DENDOGRAM

Height
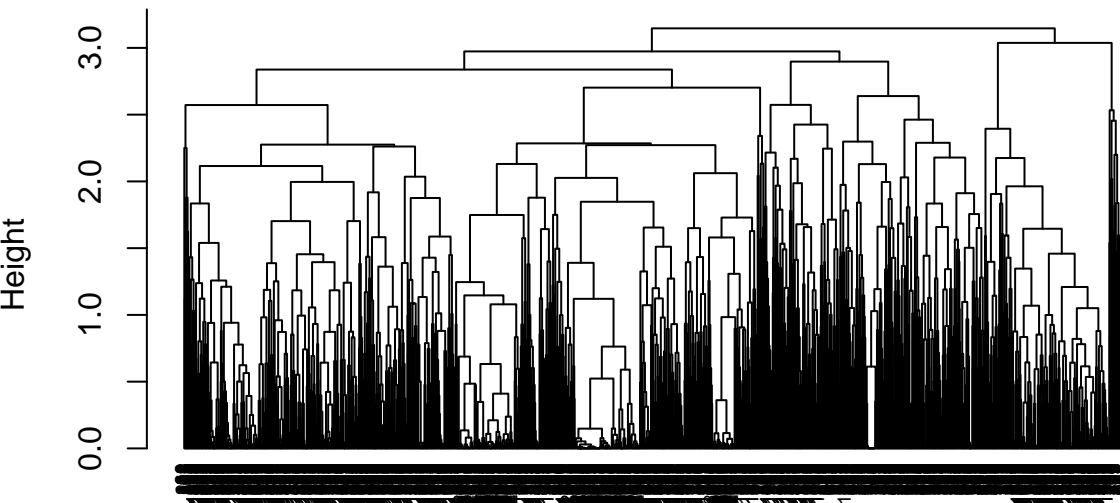
dist(obesity.cluster[, 1:15], method = "minkowski")
hclust (*, "complete")