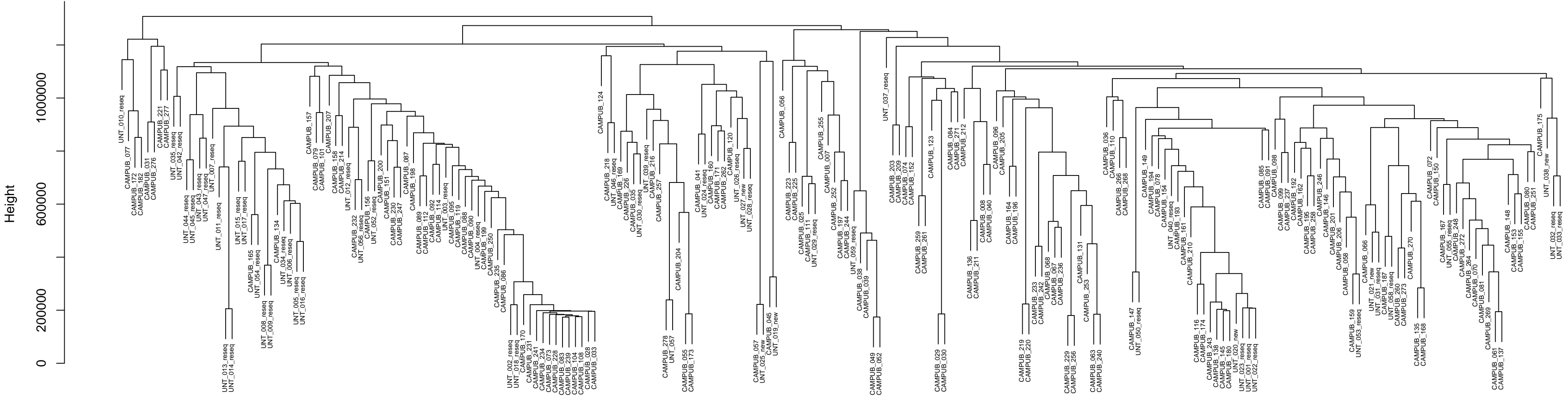


Cluster Dendrogram



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