Antimicrobial resistance profiles and genetic typing of *Salmonella* serovars from chicken embryos in China

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Supplementary Figures

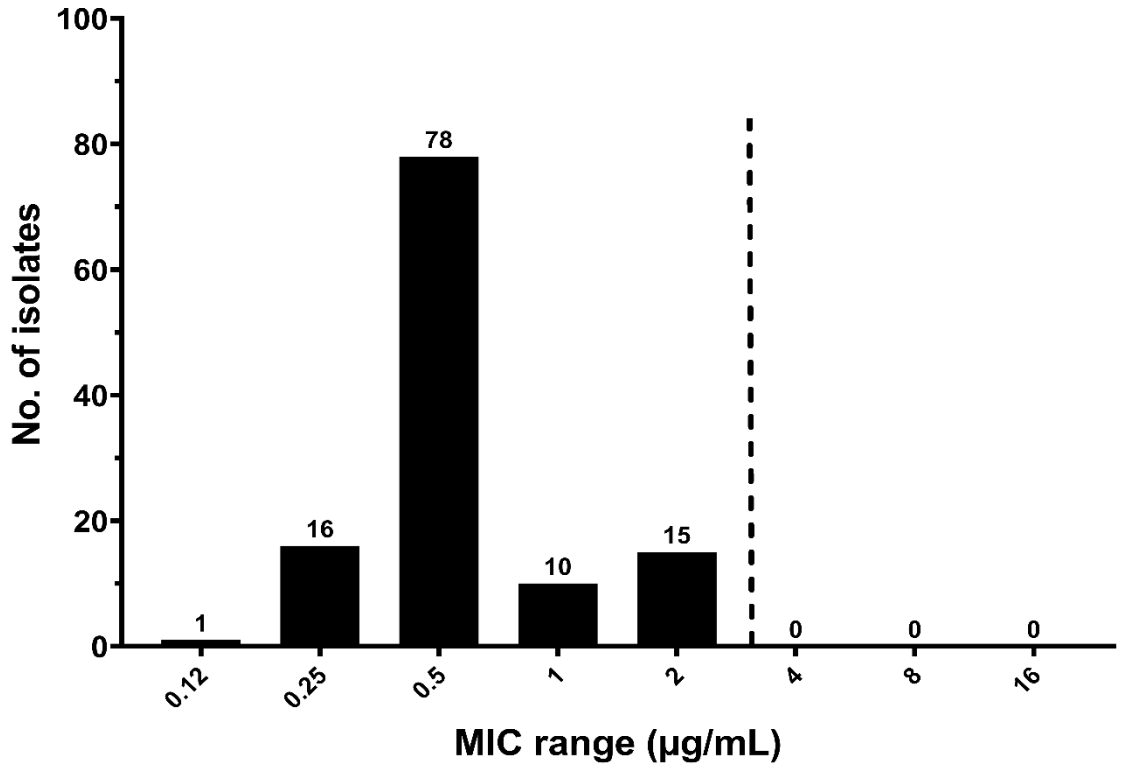

Figure S1. The distribution of minimum inhibitory concentration (MIC) values among the examined *Salmonella* isolates (n=120) against Colistin (CST). The dashed line indicates the cutoff level of the MIC, in which the lowest values correspond to susceptibility and the highest values correspond to the resistance.

Supplementary Tables

Table S1: Primers used for the detection of antimicrobial resistance genes.

Target	Forward primer (5′–3′)	Reverse primer (5′–3′)	Annealing Temperature (°C)	Product size (bp)	References
β-Lactamases and ESBLs:					
bla$_{TEM}$	GTATTCAACATTTCCGTGTCG	CCAATGCTTAATCAGTGAGGC	55	854	[1]
bla$_{SHV}$	CCCTGTTAGCCACCCTGCCG	CGTTGCCAGTGCTCGATCAGC	68	829	[1]
bla$_{CTX-M-1}$	GCTGTTGTTAGGAAGTGTGCCGC	GCCGCCGACGCTAATACATC	63	798	[1]
bla$_{CTX-M-2}$	AAATGTGCTGCTCCTTTCGTGAGC	AGGGTTCGTTGCAAGACAAGACTG	55	1122	[2]
bla$_{CTX-M-3}$	GTTACAATGTGTGAGAAGCAG	CCGTTTCCGCTATTACAAAC	55	1018	[3]
bla$_{CTX-M-8}$	TAACGCACAGACGCTCTACC	TGGCTGGGTGAAGTAAGTGAC	55	637	[4]
bla$_{CTX-M-9}$	GATCAGTCAGTGGGATAGTTT	TACTCGGCGTTAACTGATTA	47	671	[5]
bla$_{CTX-M-25}$	GCGATGTTAATGACGACAGC	AACCGTCGGTGACAATTCTG	55	847	[4]
bla$_{OXA-1}$	ACACAATACATATCAACTTCGC	AGTGTGTTTAGAATGGTGATC	55	814	[3]
bla$_{OXA-2}$	TTCAAGCCAAAGGCACGATAG	TCCGAGTTGACTGCCGGGTTG	60	703	[3]
bla$_{OXA-10}$	CGTGCTTTGTAAAAGTAGCAG	CATGATTTTGGTGGGAATGG	55	389	[3]
bla$_{PSE}$	TGCTTCGCAACTATGACTAC	AGCCTGTGTTTGAGCTAGAT	55	438	[3]
Cephalosporins:					
bla$_{MOX}$	GCTGCTCAAGGAGCACAGGAT	CACATTGACATAGGTGTGGTGC	55	520	[6]
bla$_{CIT}$	TGGCCAGAACTGACAGGCAAA	TTTCTCCTGAACGTGGCTGGC	55	462	[6]
bla$_{DHA}$	AACTTTCACAGGTGTGCTGGGT	CCGTACGCATACTGGCTTTGC	55	405	[6]
bla$_{ACC}$	AACAGCCTCAGCAGCCGGTTA	TTCGCCGCAATCATCCCTAGC	55	346	[6]
bla$_{EBC}$	TCGGTAAAGCCGATGTTGCGG	CTTCCACTGCGGCTGCCAGTT	55	302	[6]
bla$_{FOX}$	AACATGGGGTATCAGGGAGATG	CAAAGCGCGTAACCGGATTGG	55	190	[6]
Aminoglycosides:					
armA	CAAATGGATAAGAATGATGTT	TTATTTCTGAAATCCACT	55	777	[7]
rmtA	ATGAGCTTTGACGATGCCCTA	TCACTTATTCCTTTTTATCATG	55	756	[7]
rmtB	ATGAACATCAACGATGCCCT	CCTTCTGATTGGCTTATCCA	55	769	[7]
rmtC	CGAAGAAGTAACAGCCAAAG	ATCCCAACATCTCTCCCACT	55	711	[7]
rmtD	CGGCACGCGATTGGGAAGC	CGGAAACGATGCGACGAT	55	401	[7]
rmtE	ATGAATATTGATGAAATGGTTGC	TGATTGATTTCCTCCGTTTTTG	55	819	[7]
npmA	CTCAAAGGAACAAAGACGG	GAAACATGGCCAGAAACTC	55	641	[8]
aac(3)-IV	GTTACACCGGACCTTGGA	AACGGCATTGAGCGTCAG	55	674	[9]

aac6'-Ib	TTGCGATGCTCTATGAGTGGCTA	CTCGAATGCCTGGCGTGTTT	55	482	[10]
aadA1-like	GTGGATGGCGGCCTGAAGCC	ATTGCCCAGTCGGCAGCG	55	527/1916	[9]
Sulphonamides:					
sul1	TGGCGTCGCGACTGCGAAAT	TGGTGACGGTGTTCGGCATTCT	55	813	[11]
sul2	GTTTCTCCGATGGAGGCCGGT	AGCGAGGTTTCGGGAGCAGC	55	517	[11]
sul3	GATAGTTTTTCCGATGGAGG	GAAGCCCATACCCGGATCAA	55	495	[11]
Tetracyclines:					
tetA	TGGTCCGGAGGCCAGACGTG	TTCCGAGCATGAGTGCCCGC	55	867	[11]
tetG	TCTTGCAGGAGCCGCAGTCGAT	GGCCGGCATGCCAACACCC	55	721	[11]
tetX	CAATAATTGGTGGTGGACCC	TTCTTACCTTGGACATCCCG	55	468	[11]
tetB	AAATAACAGCAAACAGTAATGG	AAGTAGGGGTTGAGACGCAGCTA	55	493	[12]
tetC	TCTAACAATGCGCTCATCGT	GGTTGAAGGCTCTCAAGGGC	55	589	[12]
tetR	AGGACGACGGTGCTTGCT	ATGAGGACTGGCGGGTGT	55	295	[12]

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