ORIGINAL ARTICLE Co-occurrence of Plasmid-mediated Quinolone Resistance and Carbapenemases in *Klebsiella pneumoniae* Isolates in Assiut, Egypt

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ABSTRACT

Key words: K. pneumoniae, PMQR, Carbapenemases

*Corresponding Author: Sherine A Aly Dept. of Medical Microbiology and Immunology, College of Medicine, Assiut University, Assiut 71516, Egypt. s-aly71@windowslive.com Tel: +201097018789 Background: Co-occurrence of carbapenem and fluoroquinolone resistance amongst K. pneumoniae strains created a problem in treating infections caused by these MDR organisms. Objectives: This study was carried out to evaluate the co-existence of carbapenemases and plasmid mediated quinolone resistance (PMQR) determinants in K. pneumoniae isolates in Egypt. Methodology: Forty-three K. pneumoniae isolates were collected from patients admitted to various intensive care units at Assiut University Hospital. Genes encoding for carbapenemases and PMOR were detected by PCR and sequencing. To determine the horizontal transfer of the PMOR and /or carbapenemases positive plasmids, conjugation experiments were performed. **Results:** Carbapenemases were detected in 34/43 (79.1%) of K. pneumoniae isolates. The positive rates of bla_{KPC} and bla_{NDMI} were (48.8%) and (74.4%), respectively. PMQR determinants were detected in 100% of K. pneumoniae isolates. The positive rates of qnrB, qnrS and aac(6`)-Ib-cr were (83.7 %), (81.4 %) and (23.3%), respectively. bla NDM1 positive K. pneumoniae positive isolates co-harbored qnr S, qnrB and aac(6`)-Ib-cr at rates of (87.5%), (81.3%) and (25%) respectively, while (90.5%), (85.7%) and (9.5%) of bla_{KPC} positive isolates co-harbored qnrS, qnrB and aac(6`)-Ib-cr, respectively. qnrB and qnrS exhibited statistically significant association with bla_{KPC} and bla_{NDMI} (p<0.001). Conclusion: Our study revealed high rate of co-existence of carbapenemases and PMQR determinants in K. pneumoniae isolates in Egypt. To our best of knowledge, this is the first study to report the presence of a statistically significant relation between carbapenemases and PMQR.

INTRODUCTION

Klebsiella pneumoniae (K. pneumoniae) is an important pathogen causing a broad range of community and hospital acquired infections ¹. These infections are usually caused by multidrug resistant (MDR) strains ^{2,3}. Carbapenems or fluoroquinolones (FQs) are frequently used for the management of infections initiated by such organisms. Unfortunately, the emergence of carbapenem or FQ resistant K. pneumoniae strains has been increasing ⁴.

The most important mechanism that confers resistance to carbapenems is the spread of plasmidmediated carbapenemases of class A (KPCs) and class B (VIMs, IMPs, and NDM-1) β -lactamases⁵. Quinolone resistance may be chromosomally mediated or plasmid mediated⁶. Plasmid-mediated quinolone resistance (PMQR) determinants include: *qnr* determinants (*qnrA*, *qnrB*, *qnrC*, *qnrD* and *qnrS*) that act by protection of target enzymes from inhibition by fluoroquinolones⁷, efflux pumps (*qepA*, *oqxAB*)⁸ and aminoglycoside acetyltranferase (6[°])-Ib-cr variant enzyme that confers reduced susceptibility to ciprofloxacin by N-acetylation of its piperazinylamine⁹. An emerging co-existence of carbapenems and fluoroquinolone resistance in K. *pneumoniae* is causing major difficulty in treating infections caused by such pathogen ¹⁰.

In Egypt several studies had evaluated the presence of either carbapenemases or PMQR determinants in *K. pneumoniae* isolates ¹¹⁻¹³. As far as we know this is the first report that investigated the co-existence of carbapenemases and PMQR determinants in *K. pneumoniae* isolates in Egypt.

METHODOLOGY

Isolation and identification of K. pneumoniae:

Two hundred clinical samples were collected from patients hospitalized at Intensive Care Units of Assiut University Hospitals between May 2015 and June 2016. Seventy (35%) isolates were identified as *Klebsiella species* by conventional bacteriological methods. The API 20E system confirmed 35/70 (50%) as *K. pneumoniae* (BioMerieux, Marcy L'Etolie, France). Additionally, 8 *K. pneumoniae* isolates recovered from Pediatric ICU during a previous study were included in this study ³. A total of 43 *K. pneumoniae* isolates were tested in the study.

Antimicrobial susceptibility testing:

Antimicrobial susceptibility profile was determined for all K. pneumoniae isolates by Kirby-Bauer disk diffusion method using commercial antibiotic discs (Oxoid, UK): amoxacillin (10µg), amoxacillinclavulanic acid (20,10µg), piperacillin (100µg), cefazolin (30µg), cefpodoxime (30µg), cefoperazone (75µg), ceftriaxone (30µg), aztreonam (30µg), gentamycin (10µg), amikacin (30µg) and tetracycline (30µg). The minimum inhibitory concentrations (MICs) of imipenem and ciprofloxacin were determined using E-tests (BioMérieux, Solna, Sweden). The results were interpreted according to the Clinical and Laboratory Standard Institute guidelines ¹⁴.

Phenotypic detection of carbapenemases:

Carbapenemase production by *K. pneumoniae* isolates was screened by Modified Hodge test (MHT), as previously described ¹⁵. MBLs activity was determined by combined disk (CD) test, using EDTA as inhibitor ¹⁶ and by E-test MBLs (Liofilchem, Italy). **Detection of genes encoding carbapenemases and**

PMQR determinants by PCR:

DNA was extracted from *K. pneumoniae* isolates by boiling method ¹⁷. Screening of genes encoding carbapenemases (bla_{KPC} , bla_{VIMI} , bla_{IMP} and bla_{NDMI}) and PMQR determinants (*qnrA*, *qnrB*, *qnrS*, *qepA* and *aac*(6`)-*lb-cr*) was performed by PCR. Amplification was done in a thermal cycler (BioRadT100, USA) using hot start taq DNA polymerase (Biolabs, New England) in a 50-µl volume. Annealing temperatures and sequences of oligonucleotide primers used for PCR amplification are listed in table (1). The sequences obtained for each gene were compared with published sequences on the National Center for the Biotechnology website (http://www.ncbi.nlm.nih.gov).

After PCR amplification, the aac(6')-Ib positive products were further subjected to digestion with the restriction enzyme BtsCI (Jena Bioscience, Germany) for detection of the *cr* variant. The detection of 272-bp and 210-bp DNA fragments was suggestive of aac(6')-*Ib*, while an undigested fragment was suggestive of aac(6')-*Ib*-*cr* variant.

Table 1: Oligonucleotide	primers and annealing tem	peratures used for PCR am	plification of different genes.

Gene	Sequence (5'-3')	Size	Tm	Reference
bla _{NDM1}	F: GGTTTGGCGATCTGGTTTTC	621 bp	55 °C	[31]
	R: CGGAATGGCTCATCACGATC			
bla _{VIM1}	F: TCTACATGACCGCGTCTGTC	747 bp	58 ℃	[32]
	R: TGTGCTTTGACAACGTTCGC			
bla _{IMP}	F: CTTGATGAAGGCGTTTATGTT	610 bp	54 °C	[33]
	R: TAACCGCCTGCTCTAATGTAAG			
bla_{KPC}	F: ATGTCACTGTATCGCCGTCT	893 bp	57 °C	[34]
	R: TTTTCAGAGCCTTACTGCCC			
qnrA	F: ATTTCTCACGCCAGGATTTG	518 bp	54 °C	[35]
	R: GATCGGCAAAGGTTAGGTCA			
qnrB	F: GATCGTGAAAGCCAGAAAGG	469 bp	54 °C	[36]
	R: ACGATGCCTGGTAGTTGTCC			
qnrS	F: CAATCATACATATCGGCACC	641 bp	56 ℃	[37]
	R: TCAGGATAAACAACAATACCC			
aac(6)-Ib	F: TTGCGATGCTCTATGAGTGGCTA	482 bp	58 ℃	[38]
	R: CTCGAATGCCTGGCGTGTTT			
qepA	F: GCAGGTCCAGCAGCGGGTAG	218 bp	58 ℃	[39]
	R: CTTCCTGCCCGAGTATCGTG			

Conjugation experiment:

To determine the transmissibility of the resistance determinants, conjugation experiment was performed on all *K. pneumoniae* isolates. Horizontal transfer of antimicrobial resistance genes was investigated with the conjugation method modified from Miller (1972) ¹⁸. All isolates were grown on Luria–Bertani (LB) broth with *E. coli* J53 as the recipient. Selection of transconjugants was done on LB agar plates supplemented with sodium azide (200 µg/ml) for counter selection, and imipenem (0.5 µg/ml) or ciprofloxacin (2 µg/ml). The presence of PMQR and carbapenemases in transconjugants was determined by PCR.

RESULTS

Antimicrobial susceptibility pattern:

Forty *K. pneumoniae* isolates were found to be MDR while 3 isolates expressed Pan Drug resistant (PDR) phenotype (resistant to all drug classes). Twenty-three isolates were resistant to imipenem when tested by E-test®. Carbapenemase production was detected in 31/43 (72.1%) of *K. pneumoniae* isolates by MHT, while MBLs production was determined in 25/43 (58.1%) of *K. pneumoniae* isolates by both CD test and E-test MBLs.

Prevalence of carbapenemases among *K. pneumoniae* isolates:

Carbapenemase encoding genes were detected by PCR in 34/43 (79.1%) of *K. pneumoniae* isolates. bla_{KPC} was detected in 21 (48.8%), while bla_{NDM1} was detected in 32 (74.4%) of the isolates (Table 2). bla_{KPC} and bla_{NDM1} co-existed in 19/34 (55.9%) of carbapenemase

positive isolates. None of the isolates harbored bla_{VIMI} or bla_{IMP} . Both bla_{KPC} and bla_{NDMI} were statistically significant positively correlated to imipenem resistance pattern (r=0.44, p=0.004), (r=0.39, p=0.010) respectively (Table 3). There was statistically significant positive weak correlation between bla_{NDMI} and bla_{KPC} (r=0.36, p=0.018).

Table 2: Carbapenemases, PMQR determinants and associated MICs for imipenem and ciprofloxacin in *Klebsiella pneumoniae* isolates (n=43)

ID	MIC IMP	Carbapene	mases	MIC CIP	PMQR genes	Phenotype
Carbapene	em resistant isola					
K1	>32 (R)	KPC, NDM1	>32(R)	q	nrB, qnrS	MDR ^{1,2,3,4,7,8}
K2	>32 (R)	KPC, NDM1	>32(R)	q	nrB, qnrS	PDR
К3	>32 (R)	KPC, NDM1	>32(R)	q	nrB, qnrS	MDR ^{1,2,3,4,5,7,8,9}
K4P	>32 (R)	KPC, NDM1	>32 (R)	q	nrB, qnrS	MDR ^{1,2,3,4,7,8,9}
K5	>32 (R)	KPC, NDM1	>32 (R)	q	nrB, qnrS	MDR ^{1,2,3,4,5,7,8,9}
K6	>32 (R)	KPC, NDM1	>32(R)	an B, an	erS, aac(6`)-Ib-cr	PDR
K7	16 (R)	NDM1	>32(R)		aac(6`)-Ib-cr	MDR ^{1,2,3,4,6,7,8}
K8	16 (R)	KPC, NDM1	>32 (R)	1 ,	qnrS	MDR ^{1,2,3,4,7,8,9}
K9	12 (R)	NDM1	>32(R)	anrB. an	urS, aac(6`)-Ib-cr	MDR ^{1,2,3,4,6,7,8}
K10P	12 (R)	KPC, NDM1	>32 (R)		nrB, qnrS	MDR ^{1,2,3,4,7,8,9}
K101 K11	12 (R) 12 (R)	NDM1	1(S)	9	qnrB	MDR ^{1,2,3,4,5,6,8,9}
K12	8 (R)	KPC, NDM1	0.125 (S)	a	nrB, qnrS	MDR ^{1,2,3,4,6,8,9}
K12 K13	8 (R)	KPC, NDM1	>32 (R)	-	-	MDR MDR ^{1,2,3,4,7,8,9}
K15 K14	6 (R)	KPC, NDM1 KPC, NDM1	>32 (R) >32(R)		nrB, qnrS aac(6`)-Ib-cr	MDR MDR ^{1,2,3,4,7}
		,		qnrs,		MDR ^{1,2,3,4,7,8,9}
K15	6 (R)	NDM1	>32 (R)	ת	qnrS	MDR ^{1,2,3,4,5,7,9} MDR ^{1,2,3,4,5,7,9}
K16	6 (R)	NDM1	>32 (R)		nrS, aac(6`)-Ib-cr	MDR ^{1,2,3,4,6,7,8,9} MDR ^{1,2,3,4,6,7,8,9}
K17	4 (R)	NDM1	>32(R)	-	aac(6)-Ib-cr	
K18	4 (R)	NDM1	>32(R)	qnrS,	aac(6`)-Ib-cr	PDR
K19	4 (R)	NDM1	>32 (R)		qnrS	MDR ^{1,2,3,4,7,8,9}
K20	4 (R)	-	>32(R)	qnrB, qn	ırS, aac(6`)-Ib-cr	MDR ^{1,2,3,4,5,7,8,9}
K21	4 (R)	-	6(R)		qnr B	MDR ^{1,2,3,4,7,8,9}
K22	4 (R)	NDM1	1(S)	q_1	nr B, qnrS	MDR ^{1,2,3,4,5,6,8}
K23	4 (R)	KPC, NDM1	0.75(S)	q_{I}	nr B, qnrS	MDR ^{1,2,3,4,5,6,8}
K24	4 (R)	KPC, NDM1	0.75(S)		qnrS	MDR ^{1,2,3,4,8,9}
K25P	4 (R)	KPC, NDM1	0.38 (S)	q	nrB, qnrS	MDR ^{1,2,3,4,8,9}
K26P	4 (R)	KPC, NDM1	0.06 (S)		nrB, qnrS	MDR ^{1,2,3,4,8,9}
K27P	4 (R)	KPC, NDM1	0.06 (S)		nr B, qnrS	MDR ^{1,2,3,4,8,9}
K28P	4 (R)	KPC, NDM1	0.04(S)	1	qnrB	MDR ^{1,2,3,4,8,9}
	em intermediate		0.0 ((5)		9.02	
K29	3 (I)	-	>32(R)	anrS	, aac(6`)-Ib-cr	MDR ^{1,2,3,5,7,8}
K2) K30	3 (I) 3 (I)	_ NDM1	>32(R)	1	rS, aac(6`)-Ib-cr	MDR ^{1,2,3,5,7,8}
K30 K31P	3 (I) 3 (I)	KPC, NDM1	0.75(S)	1 1	nr B, qnrS	MDR ^{1,2,3,8,9}
K311 K32	3 (I) 2 (I)	-	4(R)	q_{I}	an B, quis anr B	MDR ^{1,2,3,5,6,7,8,9}
K32 K33	2 (I) 1.5 (I)	-		~	qnr Б nr B, qnrS	MDR MDR ^{1,2,3,5,6,7,8,9}
кзз К34Р		- KPC, NDM1	6(R)	-	-	MDR ^{1,2,3,6,8,9}
	1.5 (I)		0.12 (S)	q	nr B, qnrS	MDK MDK
	em sensitive isola		(1)		D C	MDR ^{1,2,3,5,6,7,8,9}
K35	1 (S)	NDM1	>32(R)	q_1	nr B, qnrS	MDR ^{1,2,3,5,6,7,8,9}
K36	0.75 (S)	KPC	>32(R)		qnr B	MDR ^{1,2,5,5,6,7,6,9}
K37	0.75 (S)	KPC	>32(R)	1	nr B, qnrS	MDR ^{1,2,3,5,6,7,8,9}
K38	0.75 (S)	NDM1	>32(R)	q_{1}	nr B, qnrS	MDR MDR ^{1,2,3,5,6,7,8,9}
K39	0.5 (S)	-	8(R)		qnr B	MDR ^{1,2,3,5,6,7,8,9}
K40	0.25 (S)	-	4 (R)	q_1	nr B, qnrS	MDR ^{1,2,3,5,6,7,9}
K41	0.25 (S)	-	6 (R)	q	nr B, qnrS	MDP ^{1,2,3,5,6,7,8,9}
K42	0.125 (S)	NDM1	4 (R)	-	nr B, qnrS	MDR ^{1,2,3,5,6,7,8,9}
K43	0.125 (S)	-	2 (I)	-	nr B, qnrS	MDR ^{1,2,3,5,6,9}

Abbreviations: IMP, imipenem; CIP, ciprofloxacin; MDR, multi drug resistance; PDR, pan drug resistance.

Key: 1=penicillins; 2=cephalosporins; 3=monobactams; 4= carbapenems; 5=tetracyclines; 6=chloramphenicol; 7=fluoroquinolone; 8= trimethoprim sulfonadmide; 9 = aminoglycosides.

isoluces.						
Carbonomono	Total (43)	R (28)	I (6)	S (9)	-	
Carbapenemases	No. (%)	No. (%)	No. (%)	No. (%)	– r	p value
bla_{KPC}	21 (48.8%)	17 (60%)	2 (33.3%)	2 (22.2%)	0.44	0.004**
blaNDM1	32 (74.4%)	26 (92.9%)	3 (50%)	3 (33.3%)	0.39	0.010
R=Resistant	I=Intermedi	iate	S=Sensit	tive		

Table 3: Distribution of carbapenemases and its relation to carbapenem resistance pattern in K. pneumoniae isolates.

R=Resistant

** Statistically significant correlation (p<0.01)

Prevalence of PMQR among K. pneumoniae isolates: One or more of PMQR determinants were detected in each of K. pneumoniae isolates. Thirty six (83.7%) isolates harbored qnrB, 35 (81.4%) isolates harbored qnrS, while 10 (23.3%) isolates harbored aac(6`)-Ib-cr (Table 2). No qnrA or qepA was detected in any isolate. qnrB and qnrS were present in (90.9%), (81.8%) of

ciprofloxacin susceptible isolates, respectively. Statistically significant weak positive correlation between aac(6)-*Ib-cr* and ciprofloxacin resistance (r=0.49, p=0.001) was detected, while the presence of qnrB and qnrS had no relation to ciprofloxacin resistance pattern (Table 4).

Table 4: Distribution of PMQR determinants and their relations to ciprofloxacin resistance pattern in the K. *pneumoniae* isolates

PMQR	Total (43)	R (31)	I (1)	S (11)		n voluo
	No. (%)	No. (%)	No. (%)	No. (%)	- r	p value
qnrB	36 (83.7%)	25 (80.6%)	1 (100%)	10 (90.9%)	-0.26	0.096
qnrS	35 (81.4%)	25 (80.6%)	1 (100%)	9 (81.8%)	0.15	0.350
R=Resistant	I=Intermedi	ate	S=Sensit	tive		

R=Resistant

S=Sensitive

** Statistically significant correlation (p<0.01)

Co-existence of carbapenemases and PMOR determinants in K. pneumoniae isolates:

The prevalence of PMQR determinants amongst carbapenemase positive K. pneumoniae isolates (34) was 100%. Specifically, (87.5%), (81.3%) and (25%) of bla_{NDM1} positive isolates co-harbored qnrS, qnrB and

aac(6`)-Ib-cr, respectively, while (90.5%), (85.7%) and (9.5%) of *bla_{KPC}* positive isolates co-harbored *qnrS*, qnrB and aac(6)-*Ib-cr*, respectively. The presence of both bla_{NDMI} and bla_{KPC} was statistically significant higher in isolates containing qnrS or qnrB (Table 5).

Table 5: Co-existence of carba	penemases and PMO	R determinants among	K. <i>pneumoniae</i> isolates

PMQR No (%)					
qnrS	p value	qnrB	p value	aac(6`)-Ib-cr	
19 (90.5%)	< 0.001**	18 (85.7%)	< 0.001**	2 (9.5%)	
28 (87.5%)	< 0.001**	26 (81.3%)	< 0.001**	8 (25%)	
	19 (90.5%)	19 (90.5%) <0.001**	qnrS p value qnrB 19 (90.5%) <0.001**	qnrS p value qnrB p value 19 (90.5%) <0.001**	

** Statistically significant difference (p<0.01)

Conjugative transfer of resistance genes:

PMQR determinants (qnrB, qnrS and aac(6`)-Ib-cr) and carbapenemases $(bla_{KPC} \text{ and } bla_{NDM1})$ were successfully transferred by conjugation from all K. pneumoniae isolates to the recipient (E. coli J53).

DISCUSSION

Serious infections caused by MDR K. pneumoniae are frequently treated by carbapenems or fluoroquinolones¹⁹. The co-existence of PMQR determinants together with the carbapenemases among

K. pneumoniae isolates limits the treatment options for these drug-resistant strains 20 . In the present study, carbapenemases and PMQR coexisted in 34/43 of K. pneumoniae recovered from Assiut University Hospitals. As far as we know, the present study is the first to report the presence of statistically significant correlation between the occurrence of PMQR and carbapenemases as well as the co-existence of qnrB, qnrS, aac(6)-Ib-cr, bla_{KPC} and bla_{NDM1} in the same K. pneumoniae isolate.

Carbapenem resistance resulting from production of carbapenemases is disseminating worldwide²¹. High

^{*} Statistically significant correlation (p<0.05)

prevalence of carbapenemases (79.1%) among K. pneumoniae isolates was detected in our study. bla_{NDM1} was the most prevalent carbapenemase (74.4%), followed by bla_{KPC} (48.8%), while neither bla_{VIM1} nor bla_{IMP} was detected in any of K. pneumoniae isolates. In agreement with our results, previous studies reported similar prevalence of bla_{NDM1} and bla_{KPC} among K. pneumoniae^{22,23}. However, in a recent Egyptian study, lower prevalence of bla_{NDMI} was reported in K. pneumoniae isolates ²⁴. The elevated frequency of carbapenemase positive isolates in the present study might be due to the excessive use of carbapenem in our hospitals.

Although bla_{KPC} and bla_{NDM1} were detected in (22.2%), (33.3%) of the imipenem sensitive K. pneumoniae isolates, both bla_{KPC} and bla_{NDM1} were statistically significant positively correlated to imipenem resistance pattern (r=0.44, p=0.004), (r=0.39, p=0.010), respectively (Table 2). This finding contradicts the previous assumption that bla_{NDM1} or bla_{KPC} alone mediate only reduced susceptibility to carbapenems and other mechanisms, as porin loss, are usually required for full resistance to appear²⁵.

In an attempt to understand the relation between the existence of PMQR and carbapenemases, we tested K. pneumoniae isolates for the presence of PMQR by PCR. Our study reported high prevalence of PMQR determinants (100%) in K. pneumoniae isolates which is similar to the results of a previous Egyptian study ¹². Interestingly, all K. pneumoniae isolates (even those isolated from PICU where FOs are contraindicated) have at least one PMQR, which implies that quinolone resistant plasmids are endemic and circulating in Assiut University Hospital. qnrB contributed to the majority (83.7%) of PMQR determinants detected in this study, followed by qnrS, which was detected in (81.4 %) of K. pneumoniae isolates, while aac(6)-Ib-cr showed the lowest prevalence (23.3%).

Coexistence of resistance genes in the same isolate seemed to be the primary cause of the appearance of MDR or even PDR strains²⁶. Co-occurrence of carbapenemases and PMQR in the same isolate was previously described in *Klebsiella pneumoniae*^{27,28}. In the present study, (79.1%) of K. pneumoniae isolates co-harbored carbapenemases and PMQR determinants. bla_{KPC} and bla_{NDM1} co-existed with qnrS in (90.5%), (87.5%) of K. pneumoniae, respectively and with qnrB in (85.7%), (81.3%), of K. pneumoniae respectively. Excitingly, our results indicated statistically significant relations between different carbapenemases and PMQR determinants amongst K. pneumoniae isolates (p<0.001), which increase the possibility that the multiple resistance determinants might be co-expressed on the same plasmid.

The mobility of plasmid-mediated resistance carries additional risks of spread of resistance determinants between different species²⁹. In this study, PMQR determinants and carbapenemases were successfully

transferred via conjugation in all K. pneumoniae isolates. Our results are in accordance with what reported by previous studies ^{27,30}. This high transmissibility of PMQR genes and carbapenemases points to the risk of potential spread of these resistance determinants among other pathogens in the hospital, which intensify the need for implementation of strict infection control measures to limit their dissemination between strains.

It is prudent to say that all K. pneumoniae isolates recovered from PICU have the same combination of PMQR and carbapenemases (bla_{KPC}, bla_{NDM1}, qnrB, qnrS) although they have different PFGE pattern ³. Additionally, all the resistance genes were transferred successfully by conjugation, which is suggestive of the presence of an endemic plasmid harboring the four resistance genes circulating in the PICU.

CONCLUSION

The results of the current study revealed high prevalence of carbapenemases and PMQR determinants among K. pneumoniae isolates in Egypt, as well as the co-existence of multiple resistance determinants in various combinations in the same isolate. The presence of *bla_{KPC}*, *bla_{NDM1}* is significantly related to the presence of both *qnrB* and *qnrS* which suggest a role of these PMQR in favoring selection of carbapenemases, however, this speculation needs further investigation.

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Conflicts of interest:

The authors declare no conflicts of interest.

Ethical Statement:

The Ethics Committee of the Faculty of Medicine, Assiut University approved the study according to the latest revision of the Declaration of Helsinki, and informed consent was obtained from participating patients.

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