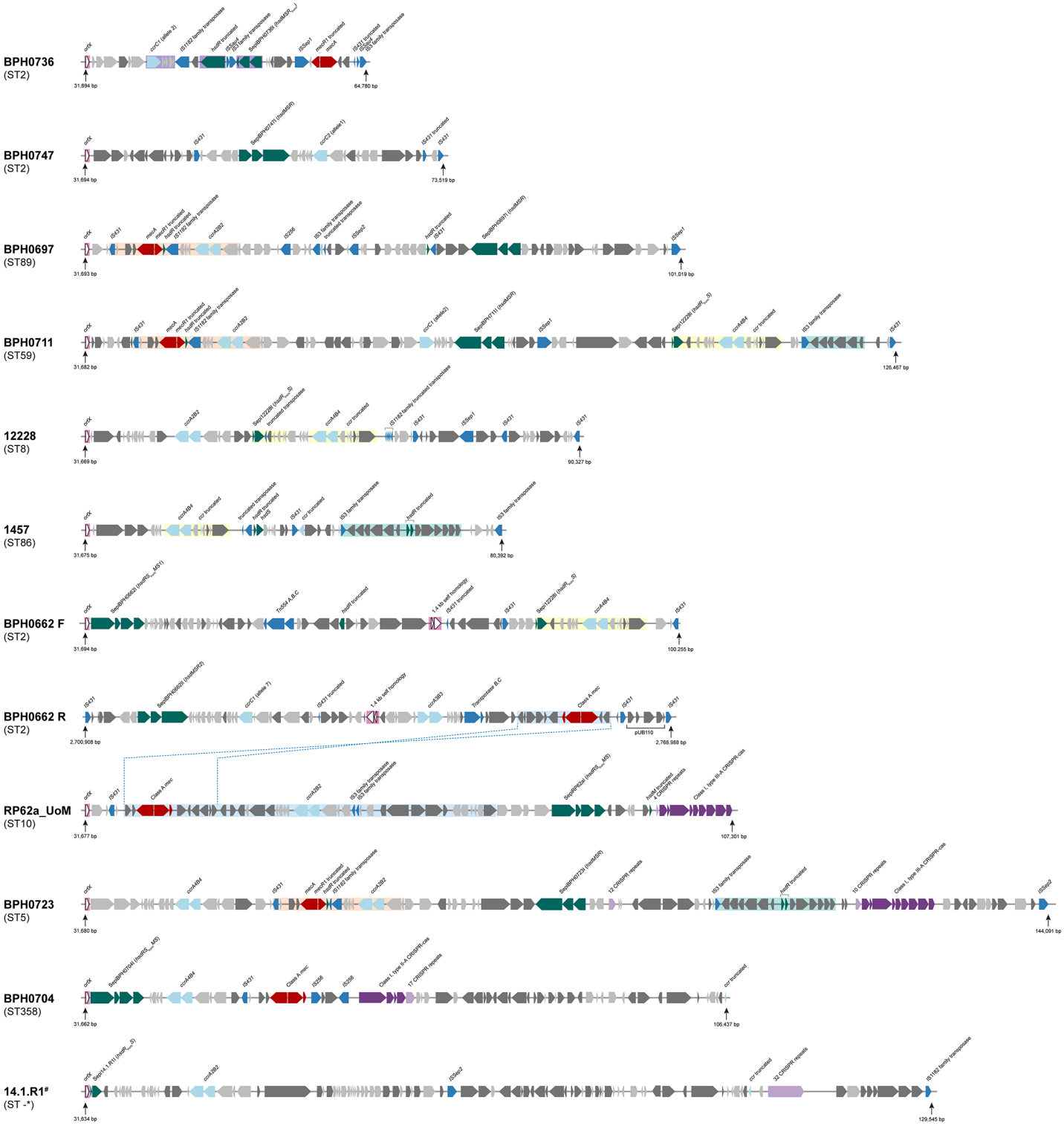
A. *Staphylococcus epidermidis* complete genomes type I RM systems



B. *Staphylococcus aureus* imported type I RM systems

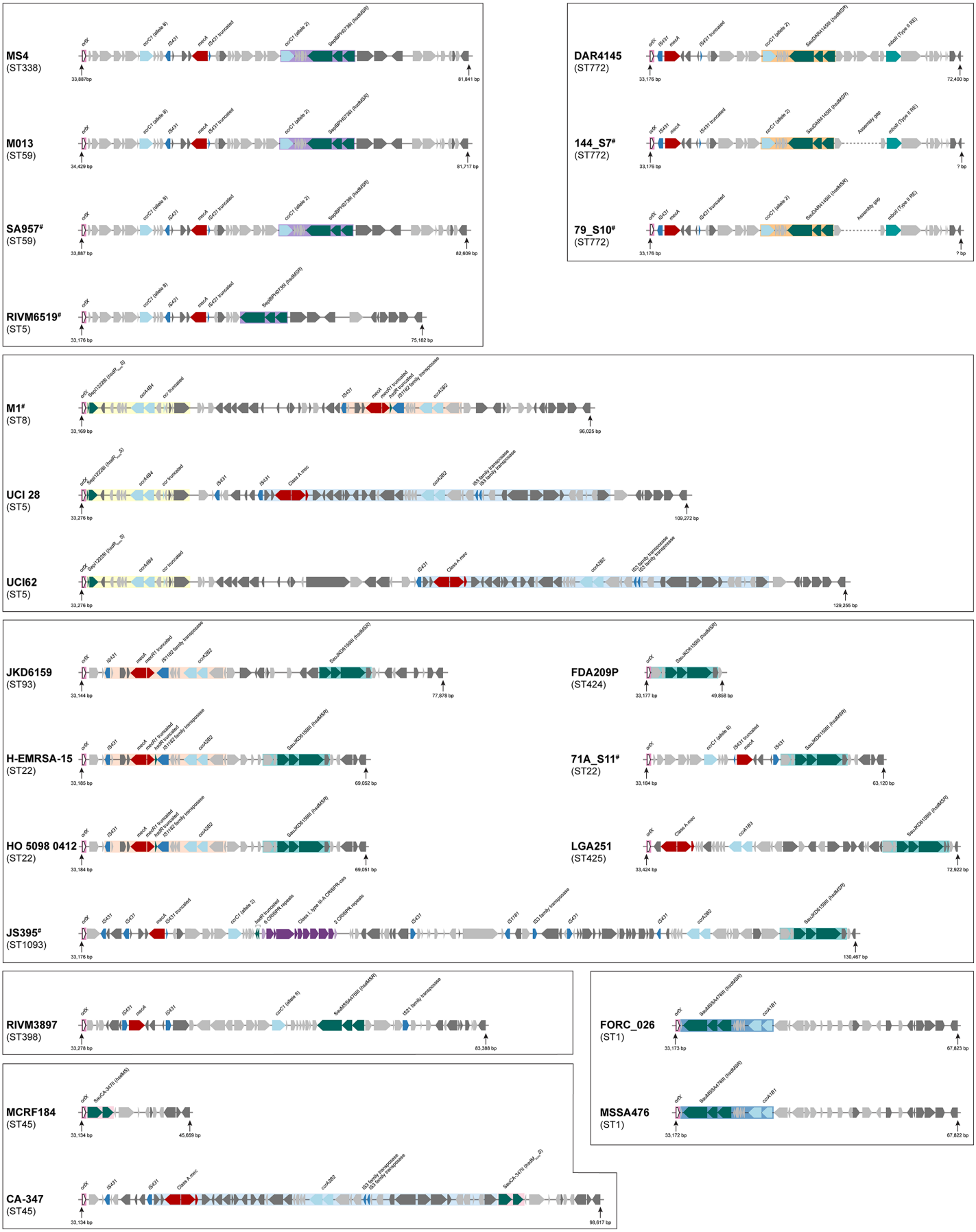


Figure S1. A hypothesised role for cassette chromosome recombinase (*ccr*) in the mobilisation of *S. epidermidis* and *S. aureus* type I restriction modification systems. A. Complete *S. epidermidis* genomes with type I RM systems. **B.** *S. aureus* genomes with imported type I RM systems. Genomes are orientated forwards starting at *dnaA*. [#]NCBI uploaded genome does not start at *dnaA*. *Strain not classifiable by existing MLST scheme.