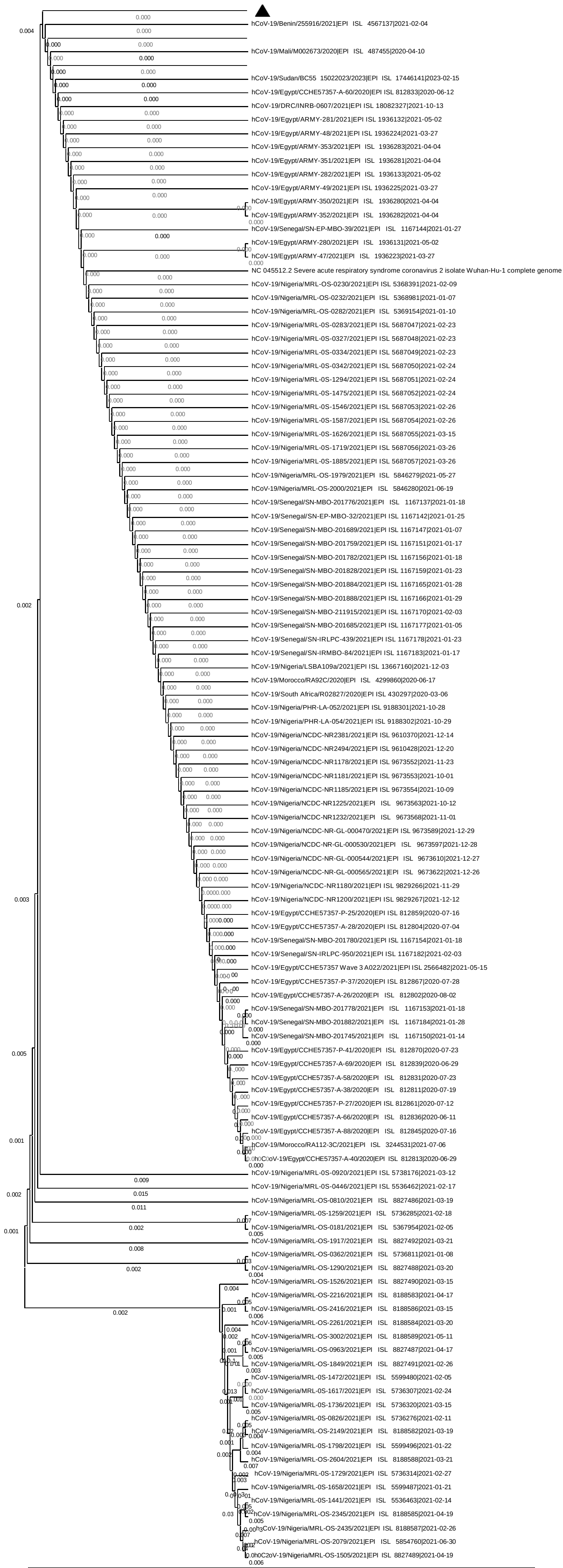

Figure S1. Phylogenetics of strains closely related to the study's 45 SARS-CoV-2 in Fig. 6a