

Supplementary Table 4. Clonal population distribution of swine ESBL-EC isolates for each MLST sequence type.

CC – Phylogenetic group (No. of isolates)	ST - Phylogenetic group	No. of isolates in cluster					Total no.
		1	2	3	4	5	
CC101-B1 (37)	ST101-B1	10	19			2	31
	ST5229-B1		3			1	4
	ST5696-B1	1					1
	ST4014-B1					1	1
CC10-A (13)	ST10-A				3		3
	ST215-A				2		2
	ST48-A				3		3
	ST617-A				4		4
	ST218-A				1		1
CC23-A (4)	ST23-A				2		2
	ST410-A				2		2
CC648-F (4)	ST648-F			4			4
CC86-B1 (4)	ST641-B1		4				4
CC205-B1 (1)	ST205-B1					1	1
CC20-B1 (1)	ST376-B1	1					1
CC12-B2 (1)	ST12-B2			1			1
Total	ST457-F			22			22
	ST75-B1	15	1			1	17
	ST224-B1					11	11
	ST2628-B1	5				1	6
	ST3944-A				3		3
	ST3076-B1					3	3
	ST744-A				2		2
	ST1642-B1					2	2
	ST3285-B1	1					1
	ST1011-E					1	1
	ST7203-A				1		1
	ST953-A	1					1
	ST767-B1					1	1
	NT-B1		1				1
	NT-A				1		1
	Total		34	28	27	24	25

CC, Clonal complex; ST, Sequence type; NT, non-typable.