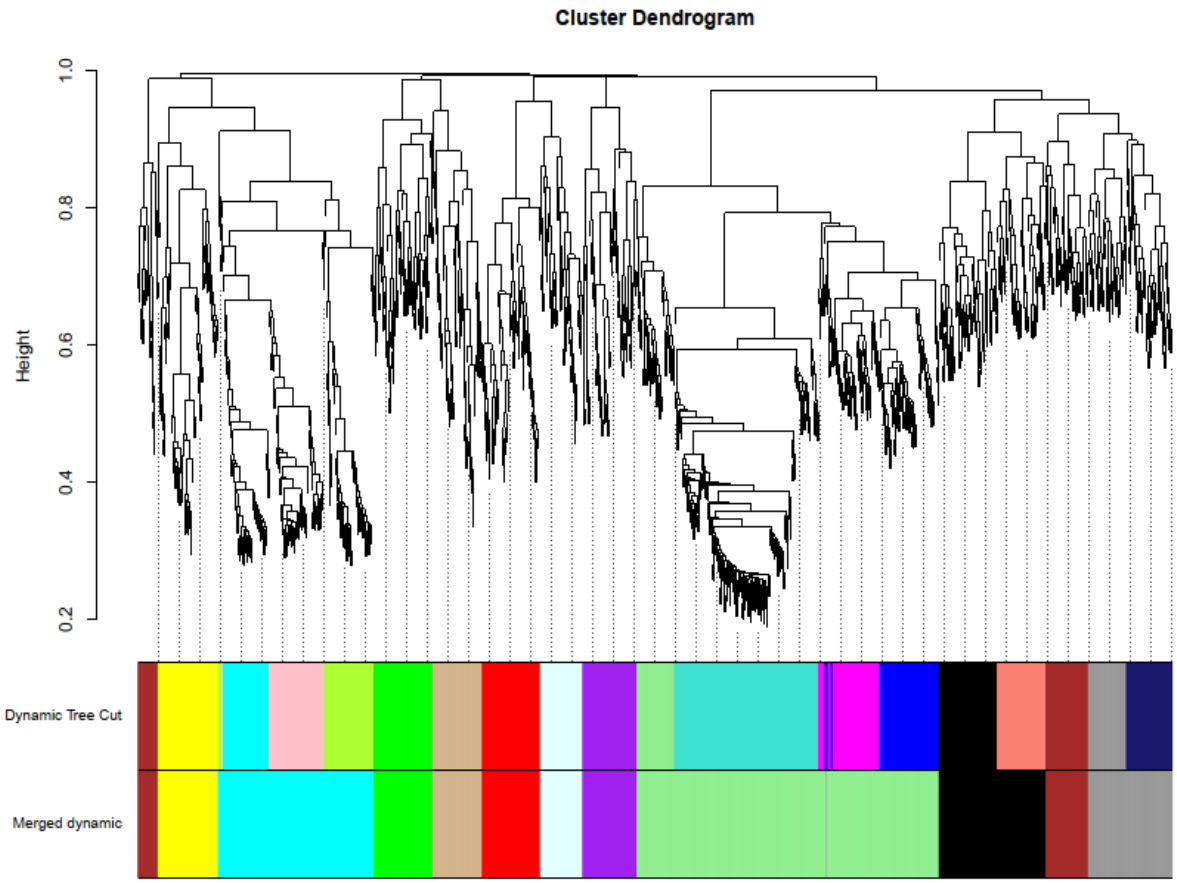


Supplementary Figure 6. Hierarchical cluster dendrogram showing correlation modules identified by WGCNA with minModuleSize = 50 and MEDissThres = 0.09. Each colored row represents a module containing a group of highly connected genes.