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## Frequency of PmrA, PmrB and mcr-1 genes in pseudomonas aeruginosa isolates from cystic fibrosis patients of mofid children hospital, Tehran, Iran

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Introduction: The emergence of multidrug- resistant (MDR) in gram negative pathogens such as P. aeruginosa has become an important challenge to worldwide public health, recently (1). Colistin as a last line of treatment against MDR gram negative bacteria has been propounded (2). Recent studies have been demonstrated that colistin resistance is adaptive and regulated via PmrA and PmrB as a two component regulatory system (3). The plasmid borne colistin resistance gene mcr-1 was primarily recognized from China (4). Due to the raising of antibiotic resistance, in this study prevalence of mcr-1 gene and the mutation in pmrA and pmrB genes in P. aeruginosa isolates from sputum of Cystic Fibrosis (CF) patients was investigated.

Methods: 41 isolates of P. aeruginosa from sputum of CF patients in Mofid Children Hospital during Apr-Sep 2017 were collected. According to CLSI guideline 2017, antibiotic susceptibility test (AST) was applied by using the disk diffusion method. Detection of pmrA, pmrB and mcr-1 genes was performed by PCR and further sequencing was administrated for finding the mutations.

Results: Among 41 isolates of P. aeruginosa 22 (53.65%) were resistant to Amikacin, 21 (51.21%) to Ofloxacin and Cefepime, 20 (48.78%) to Imipenem, 19 (46.34%) to Ceftazidime and Ciprofloxacin, 18 (43.9%) to Aztreonam, 17 (41.46%) to Piperacillin, 16 (39.02%) to Gentamicin, 14 (34.14%) to Pipracillin-Tazobactam and 1 (2.43%) to Colistin. PCR results showed that all of the isolates had *pmrA* and *pmrB* genes and all of them were negative for *mcr-1*. One of the isolates show some mutations in pmrB gene.

Conclusion: The results of this study showed that colistin is the best choice for treatment. Using the molecular tests is necessary and have a determinative role to prescription of antibiotics by physicians.

## **Biography**

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