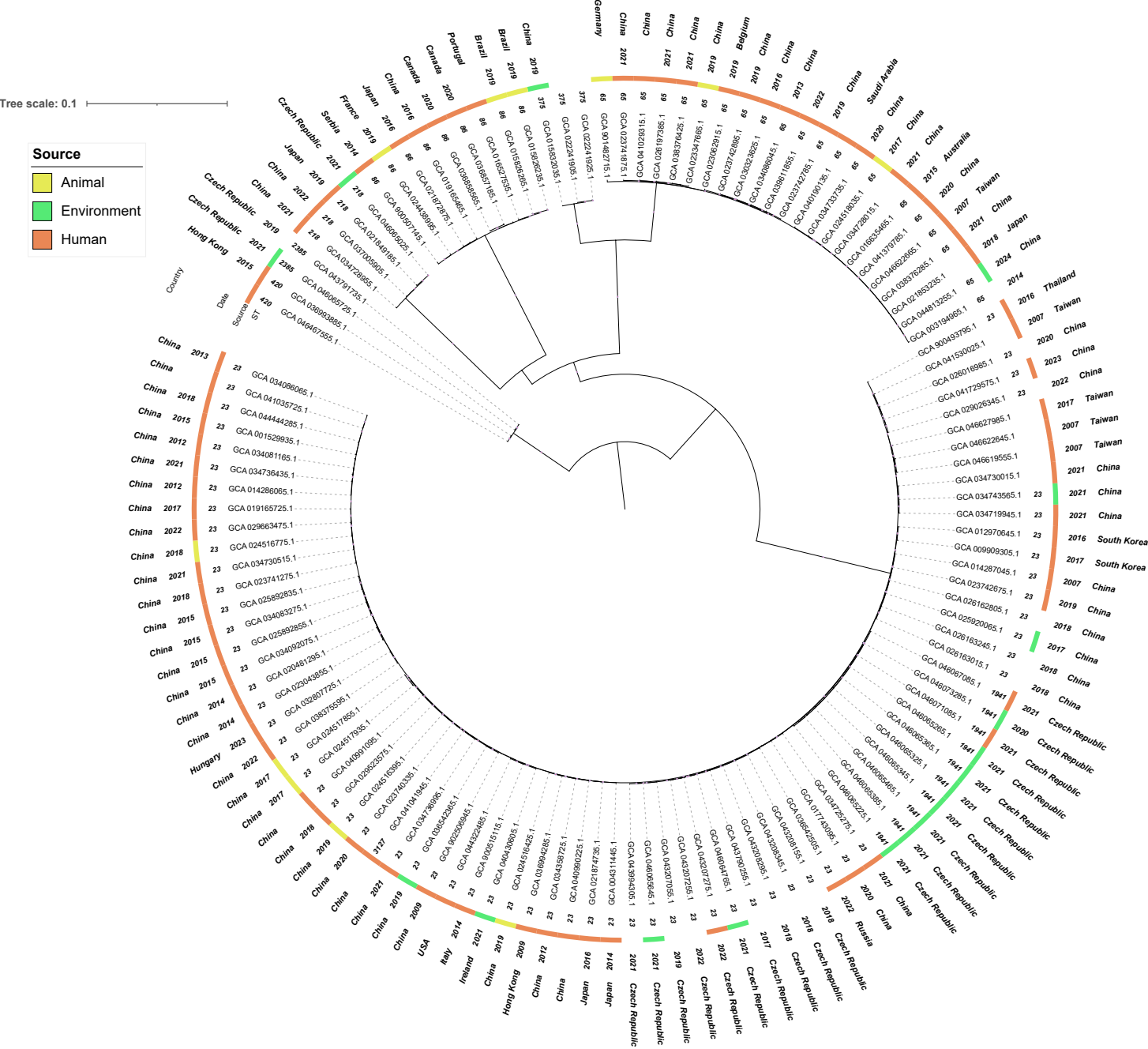


Figure S1. Phylogenomic relationship of hvKp genomes across One Health compartments. The maximum-likelihood tree illustrates the genetic relatedness of hvKp isolates recovered from human, animal, and environmental sources. Colored strips indicate the source for each genome. The tree branch lengths were ignored in this tree. Red circles on branches represent >70% bootstrap.