SDC2: Supplementary Figures for Detection of plasmid-mediated tigecycline resistance gene *tet*(X4) in a *Salmonella enterica* serovar Llandoff isolate

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# YW, FL and XX contributed equally to this work.

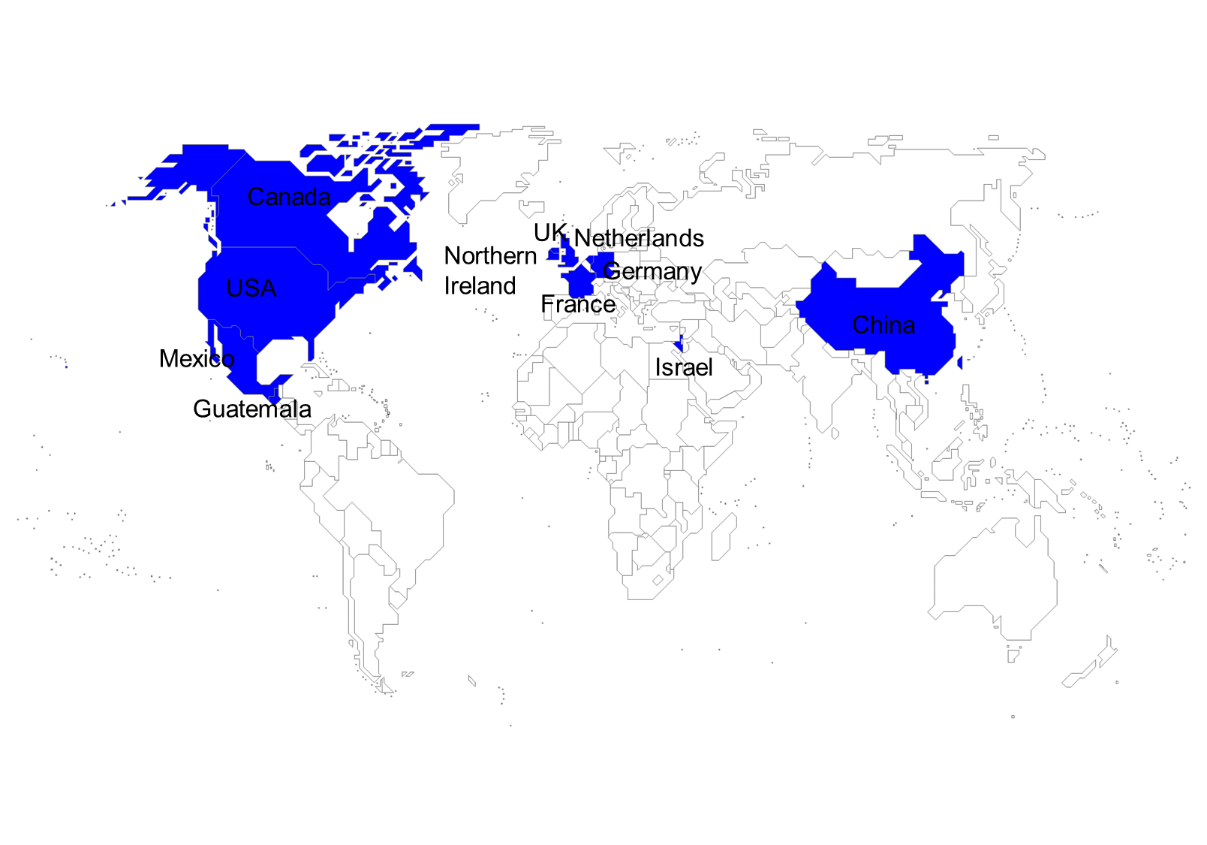


Figure S1. Source location of *Salmonella enterica* serovar Llandoff strain genomes used in this study. The blue color indicates the presence of this serovar.

E:\文件资料\微生物所\细菌单菌株-20190621\manuscript-202008\20210807-IMD--Infectious Microbes & Disease\Figure S2 The phylogenomic relationship among the Salmonella Llandoff isolates-20210830-01.tif

Figure S2. The phylogenomic relationship among 20 *Salmonella enterica* serovar Llandoff strains. The 20 *S.* Llandoff isolates were distributed in 11 countries on four continents and grouped into different sublineages. SH16G3606 was closer to the *S.* Llandoff strains in Guatemala, the USA, and Mexico in the evolutionary tree. “100” represent bootstrap value and “0.05” represents distance scale.

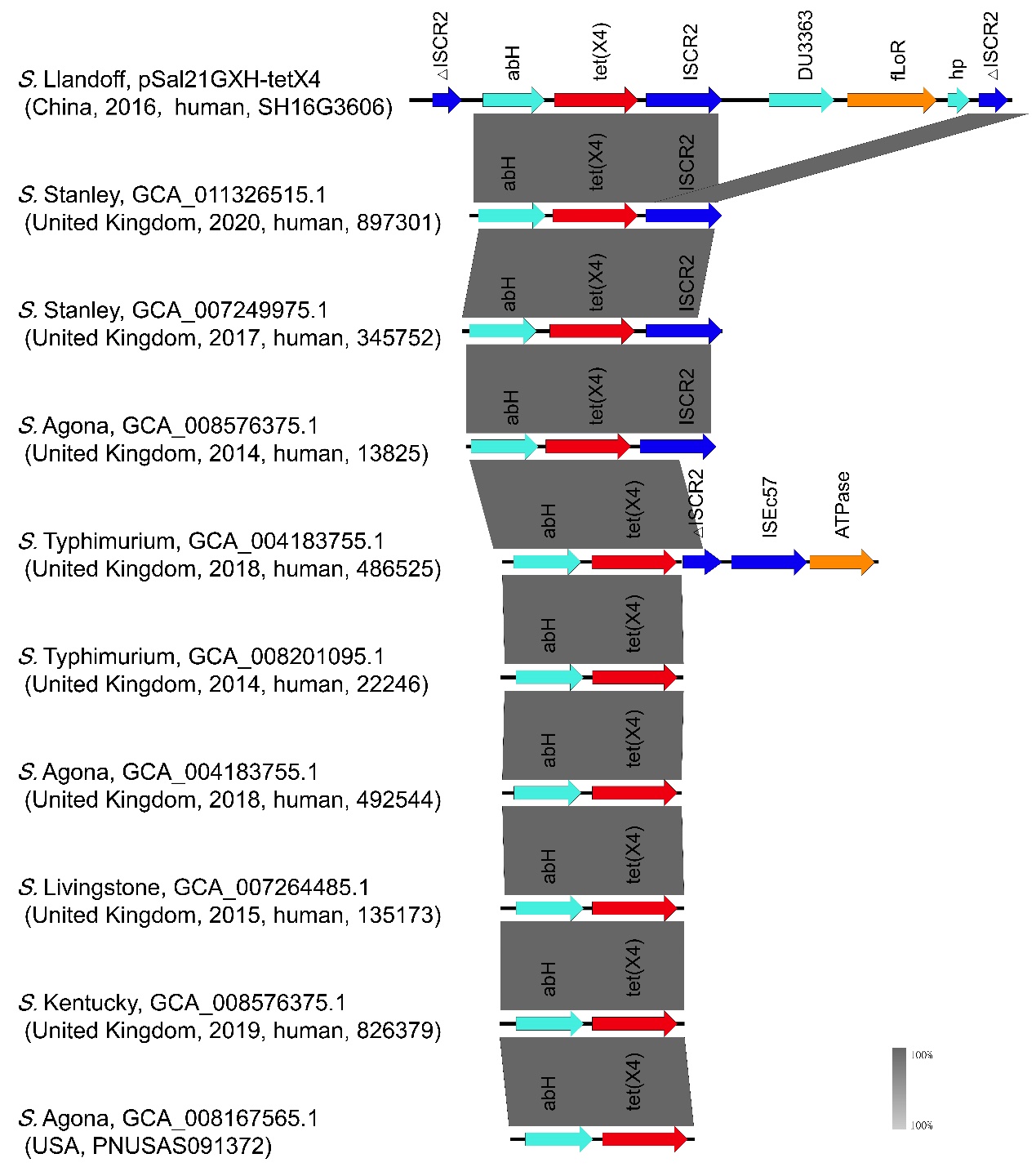


Figure S3. Comparison analysis of *tet*(X4)-positive *Salmonella enterica* contigs.