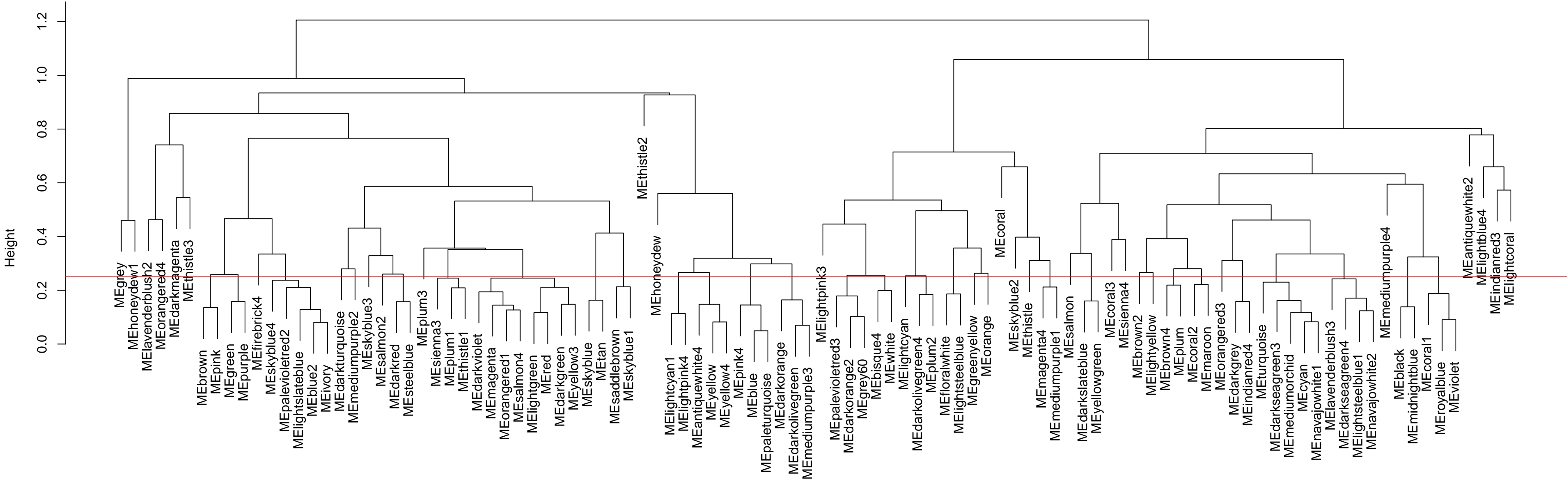


Clustering of Module Eigengenes



as.dist(MEDiss)
hclust (*, "average")