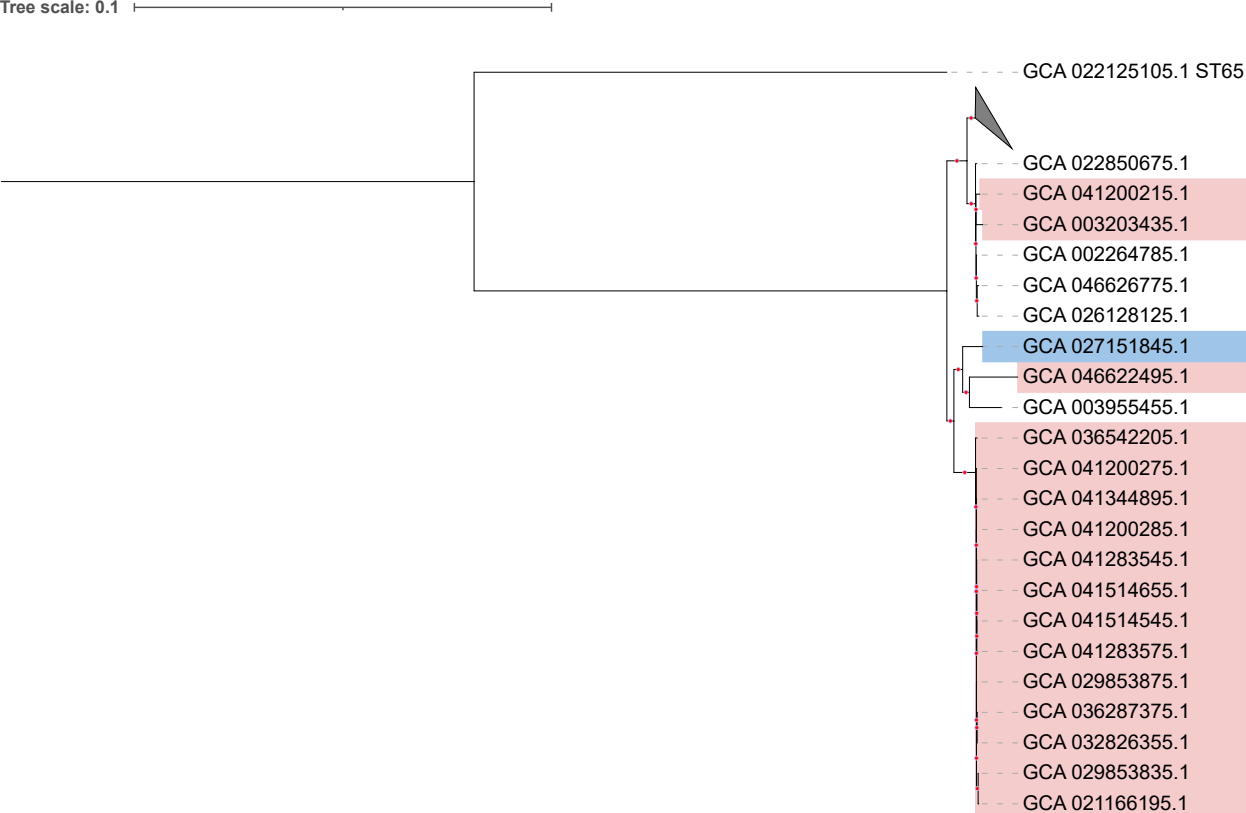


Figure S2. Phylogeny of hvKp ST11 genomes. Distinct MDR lineages are observed, with most genomes from China (collapsed node) clustering separately from those from Taiwan (red background) and Italy (blue background). A ST65 genome was included as the outgroup.