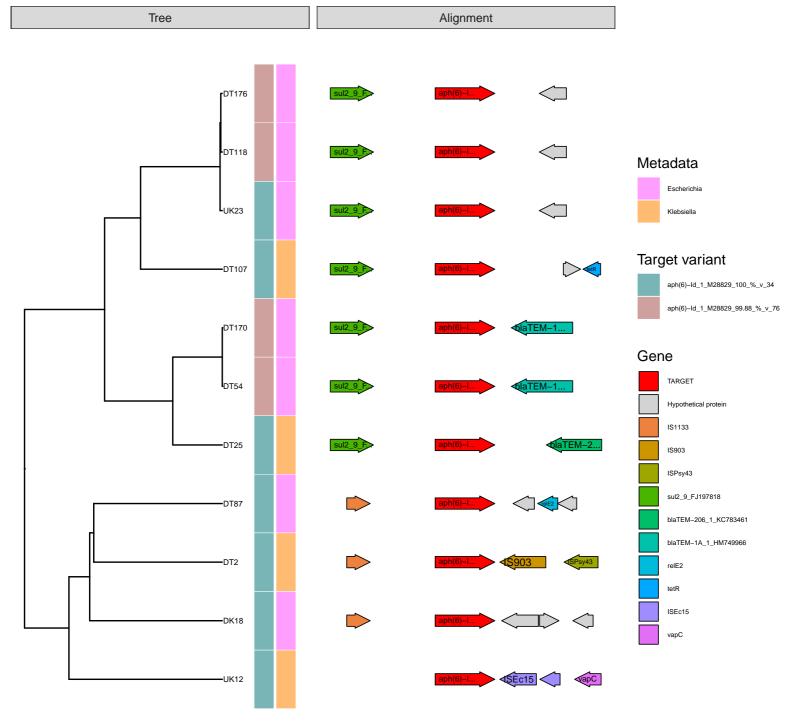
Cluster 33_aph_6__Id_1_M28829 – distance tree based on target and flanking region sequences



VARIANT METADATA