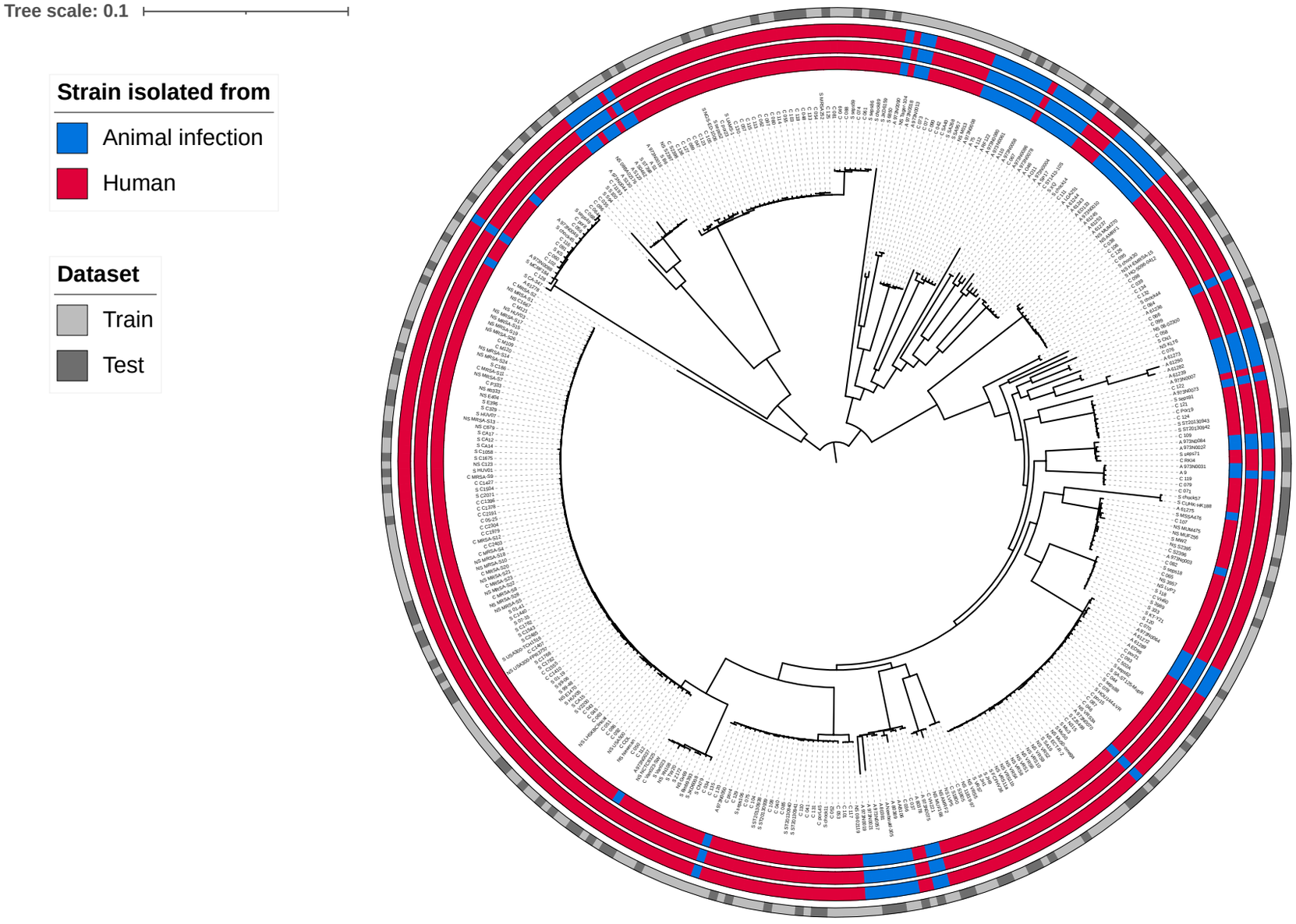

Figure S10: Predictions obtained using random forest to discriminate animal and human isolates plotted on phylogenetic reconstruction of multiclonal Staphylococcus aureus. The first outer circle represents the actual phentotype. The second and third outer circle indicates prediction using CDS or IGRs, respectively. The fourth outer circle refer to strains used for the training or test sets, respectively.