عنوان مقاله:

Prevalence of Virulence Genes and Antibiotic Resistance Pattern in Enterococcus Faecalis Isolated from Urinary Tract Infection in Shahrekord, Iran

محل انتشار:

مجله گزارش های بیوشیمی و زیست شناسی مولکولی, دوره 10, شماره 1 (سال: 1400)

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نویسندگان:

Roya Jafarzadeh Samani - Department of Microbiology, Faculty of Basic Science, Shahrekord Branch, Islamic Azad .University, Shahrekord, Iran

Elahe Tajbakhsh - Department of Microbiology, Faculty of Basic Science, Shahrekord Branch, Islamic Azad .University, Shahrekord, Iran

Hassan Momtaz - Department of Microbiology, Faculty of Basic Science, Shahrekord Branch, Islamic Azad University, .Shahrekord, Iran

Mohsen Kabiri Samani - Internal Medicine Department, Shahrekord University of Medical Siences. Shahrekord, Iran & .Internal Medicine Department, Shahrekord University of Medical Sciences. Shahrekord, Iran

خلاصه مقاله:

Background: This study aims to specify the antimicrobial resistance pattern and virulence genes of Enterococcus faecalis isolated from urinary tract infections in Shahrekord, Iran. Methods: Urine samples of 1000 people suspected of having urinary tract infections referred to Shahrekord medical diagnostic laboratories were examined. Biofilm assays were performed by microtiter plate test through reading the ODF9. Polymerase Chain Reaction (PCR) was applied to study the virulence factors. Results: Enterococcus faecalis was detected in 50 samples. After performing microbiological tests, all samples were positive in the molecular analysis. Strong, moderate and weak biofilm reactions reported ۶۶.۶۷%, Υ۵%, and Λ.٣٣% respectively. The most resistance reported to cotrimoxazole, vancomycin and amikacin and the lowest resistance to nitrofurantoin (٨.٣٣%) was reported. Statistical analysis with Fisherchr('٣٩')s exact test showed a statistically significant relationship between biofilm production and resistance to cotrimoxazole, vancomycin and cefotaxime. Prevalence of efe A, ace, gel E, esp, cyl M, agg, cyl A and cyl B in strong biofilm formation isolates was reported ۱۰۰%, ۸۲.۵%, ۸۲%, ۶۲.۵%, ۵۵%, ۳۷.۵% ۲۵% and ۲۲.۵% respectively. There was a significant relationship between the frequency of efa A and strong biofilm reaction. Conclusions: The presence of E. faecalis strains resistant to co-trimoxazole and vancomycin and present of some virulence factors is alarming the researchers. Since antibiotic resistance genes are probably transmitted among enterococci, and Staphylococci, controlling infections made by enterococci as well as the appropriate administration of antibiotics could treat the .nosocomial infections effectively

کلمات کلیدی:

.Antibiotic Resistance, Enterococcus faecalis, Urinary Tract Infection, Virulence genes

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