



**Fig S5. *cmrRST* and upstream regulatory sequences are highly conserved across *C. difficile* strains and ribotypes.** Heat map showing percent sequence identity of 71 sequenced *C. difficile* strains in the NCBI database compared to the R20291 reference sequence (NCBI accession number FN545816.1). If the strain has been ribotyped, the ribotype is indicated on the left. For the *cmr* switch, both orientations were used as the reference and the highest percent sequence identity is shown. The following nucleotides were used as the reference sequence from R20291: riboswitch, 3,736,512-3,736,863; *cmr* switch, 3,735,969-3,736,513; *cmrRST*, 3,732,474-3,735,968.