

Table S1. Oligonucleotide primer sequences used for PCR assays.

Target Gene	Primer	Oligonucleotide Sequence (5'→3')	Product Size (bp) [§]	Annealing Temperature (°C)	References
<i>K. pneumoniae</i> 16S-23S ITS*	<i>K. pneumoniae</i> Pf	ATTGAAAGAGGTTGCAAACGAT	130	57	[23]
<i>bla</i> _{OXA-1}	<i>K. pneumoniae</i> Pr1	TTCACTCTGAAGTTTCTGTGTTG			
	OXA-F	GGCACCAAGATTCAACTTCAAG	564		
<i>bla</i> _{CTX-M1}	OXA-R	GACCCCAAGTTCCCTGTAAGTG			
	CTX-M-1-F	TTAGGAAGTGTCCCCGTGTA	655		
<i>bla</i> _{SHV}	CTX-M-1-R	CGGTTTATCCCCAACAC		61	[33]
	SHV-F	AGCGCTTGAGCAAATTAAAC	713		
<i>bla</i> _{TEM}	SHV-R	ATCCCGCAGATAAATCACCAC			
	TEM-F	CATTTCCGTGTCGCCCTTATTG	800		
	TEM-R	CGTTCATCCATAGTTGCCTGAC			
<i>bla</i> _{IMP}	IMP1-F	CATGGTTGGTGGTTCTTGT	488	53	[34]
	IMP1-R	ATAATTGGCGGACTTTGGC			
<i>bla</i> _{VIM}	VIM-F	AGTGGTGAGTATCCGACA	280	52	[35]
	VIM-R	ATGAAAAGTGCCTGGAGAC			
<i>bla</i> _{NDM1}	NDM-1-F	GGCGGAATGGCTCATCACGA	287	55	[36]
	NDM-1-R	CGCAACACAGCCTGACTTTC			
<i>mcr-1</i>	CLR5-F	CGGTCACTCCGTTGTT	309	55	[37]
	CLR5-R	CTTGGTCGGTCTGTAGGG			
ERIC consensus	ERIC1R	ATGTAAGCTCCTGGGGATTAC		Variable	52
	ERIC2	AAGTAAGTGAATGGGTGAGCG			[38]

* ITS, Internal transcribed spacer; [§]bp, base pair.

Table S2. Frequency distribution of *K. pneumoniae* isolates recovered from broiler farms and human workers in the study area.

Farms	C	Chicken				TC	Cw**		
		Ce*							
		Water	Food	Litter	Total				
1	2	1	0	1	2	4	1		
2	0	0	0	0	0	0	0		
3	3	2	1	1	4	7	2/3		
4	1	1	0	0	1	2	1		
5	1	0	0	1	1	2	0		
6	1	0	0	1	1	2	1		
7	0	0	0	0	0	0	0		
8	0	0	0	0	0	0	0/3		
9	0	0	0	0	0	0	0		
10	1	0	0	1	1	2	0		
Total (%)	9/100 (9)	4/20 (20)	1/20 (5)	5/20 (25)	10/60 (16.7)	19/160 (11.9)	5/22 (22.7)		

C: diseased chicken; Ce: chicken environment samples; TC: total chicken samples.

* Two pooled (a pool of three) samples for each type per farm; ** Cw: stool samples of chicken workers in broilers farms

Table S3. Source associated variations in phenotypic and genetic antibiotic resistance traits of *K. pneumoniae* isolates from chickens and humans in this study.

Variables	Source	Positive	OR	*p-Value	95% CI
Carbapenemase encoding gene	<i>bla</i> _{VIM}	Chicken 1/19 (5.3) Human 12/18 (66.7)	- 36	- 0.002	- 3.8–337.9
	<i>bla</i> _{NDM}	Chicken 2/19 (10.5) Human 10/18 (55.6)	- 10.6	- 0.01	- 1.9–60.2
	Imipenem	Chicken 3/19 (15.8) Human 14/18 (77.8)	- 18.7	- 0.001	- 3.6–98.2
	Gentamicin	Chicken 5/19 (26.3) Human 11/18 (61.1)	- 4.4	- 0.04	- 1.1–17.7
Phenotypic resistance	Aztreonam	Chicken 4/19 (21.1) Human 10/18 (55.6)	- 4.7	- 0.04	- 1.1–19.8
	Azithromycin	Chicken 7/19 (36.8) Human 17/18 (94.4)	- 29.1	- 0.003	- 3.2–268.8

* There was no significant association with regard to other antibiotic resistance genes or tested antimicrobials among the detected isolates. OR: Odds ratio; CI: Confidence interval. *p*-value is considered significant at ≤ 0.05 .

References

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