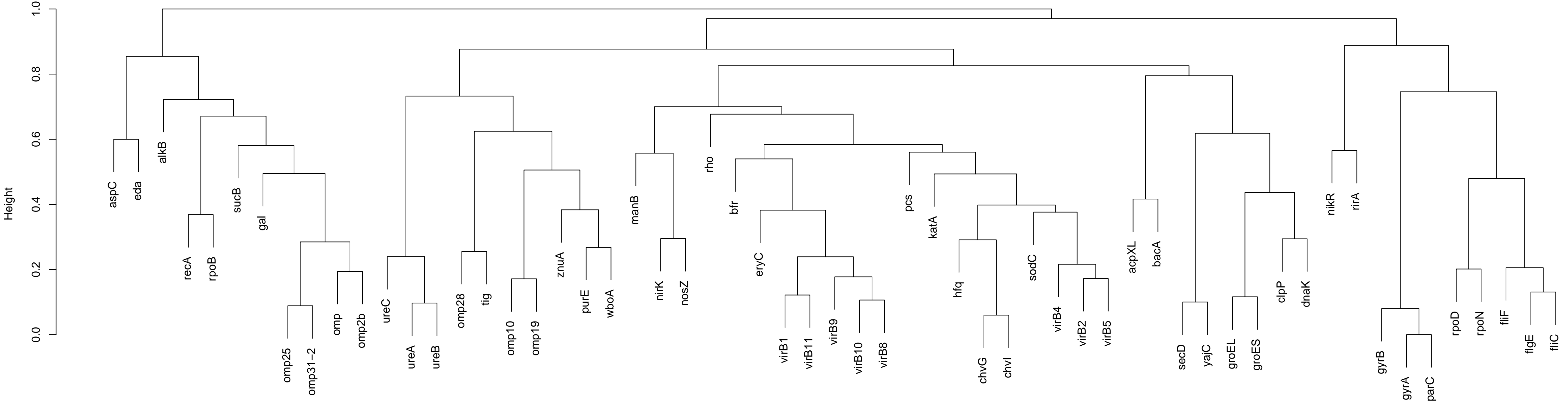


Brucella cluster result (Sqrt\_Cosine\_coefficient)



as.dist(t(a))  
hclust (\*, "complete")