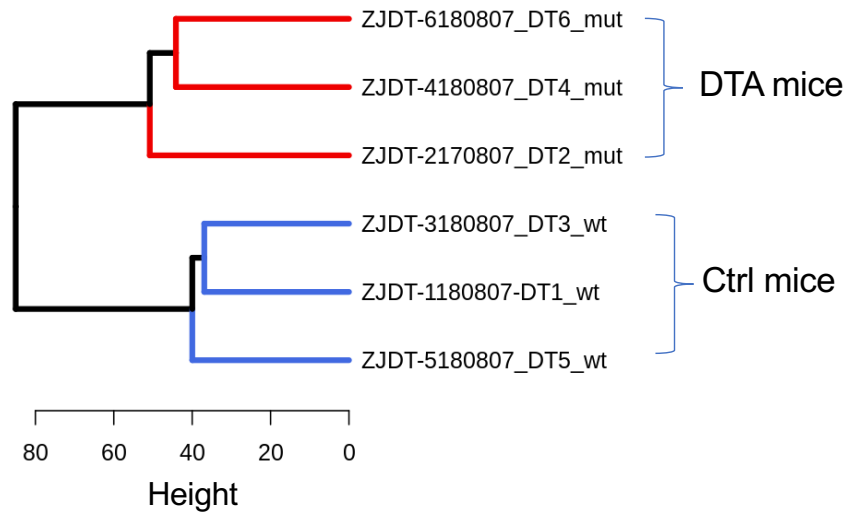


Supplementary Figure 1



Supplementary Figure 1. Cluster analysis of transcriptome data. Hierarchical clustering shows that samples fall into two clear groups, i.e. DTA mutant mice (in red) and littermate control mice (in blue).