

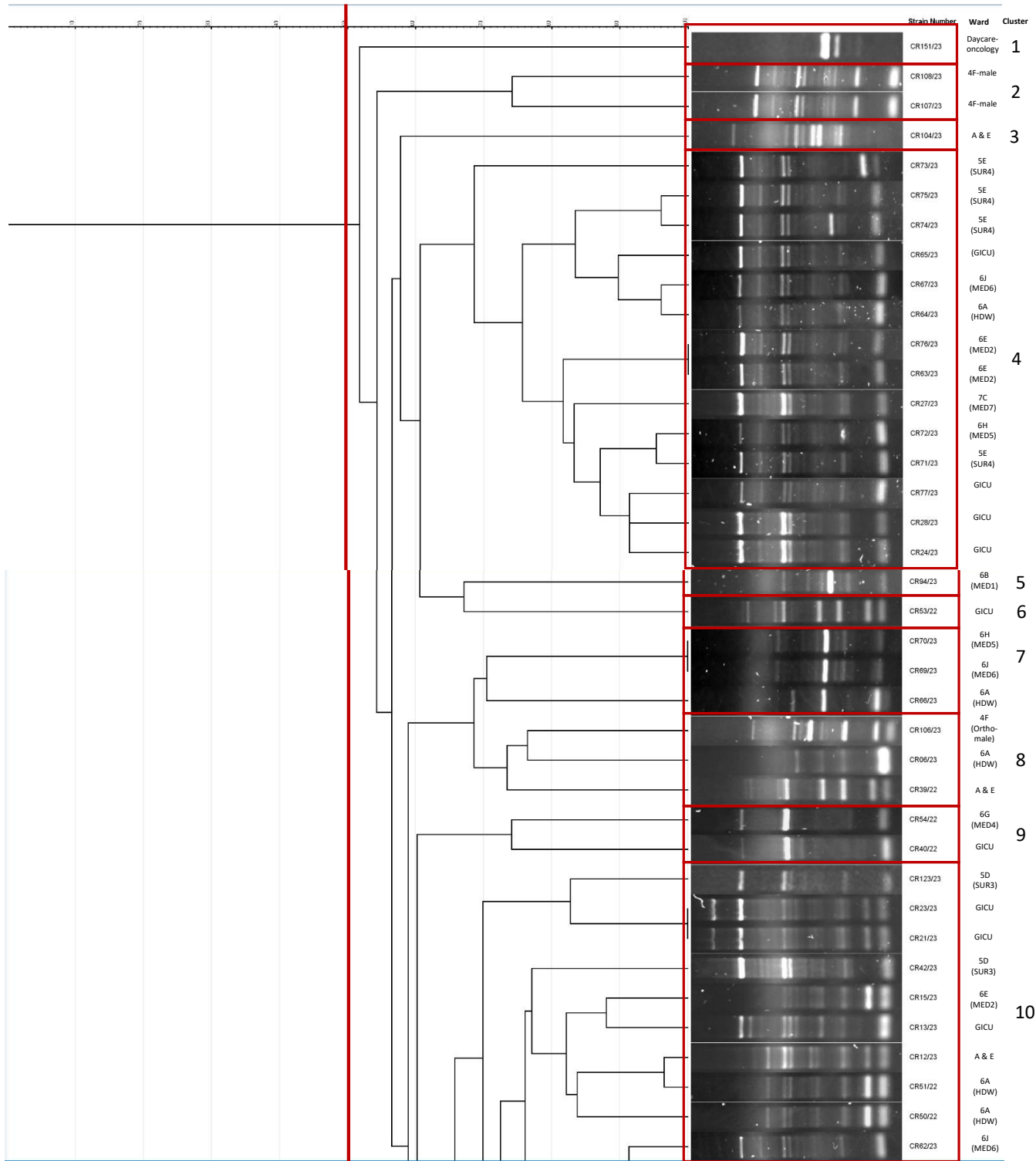


Figure S2. Clustering of enterobacterial repetitive intergenic consensus polymerase chain reaction genotypes for study isolates. A total of 24 clusters were identified as having > 50% similarity. Strains in clusters 10, 11, 22, and 24 appeared to have closer similarity (> 60%). No cluster was distinct to a single ward, signifying transmission of various carbapenem-resistant *Klebsiella pneumoniae* genotypes across wards of the hospital. (Continued on the next page.)

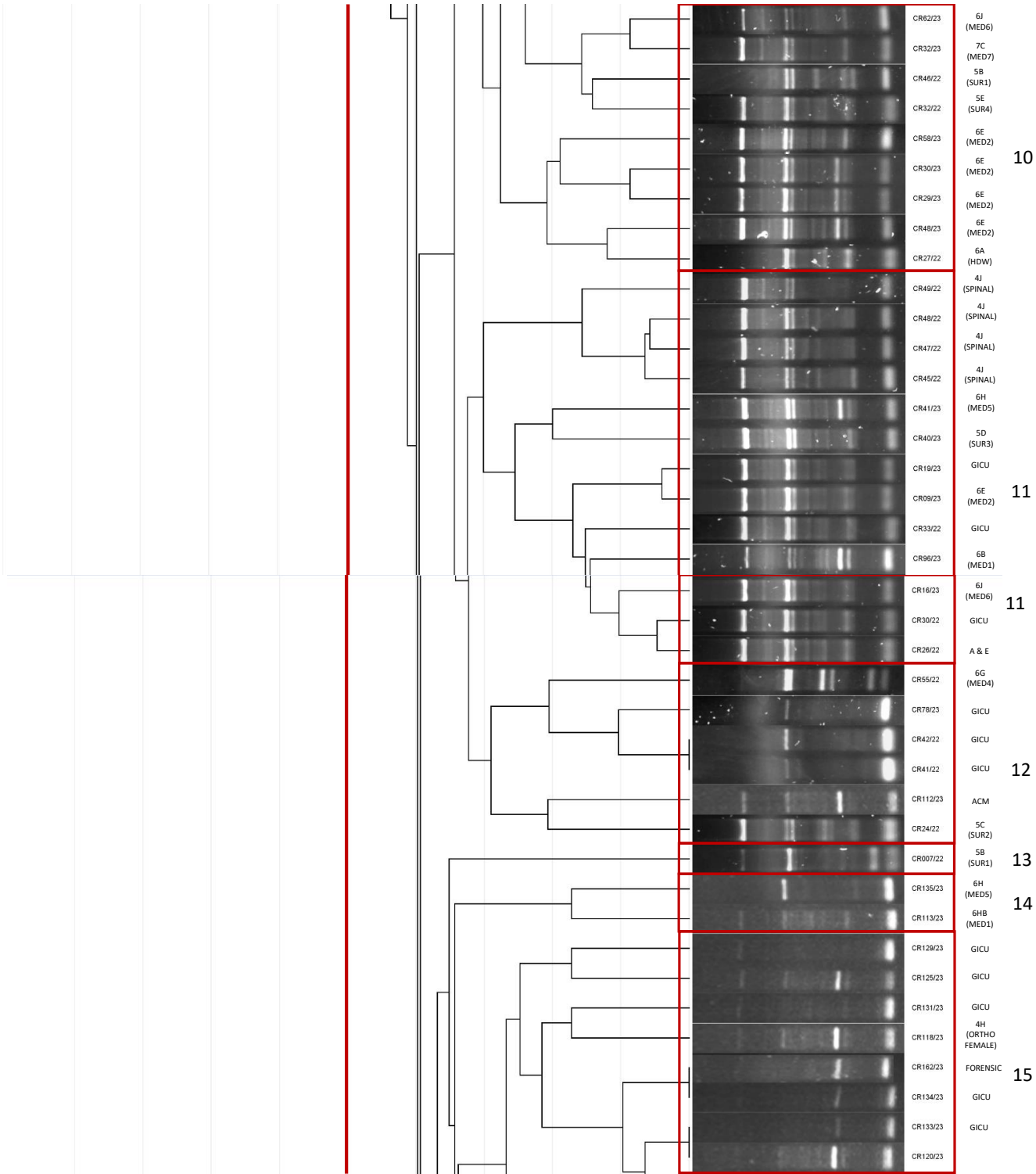


Figure S2. (Continued; caption shown on previous page).

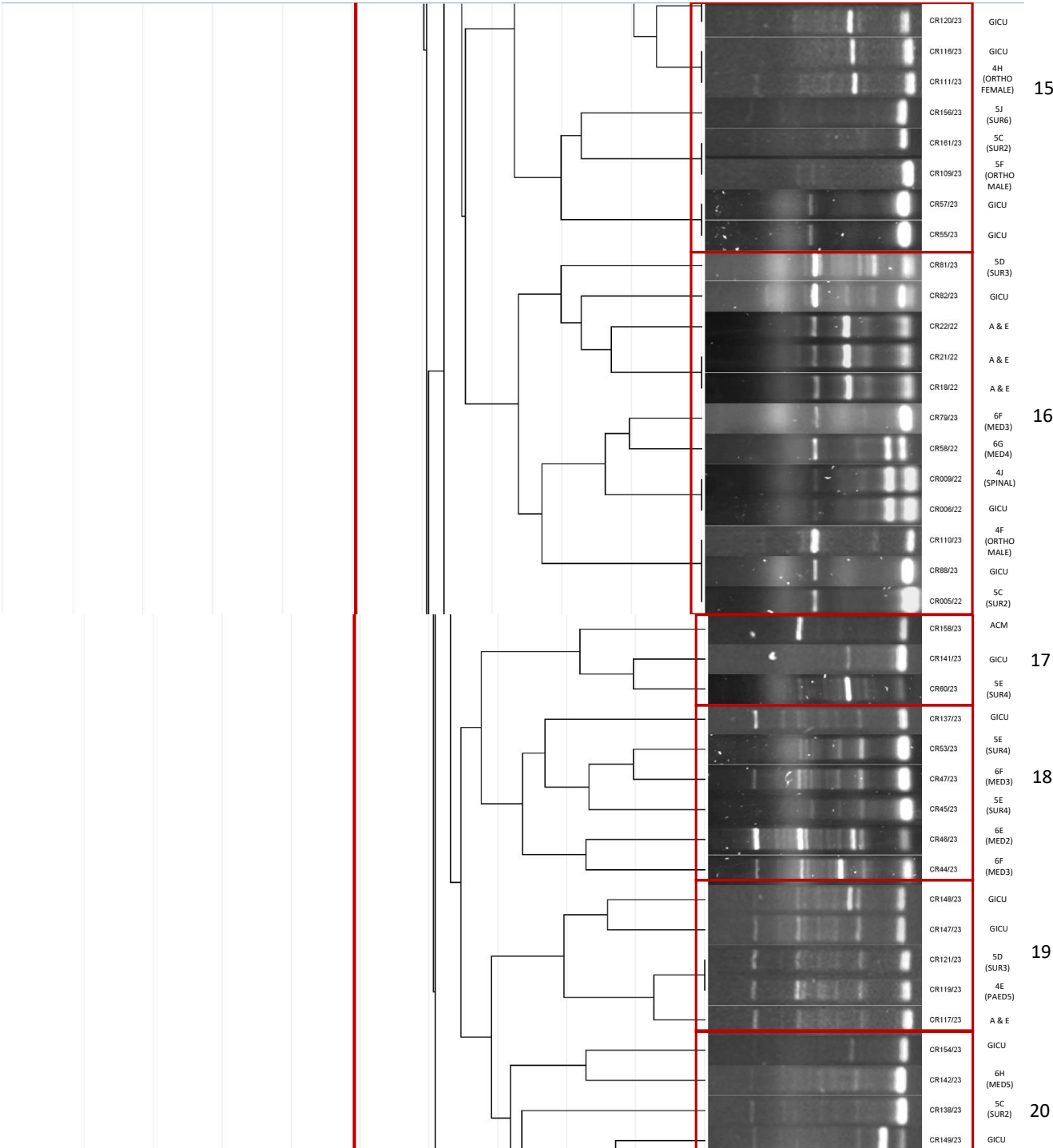


Figure S2. (Continued; caption shown on previous page).

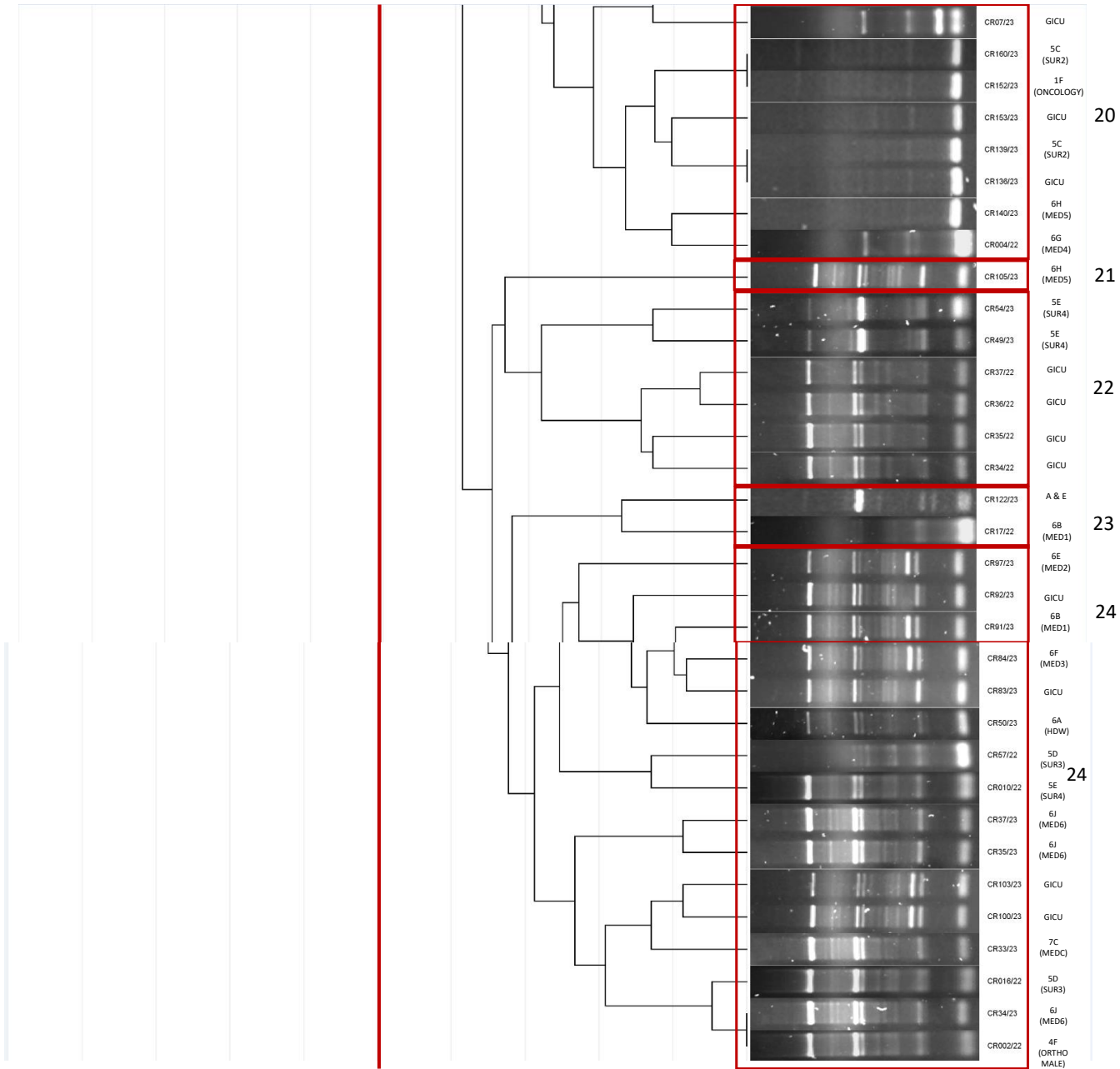


Figure S2. (Continued; caption shown on previous page).