

Description of additional supplementary files

Title: Supplementary Data 1.

Description: Excel file containing tabs for 1) Genome information used in the study, 2) Genome comparison showing the similarity between flight isolates 3) Intact prophage clusters, 4) Long-read assembly prophage mapping, 5) Complete functional annotation of the prophage-encoded proteins, 6) Functional annotation of prophage-encoded proteins that are non-redundant with host genome functions, 7) AMR genes corroborated with ABRicate, 8) NCBI Accessions for Terrestrial genomes, 9) sequence information for all prophage regions.