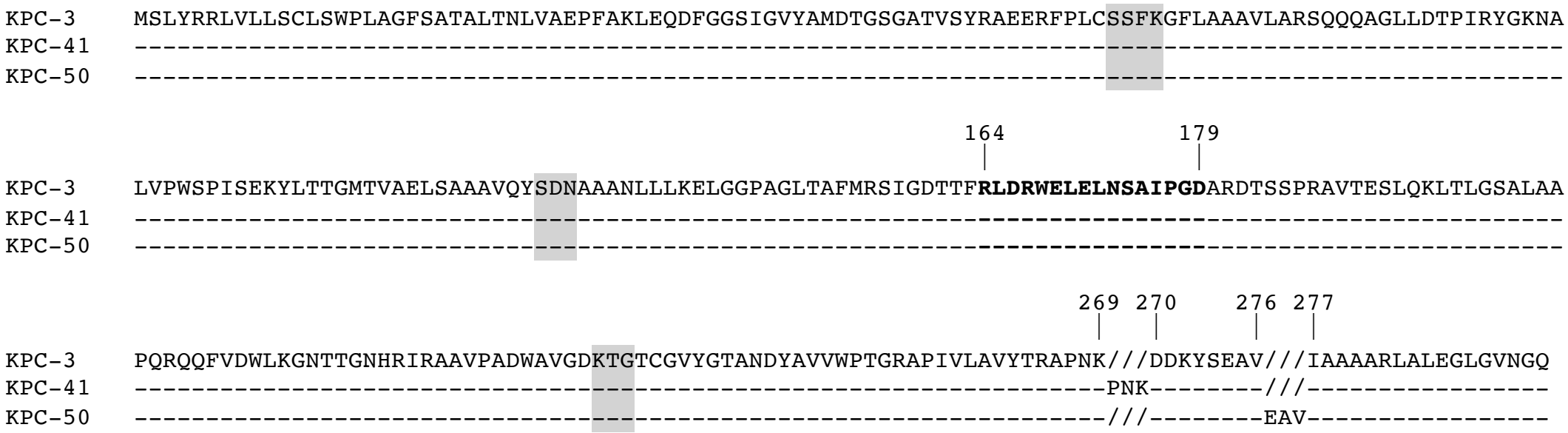Figure S1. Alignment of the KPC-50 amino acid sequence in comparison with the KPC-3 ancestor and the KPC-41 variant conferring resistance to CZA. Dashes indicate identical amino acids between sequences, and slashes indicate absence of amino acids at the corresponding positions. Shaded amino acids indicate conserved motifs found within class A ß-lactamases. Boldened amino acids indicate the omega-loop of KPC-type enzymes.