Detection of plasmid-mediated colistin resistance in carbapenem-resistant \textit{Escherichia coli} and \textit{Klebsiella pneumoniae} isolates in Suez Canal University Hospitals

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\textbf{Introduction}

The polymyxins including colistin are considered the last choice of antibiotics for treatment of infections with carbapenem-resistant \textit{Enterobacteriaceae} (CRE). In 2012 colistin was reclassified by the World Health Organization as critically important for human medicine [1]. Colistin interacts with the bacterial outer membrane by removing divalent cations from the negatively-charged phosphate groups of the Lipid A of the lipopolysaccharide membrane which leads to cell lysis [2].

Previously, colistin resistance was considered only due to chromosomal mutations in the genes encoding the PmrA/PmrB and PhoP/PhoQ systems. However, lately, transferable colistin resistance mediated by the \textit{mcr}-genes has been described in \textit{Enterobacteriaceae}. The testing of colistin susceptibility is challenging.

The aim of the work is to detect the occurrence of \textit{mcr}-1 and \textit{mcr}-2 genes in phenotypically colistin-resistant carbapenem-resistant \textit{Escherichia coli} (\textit{E. coli}) and \textit{Klebsiella pneumoniae} (\textit{K. pneumoniae}) isolates using conventional polymerase chain reaction (PCR).

\textbf{Methods:} One hundred and sixteen carbapenem-resistant \textit{E. coli} and \textit{K. pneumoniae} isolates were collected from patients admitted to different wards of Suez Canal University Hospitals (SCUHs) in Ismailia. Urine, endotracheal aspirates, blood, pus and sputum specimens were collected from different patients. Minimal inhibitory concentration (MIC) by broth microdilution method was done to assess phenotypic colistin resistance. The colistin resistant \textit{E. coli} and \textit{K. pneumoniae} isolates were tested by conventional PCR to detect plasmid mediated \textit{mcr}-1 and \textit{mcr}-2 genes. Chi-square test was applied and \textit{p}-value < 0.05 was considered statistically significant.

\textbf{Results:} Forty three isolates out of 116 carbapenem resistant isolates (37.1\%) were colistin resistant as shown by MIC by broth microdilution method. The 43 colistin resistant \textit{E. coli} and \textit{K. pneumoniae} isolates were tested by conventional PCR to detect plasmid mediated \textit{mcr}-1 and \textit{mcr}-2 genes. Two \textit{mcr}-2 genes and one \textit{mcr}-1 gene were detected.

\textbf{Conclusion:} Results revealed that the prevalence of colistin resistance among carbapenem-resistant \textit{E. coli} and \textit{K. pneumoniae} in SCUHs is concerning; further limiting potential therapeutic options. Plasmid mediated colistin resistance genes \textit{mcr}-1 and \textit{mcr}-2 is emerging in SCUHs which refer to a problem in the hospital as by horizontal transfer of this plasmid, the resistance can spread to many isolates in the hospital.
signalling systems or the negative regulator MgrB. These mutations cause modifications to the Lipid A molecule [2]. Recently, transferable resistance of the polymyxins mediated by the mcr- genes has been detected. These mcr- genes were located on the plasmid which raises the concern that the resistance may be transferred to Gram-negative bacteria. The mcr-1 gene firstly discovered in Escherichia coli (E. coli) and Klebsiella pneumoniae (K. pneumoniae) in China [3], after that several studies across the world reported the presence of the mcr-1 gene in various bacteria from many other countries such as United States, European Union, Turkey, South Africa, Malaysia, Greece, Italy, Algeria, Tunisia and Kuwait [4]. The first study carried out in Egypt in 2016 reported the presence of mcr-1 gene in E. coli strain taken from an intensive care unit (ICU) patient [5].

In 2016, the mcr-2 plasmid-mediated colistin resistance was detected in E. coli first in Belgium from pigs and then from patients [6] Mcr-2 gene which is a phospho-ethanolamine transferase, modifies the lipopolysaccharide and shares 80.6% identity with mcr-1. After that, transferable colistin resistance gene mcr-3 has been identified in E. coli and K. pneumoniae from Asia and the United States and then some studies found mcr 4, 5, 6, 7, 8 and 9 [7].

The transfering of plasmid-borne mcr genes will diminishes the clinical choice of the last-resort colistin, which may lead to more treatment failures and increased morbidity and mortality. As a result, the European Medicines Agency, is currently updating the guidelines of using colistin. They considered upgrading of colistin as an antibiotic category 2 used for critical medicine and only if no other options actually found [8].

Colistin susceptibility testing is considered challenging due to recommendation updates by Clinical and Laboratory Standards Institute (CLSI) and European Committee on Antimicrobial Susceptibility Testing (EUCAST) which were declared in 2016 [9] that stated standard broth microdilution (BMD) method to test colistin. As a result, colistin susceptibility is probably neither tested routinely nor correctly against CRE isolates in most diagnostic laboratories in Egypt, and the frequency with which colistin resistance occurs is difficult to estimate from local data.

The purpose of the present study is to evaluate the frequency of colistin resistance among E.coli and Klebsiella carbapenem resistant isolates obtained from patients in Suez Canal University Hospitals (SCUHs), and to determine how often mcr-1 and mcr-2 genes are detected among the colistin-resistant subset.

**Patients and Methods**

A cross-sectional descriptive study was conducted on patients admitted to SCUHs in Ismailia. One hundred and sixteen carbapenem resistant isolates of E.coli and K. pneumoniae were collected from those patients. The study was conducted on patients admitted to the urology, internal medicine, surgery wards, intensive care unit (ICU), and neonatal intensive care unit.

**Inclusion criteria:**
- Both sexes.
- All age groups.
- Patients showing signs of sepsis or pyogenic infections.
- Patients having risk factors for acquisition of infection as:
  - Patients having high-risk procedures (such as surgery, urinary tract instrumentation, indwelling urinary or intravenous catheters and patients admitted to the burn unit).
  - Immunocompromised and debilitated patients and patients having serious underlying disease.

**Exclusion criteria:**
- Patients refusing to participate in the study.

**Specimens:**

Different microbial specimens were collected on clinical suspicion of infection according to standard definitions. Specimens included urine from catheterized and non-catheterized patients, respiratory specimens (from both intubated and non-intubated patients), blood and swab specimens.

**Methods:**

1) **Collection of specimens:**
Specimens were collected under aseptic conditions and transported in suitable transport media (when needed) to be processed in the Microbiology laboratory Department, Faculty of Medicine, Suez Canal University for the isolation and identification of E. coli and K. pneumoniae.
2) **Processing of specimens:**

Collected specimens were inoculated onto blood agar plates and MacConkey agar plates. The plates were incubated aerobically at 35 ± 2°C for 24 hours.

A. **Identification of E.coli and K. pneumoniae** [10]:

They were non –spore forming Gram negative rods. On blood agar: they appeared as moist smooth and greyish-white colonies. On MacConkey agar: they appeared as pink colored colonies due to lactose fermentation. *K.pneumoniae* strains produced mucoid colonies. They were catalase positive, oxidase negative, ferment glucose with acid production and reduce nitrate to nitrite.

B. **Suspected isolates were screened for genus and species by biochemical reactions based on metabolic characteristics** [11]

*IMVC tests (Indole test, Methyl red, Voges-Proskauer test (VP), and Citrate test), ornithine decarboxylation and motility tests were done for lactose fermenting colonies.

3) **Antimicrobial susceptibility testing:**

Disk diffusion susceptibility test was performed on each isolate according to CLSI guidelines (2018) [12]. Antibiotic disks were purchased from (Oxoid, UK). The following antimicrobial agents were included:

- Amoxacillin-clavulanate(20/10 µg), Cefotaxime (30µg), Ceftazidime (30µg), Cefepime(30µg), Imipenem (10µg), Meropenem (10µg).
- Ciprofloxacins (5µg), Levofloxacin (5µg).
- Gentamycin (10µg), Amikacin (30µg).
- Trimethoprim-sulfamethoxazole (1.25/3.75µg).

4) **Screening & confirmatory tests for suspected carbapenemase production in Enterobacteriaceae isolates** [12]:

*E.coli* and *K.pneumoniae* isolates that showed inhibition zone diameter ≤ 22 mm for either meropenem or imipenem were considered carbapenem nonsusceptible (intermediate or resistant) and were subjected to the following:

❖ **Modified carbapenem inactivation method (mCIM).**

One microlitre loopful of the test isolate was suspended in 2 ml of trypticase soy broth (TSB), and meropenem disk was added into the suspension. The culture was incubated for 4 h at 35 °C. *Escherichia coli* ATCC 25922 (a carbapenem –sensitive type strain) was adjusted using McFarland standard no. 0.5 in sterile saline and streaked in three directions on the Mueller-Hinton Agar (MHA) plates to ensure even and standardized cell lawn. The disk was transferred from TSB suspension onto MHA plate, and the plate was incubated overnight at 35 °C. After incubation, the diameter of the inhibition-zone was measured. Carbapenemase positive test strain: if it showed inhibition zone 6-15 mm or presence of colonies within a 16-18 mm zone. Carbapenemase negative test strain: if it showed inhibition zone ≥19 mm. Intermediate test strain: if it showed inhibition zone 16-18mm.

5) **Phenotypic detection of colistin resistance:**

**Minimum inhibitory concentration method:**

Colistin susceptibility was performed by broth microdilution method according to CLSI 2012 [13]. Antibiotic stock solution was prepared from pure colistin sulfate powder (Sigma-Aldrich, Germany). The amount needed and the diluents in which it was dissolved (distilled water) was calculated by using the following formula to determine the amount of antimicrobial powder.

\[
\text{weight(mg)} = \frac{\text{volume(ml) \times concentration(\mu g/ml)}}{\text{potency(\mu g/mg)}}
\]

Preparation of antibiotic dilution range was done according to CLSI 2018 [12]. The inoculum was prepared by making a direct broth suspension (brain heart broth) of isolated colonies selected from an 18- to 24-hour blood agar plate. The suspension was adjusted to achieve a turbidity equivalent to a 0.5 McFarland turbidity standard. This resulted in a suspension containing approximately 1 to 2 x10⁸ colony forming units (CFU)/mL for *Escherichia coli* ATCC®a 25922. The adjusted inoculum suspension in broth was diluted to obtain a final concentration after inoculation approximately 5 x 10³ CFU/mL. Within 15 minutes after the inoculum has been standardized as described above, 100 µl of the adjusted inoculum was added to each well containing 100 µl of antimicrobial agent in the dilution series (and a positive control tube containing only broth), and mixed. The inoculated microtiter plate was incubated at 35 ± 2°C for 16 to 20 hours in an ambient air incubator. The amount of growth in the wells containing the antimicrobial agent was compared with the amount of growth in
the growth-control wells (no antimicrobial agent). For a test to be considered valid, acceptable growth (≥ 2 mm button or definite turbidity) must occur in the growth-control well.

The lowest concentration at which the isolate is completely inhibited (as evidenced by the absence of visible bacterial growth) is recorded as the minimal inhibitory concentration or MIC.

6) Genotypic detection of plasmid-mediated mcr-1 and mcr-2 colistin resistance genes.

Plasmid DNA was extracted from the test isolates by using QIAprep Mini-Prep Kit (QIAGEN, Germany). Polymerase chain reaction was performed and the product was analyzed according to the laboratory protocol optimized by National Food Institute, Denmark, for detection of plasmid-mediated colistin resistance genes, mcr-1 and mcr-2 [14], using previously published primers. The reaction mixture was prepared in a total volume of 25μl including 3μl of template DNA, 12.5 ul of 2X ABT Red master mix (Sigma-aldrich, Egypt), and 2 μl of both forward and reverse primers. The volume completed with distilled water up to 25μl. Reaction mixtures without a DNA template served as negative controls. Amplification was carried out in a thermal cycler (Peltier Thermal cycler, MJ Research, U.S.A) with the following thermal cycling: 95°C 5 mins + 25X (95°C 30 sec + 58°C 60 sec + 72°C 60 sec) + 72°C 10min. Amplicons obtained from PCR reactions were analyzed by gel electrophoresis (Major Science, Taiwan) and finally visualized with ultraviolet light. Amplicon size (bp) of the tested genes was identified and compared to a100 bp molecular size standard DNA ladder (Sigma-aldrich, Egypt).

Results

Results of demographic and clinical data

The study was carried out during the period from 2018 till 2020. Specimens collected from patients admitted in different wards in SCUHs in Ismailia. Number of specimens collected was 536, included endotracheal aspirate, blood, urine, pus and sputum specimens. Number of K. pneumoniae isolates was 189 and E.coli was 91 isolates. Among them carbapenem-resistant K. pneumoniae were 90 isolates and carbapenem-resistant E.coli were 26 isolates.

From the 536 examined clinical specimens, E.coli and K. pneumoniae were isolated at a rate of 52.2%, while other organisms were isolated at a rate of 45.4%. No growth was detected in 2.4% of the cases (Figure 1).

Number of E.coli and K.pneumoniae strains isolated was 280. Among them 116 (26 E.coli & 90 K.pneumoniae) were carbapenem resistant isolates (41.4%). The 116 carbapenem resistant E.coli and K.pneumoniae isolates were collected from 64 male patients (55.2%) and 52 female patients (44.8%).

Table 1 shows that 46.6% of the studied carbapenem-resistant strains were isolated from endotracheal aspirates. Other carbapenem-resistant strains were isolated from blood (21.6%), urine (9.4%), sputum (8.6%) and pus (13.8%) specimens.

Results of antibiotic susceptibility testing

Antibiotic susceptibility test was done to the carbapenem resistant K. pneumoniae (90, 77.6%) and E.coli (26, 22.4 %) isolates by disk diffusion method (Kirby Bauer method). It showed 100% extended spectrum beta-lactamase as all isolates were resistant to both amoxicillin -clavulanic and ceftazidime antibiotics. While gentamycin, amikacin, levofloxacin and cefipime showed 51.3%, 63%, 79.5% and 87.7% resistance respectively. (Figure 2).

Modified carbapenem inactivation method was done to assess carbapenemases activity among the resistant isolates. Eighty two isolates were carbapenemases positive (70.7%) (Figure 3). Among them, 14 out of 26 E.coli isolates were carbapenemase positive (53.8%). While 68 isolates out of the 90 carbapenem-resistant K.pneumoniae isolates were carbapenemase positive (75.6%).

Phenotypic colistin resistance detection was done to the 116 CR isolates by minimal inhibitory concentration method using broth microdilution method (Figure 4). Forty three isolates (37.1%) were colistin resistant (Non-wild type). Thirty nine colistin resistant isolates were collected from the ICU, while two isolate were from the internal medicine department and two from the surgery department. Out of the 43 isolates, 29 isolates were obtained from endotracheal aspirate specimens, 6 isolates were obtained from blood, 4 from urine and 4 from pus.

The 43 colistin resistant E.coli and K.pneumoniae isolates were tested by conventional PCR and agarose gel electrophoresis for detection of plasmid mediated colistin resistance genes mcr-1 and mcr-2 (Figure 5). Two mcr-2 genes and one mcr-1 gene were detected among the 43 isolates (Table 2).
Regarding the three positive mcr isolates; all of them were *K. pneumoniae* isolates, two were collected from endotracheal tube and all of them were from ICU. They showed high level of antibiotic resistance and they were phenotypically colistin resistant (Table 3).

**Figure 1.** Frequency distribution of *E. coli* and *K. pneumoniae* in relation to other organisms (N = 536).

**Figure 2.** Frequency of antibiotic resistance pattern of the studied carbapenem resistant *E. coli* and *K. pneumoniae* isolates (N = 116).

**Figure 3.** Frequency distribution of carbapenemases activity among *E. coli* and *K. pneumoniae* isolates by mCIM test (N = 116).
Figure 4. MIC (BMD) of Colistin antibiotic among *E.coli* and *K.pneumoniae* isolates according to CLSI 2019.

Row A shows MIC result=8, Row B MIC result=2, Row C MIC result=2, Row D MIC=8, Row E MIC=32, Row F MIC=16.

Figure 5. Detection of *mcr-2* gene by agarose gel electrophoresis (567 bp).

Lane M shows 100 bp molecular size standard DNA ladder. Lane 1 shows a positive control strain, lane 2 shows a negative control strain Lanes 5 shows a positive isolate.

Table 1. Distribution of the isolated carbapenem-resistant *E.coli* and *K.pneumoniae* isolates according to the type of the specimen.

<table>
<thead>
<tr>
<th>No. of specimens (N=280)</th>
<th>No. CR. (^1) isolates (N=116)</th>
<th>Percent of CR. specimens</th>
<th>Percent to all CR. strains.</th>
</tr>
</thead>
<tbody>
<tr>
<td>ETT(^2) (80)</td>
<td>54</td>
<td>67.5% (54/80)</td>
<td>46.6% (54/116)</td>
</tr>
<tr>
<td>Blood (45)</td>
<td>25</td>
<td>55.5% (25/45)</td>
<td>21.6% (25/116)</td>
</tr>
<tr>
<td>Urine (89)</td>
<td>11</td>
<td>12.3% (11/89)</td>
<td>9.4% (11/116)</td>
</tr>
<tr>
<td>Sputum (24)</td>
<td>10</td>
<td>41.6% (10/24)</td>
<td>8.6% (10/116)</td>
</tr>
<tr>
<td>Pus (42)</td>
<td>16</td>
<td>38% (16/42)</td>
<td>13.8% (16/116)</td>
</tr>
</tbody>
</table>

\(^1\) Carbapenem resistant  \(^2\) Endotracheal tube
Table 2. Frequency of *mcr*-1 & *mcr*-2 genes among phenotypic colistin resistant isolates (N=43).

<table>
<thead>
<tr>
<th></th>
<th>Number</th>
<th>Percentage</th>
<th>P value</th>
</tr>
</thead>
<tbody>
<tr>
<td>MCR-1 gene</td>
<td>1/43</td>
<td>2.3%</td>
<td>&gt;0.05</td>
</tr>
<tr>
<td>MCR-2 gene</td>
<td>2/43</td>
<td>4.6%</td>
<td>&gt;0.05</td>
</tr>
</tbody>
</table>

Table 3. Data about the three colistin resistant isolates by PCR.

<table>
<thead>
<tr>
<th>Specimen</th>
<th>Department</th>
<th>Organism</th>
<th>Antibiotic susceptibility</th>
<th>Carbapenemase result</th>
<th>MIC result</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mcr-1 positive</td>
<td>Endo-tracheal tube</td>
<td><em>K.pneumoniae</em></td>
<td>Resistant to all antibiotics used except gentamycin and amikacin</td>
<td>Negative</td>
<td>16</td>
</tr>
<tr>
<td>Mcr-2 Positive</td>
<td>Pus</td>
<td><em>K.pneumoniae</em></td>
<td>Resistant to all except gentamycin, amikacin and ciprofloxacin</td>
<td>Negative</td>
<td>128</td>
</tr>
<tr>
<td>Mcr-2 Positive</td>
<td>Endo-tracheal tube</td>
<td><em>K.pneumoniae</em></td>
<td>Resistant to all antibiotics used</td>
<td>Positive</td>
<td>16</td>
</tr>
</tbody>
</table>

Discussion

With the increased spread of carbapenem resistance, treatment options are decreasing. Colistin is considered as a drug of choice in the treatment of carbapenem resistant isolates. Unfortunately, colistin resistance has emerged and became a big problem [3].

*Klebsiella pneumoniae* carbapenem resistant isolates were 90 out of 189 *K. pneumoniae* isolates (47.6%) while *E.coli* carbapenem resistant isolates were 26 out of 91 *E.coli* isolates (28.6%). This result shows accordance with El-Sweify *et al.* [15] who reported carbapenem resistance in 44.3% of *K. pneumoniae* isolates in SCUHs. Higher results were detected by Melake *et al.* [16] and Amer *et al.* [17] as they reported 62.5% and 62.7% carbapenem-resistance among *K.pneumoniae* and *Enterobacteriaceae* isolates. On the other hand, lower results of carbapenem resistance were also reported in different studies as Khare *et al.* [18] and Okoche *et al.* [19] reported 37.9% and 28.6% carbapenem resistance. Khalifa *et al.* [20] reported 15.7% *E.coli* carbapenem resistant isolates. The difference of prevalence among different studies may be due to the difference in application of infection control measures and antibiotic policies in different hospitals. Also the variety of specimens obtained from different wards all over the hospitals may contribute to the difference of these results. The high trend of carbapenem resistant *K.pneumoniae* in our study could be attributed to the frequent use of carbapenems as an empiric therapy in our hospital, as well as lack of implementation of antimicrobial stewardship program.

The current study showed that 46.6% of the collected carbapenem-resistant isolates were in endotracheal aspirates obtained from the ICU. That shows accordance with the study of Saeed *et al.* [21] as they collected most of the carbapenem resistant isolates from endotracheal aspirates in ICU. The high prevalence of carbapenem-resistant isolates found in the ICU with highest prevalence among endo-tracheal aspirates may be related to the low immunity status of patients and the respiratory tract invasive procedures done in ICU. It may also reflect cross-infection with multi-resistant clones or long-term exposure of respiratory tract microbiota to antibiotics, causing accumulation of resistance determinants in strains causing respiratory tract infection. Also, the presence of the tube for long period may cause inflammation and edema of the respiratory tract which aids in repeated infections.

Antibiotic susceptibility testing was done to the carbapenem resistant *K.pneumoniae* and *E.coli* isolates. It showed 100% resistance to β-lactam antibiotics due to ESBL production, while gentamycin, amikacin, levofloxacin and cefepime showed 51.3%, 63%, 79.5% and 87.7% resistance.
respectively. In a study done by Lin et al. [22] antibiotic susceptibility to carbapenem-resistant isolates showed resistance to amikacin 55.3%, ciprofloxacin 84.2%, amoxicillin/Clavulanic acid 97.4% and cefepime 97.4% which showed accordance with our results. Also a study done in Egypt in El-Mansoura University by Moemen and Masallat [23] reported high resistance among the carbapenem-resistant isolates as it showed ESBL production in 97.6%, amikacin 52.4%, ciprofloxacin 95.2%. Another study done in Egypt by Shawky et al. [24] reported 100% ESBL production among their isolates, 77.7% resistance to amikacin and 87.1% resistance to ciprofloxacin. The abuse of antibiotics is the basis of appearance of the high level resistance among different antibiotics, also self-medication by patients and lack of implementation of antibiotic policies in hospitals may contribute.

Modified carbapenm inactivation method was done in our study to assess the presence of carbapenemases enzymes among the resistant isolates. Among the 26 carbapenem-resistant E.coli isolates, 14 isolates were carbapenemase positive (53.8%). While 68 isolates out of the 90 carbapenem-resistant K.pneumoniae isolates were carbapenemase positive (75.6%). Similar levels of carbapenemases detection was found in a study by Laolerd et al. [25] and a study done by Qadri et al. [26] as they detected 77.7% and 67.3% carbapenemases among carbapenem resistant isolates respectively. Decreased rates of carbapenemase production among the carbapenem-resistant isolates were detected in another study conducted by Kandeel [27] who reported 18.5% carbapenemases among his isolates. In the current study many of potential risk factors for carbapenemases acquisition were observed. This is similar to previous reports describing these risk factors, including prolonged hospital stay, antimicrobials exposure, mechanical ventilation, pulmonary disease, current stay in the ICU and use of indwelling device as the most prominent ones. Also the horizontal transmission of plasmid mediated carbapenem resistance among different isolates may contribute to the high prevalence of carbapenem resistance in the hospital.

The presence of carbapenem-resistant isolates with negative carbapenemases results may be attributed to other causes as efflux pumps, drug impermeability or production of carbapenemases Class-D that can’t be detected by mCIM method (Benmahmod et al.) [28]. With the increased emergence and spread of carbapenem resistance, treatment options are decreasing.

Minimum inhibitory concentration for colistin was done by broth microdilution method. Forty three isolates out of 116 (37.1%) carbapenem-resistant isolates were colistin resistant (Non-wild type). A study done by Capone et al. [29] reported 36.1% colistin resistance among K.pneumoniae carbapenem-resistant isolates. Lower levels of colistin resistance were found in other studies as Rojas et al. [30] and Bhaskar et al. [31] detected 13% and 27.7% colistin resistant isolates among carbapenem-resistant K.pneumoniae isolates respectively. Higher result of colistin resistance was detected in a study by Monaco et al. [32] as they reported 43% colistin resistant isolates. In Alexandria, Egypt, a study done by Shawky et al. [28] reported 14% colistin resistance among carbapenem resistant isolates. Many risk factors were identified for colistin resistance in GNRs, including recent prior hospitalization, prior carbapenem resistance, prior treatment with colistin, exposure to chlorhexidine, the presence of multiple comorbidities in the patients, increasing age, male sex, length of hospitalization, and the presence of indwelling urinary catheters and all of these factors present in our hospital.

In this study the 43 colistin and carbapenem resistant E.coli and K.pneumoniae isolates were tested by conventional PCR for detection of plasmid mediated colistin resistance genes mcr-1 and mcr-2. Two isolates harbor mcr-2 gene (4.6%), meanwhile, only one isolate has mcr-1 gene (2.3%) were detected among the 43 isolates. A study done by Moosavian et al. [33] reported 1.7% mcr-1 gene. Higher prevalence of mcr-1 gene detection were found in a study done in Egypt by Zaki et al. [34] as they reported 4% mcr-1 genes. Regarding mcr-2 gene Zhang et al. [35] and Jayol et al. [36] reported mcr-2 gene with prevalence of 1.5% and 1% respectively. The presence of the plasmid mediated colistin resistant gene in the current study may be due to previous antibiotic therapy (e.g. colistin, beta-lactam/beta-lactamase inhibitors, carbapenems, glycopeptides), previous hospitalisation and previous colonisation with multidrug-resistant bacteria (e.g. KPC). Additional risk factors may be immunosuppression and the use of fluoroquinolones.

To the best of our knowledge by reviewing previous studies, plasmids that are harboring mcr
genes are different from plasmids carrying other antibiotic resistance genes. This may explain the high rate of antibiotic resistance found in our strains with the low rate of mcr genes detected. Low rates of horizontal transmission of plasmid-mediated colistin resistance genes mcr-1 and mcr-2 is still noticed in the hospital as the dissemination of mcr in K. pneumoniae is contingent on colistin selection pressures, and by extension, that by limiting colistin usage, further dissemination of the gene in K. pneumoniae could be stopped. In SCUH the usage of colistin in treatment of patients is still limited. In conclusion, this study revealed that the presence of colistin resistance among carbapenem resistant isolates in SCUHs is concerning because further limiting potential therapeutic options and inevitably leading to higher mortality rates. In addition the detection of the plasmid mediated mcr-1 and mcr-2 raises the concern and the necessity for paying attention to the horizontal transmission of this resistance which may eventually lead to pan-drug resistant isolates. Strict application of standard infection control practices in the hospitals especially in the critical areas are highly recommended to prevent the emergence and the spread of the resistant organisms. Also, we recommend the wise use of carbapenems, colistin and other antibiotics. 

Conflict of interest: The authors have no conflict of interest to declare.

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