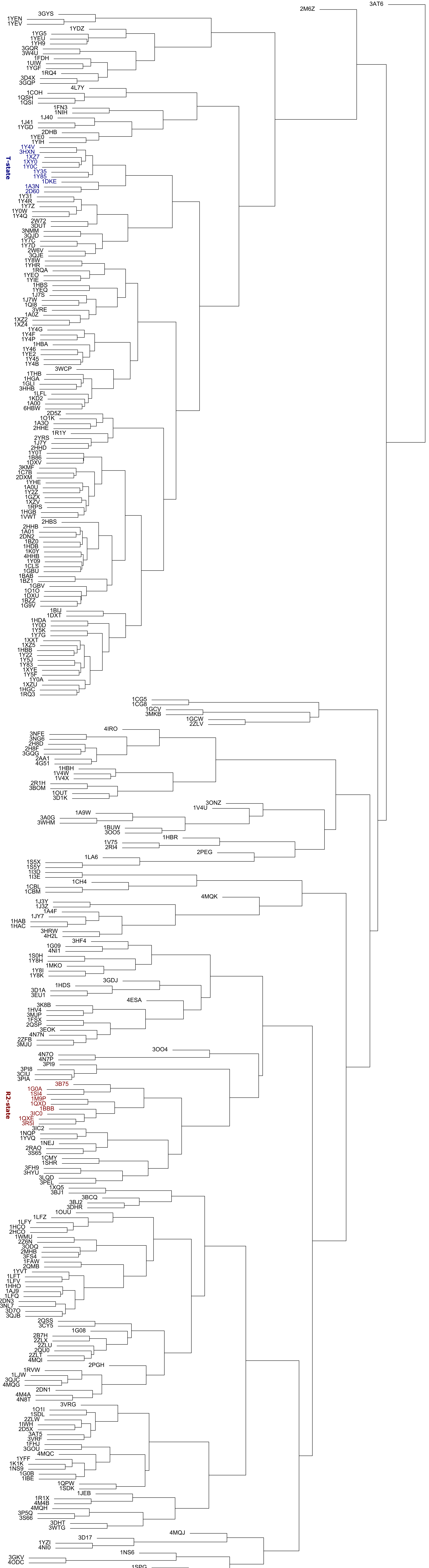


Height



T-state

R2-state

Hemoglobin (dataset2) DSS-based dendrogram

as.dist(dsim)  
clus1("complete")