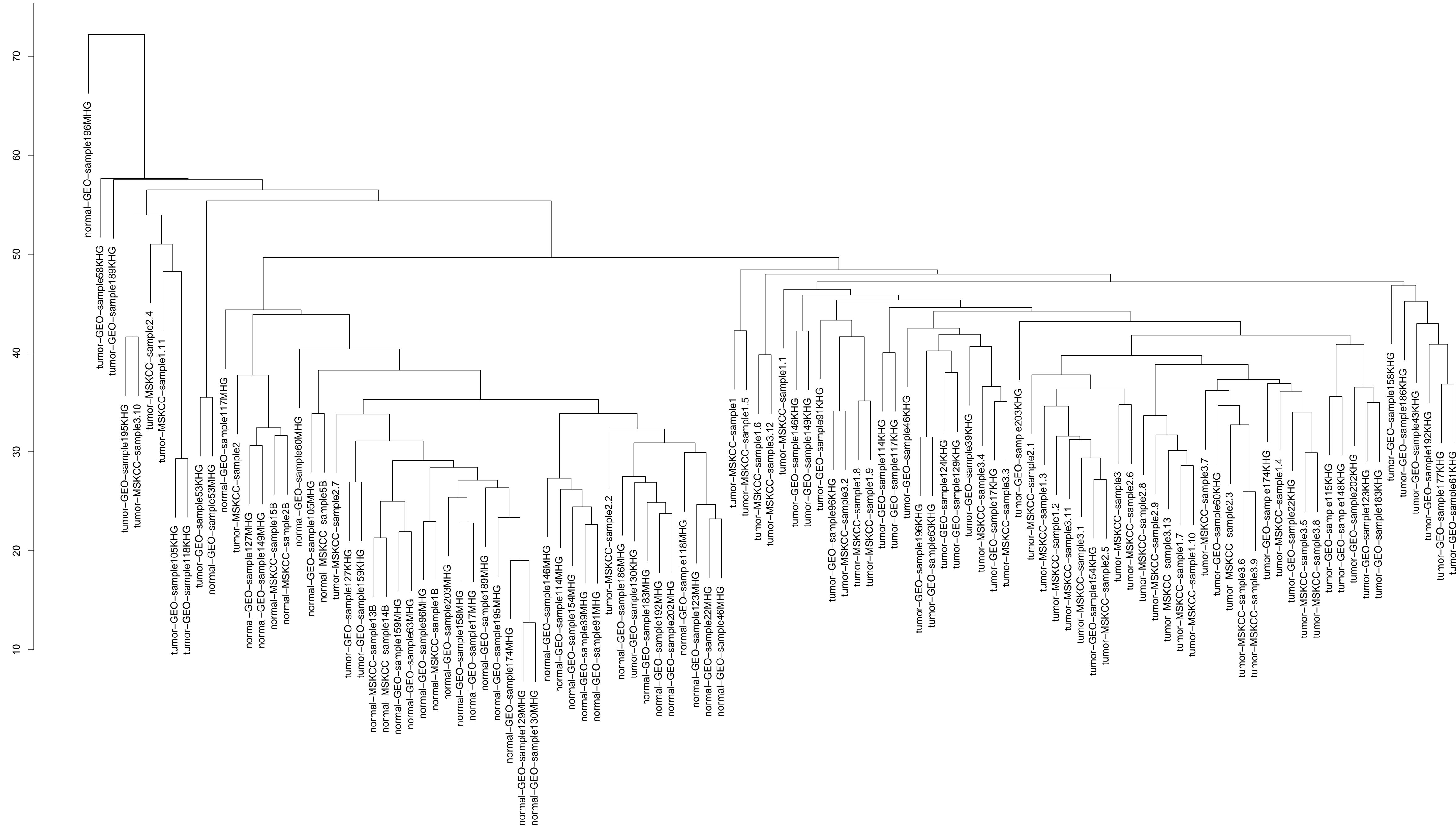


Sample relations based on 11413 genes with sd/mean > 0.1



Sample
hclust (*, "average")