

عنوان مقاله:

THE STUDY MOLECULAR OF OXA-48 AND DETERMINATION OF RESISTANCE PATTERN IN CLINICAL ISOLATE OF PSEUDOMONAS AERUGINOSA IN PATIENTS HOSPITALIZED IN BURNING WARD OF ERFAN HOSPITAL, TEHRAN

محل انتشار:

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نویسندگان:

Farid Tarafdar - Msc Student of Microbiology, Department of Microbiology, Ahar branch, Islamic Azad University, Ahar, Iran

Behboud Jafari - Assistant professor, Department of Microbiology, Ahar branch, Islamic Azad University, Ahar, Iran

خلاصه مقاله:

Background and Aim: *Pseudomonas aeruginosa* is a gram-negative, non-fermentative bacillus and one of the most common opportunistic human pathogen causing 10-15% of nosocomial and burn wound infections worldwide. In this study, the plenty of OXA-48 gene and antibiotic resistance of clinical specimens in isolates isolated from burn wound infection in patients hospitalized in the ward of Erfan hospital in Tehran was performed. **Methods:** In this study, 20 strains of *Pseudomonas aeruginosa* were isolated from patients in the burn ward of Erfan Hospital and identified by biochemical tests. Antibiotic resistance pattern was determined by disk diffusion method. The genotype of OXA-48 gene was evaluated by PCR method and analyzed by SPSS23 software. **Results:** The highest resistance to antibiotics Ceftriaxone and Cefotaxime (100%), Meropenem (93.75%), Cefepime (89.55%), Amikacin (85.40%) And the lowest resistance to other antibiotics was observed for ceftazidime (55%). In genotypic study, 14 strains (68%) of OXA-48 gene were identified. **Conclusion:** The results show that most of the samples are resistant to the drug and OXA-48 genes were observed among the strains. Therefore, rapid measurement and accurate examination of antibiotic resistance is essential.

کلمات کلیدی:

Pseudomonas aeruginosa, Antibiotic Resistance, OXA-48

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