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### **Original Research Article**

# Study of Antimicrobial Resistance among Enterococcus Species in Tertiary Care Teaching Hospital

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### **Abstract**

The current study was carried out to isolate and identify Enterococcus species from clinical samples and to analyse the antimicrobial susceptibility pattern of Enterococcus Species. All clinical samples which were received in Microbiology laboratory were included in the study. A total of 2958 samples were received. These 129 Enterococci isolates were obtained over a period of 1 year from various samples namely, urine, blood, pus. Samples were processed for microscopy and culture as per the standard guidelines. all samples were inoculated on MacConkey agar and Blood agar. Identification and antibiotic sensitivity pattern of bacterial isolates were carried out by automated system. i.e. VITEK 2 Compact. Out of 2958 samples, Total 129 samples showed growth of enterococci. Out of 129 isolates, 84 were Enterococcus spp, 18 were E.faccalis, 16 were E.faccium. Out of total 129 samples, 76 samples were of males and 53 were of females. Most common samples from which enterococci were isolated was urine accounting for 85 followed by Pus accounting for 15. Medicine department showed the highest isolates with 88. Out of 129 Enterococci 64 showed resistance to primary line of drug like Benzyl penicillin, 79 Enterococci showed resistance to Ampicillin. Out of 129 enterococci 70 showed resistance to secondary line of drug like tetracycline,59 isolates showed resistance to High level Gentamycin, 41 isolates showed resistance to Vancomycin,23 isolates showed resistance to Linezolid. The strains of Enterococci were multidrug resistant. Anitibiotic selection should be done judiciously based on the antibiotic sensitivity pattern of the organism.

Keywords: Enterococcus species, Antibiotic resistance, Antimicrobial susceptibility pattern.

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#### Introduction

Enterococci are natural inhabitants of the oral cavity, gastrointestinal tract (GIT) and the female genital tract in both humans and animals [1]. Enterococci, though commensals in adult faeces are important nosocomial pathogens [2]. The most common nosocomial infections by these organisms are urinary tract infections followed by intraabdominal and pelvic infection [3]. Enterococci are also becoming increasing important agents of human disease because of their resistant to antibiotics. E. faecalis is most common species found in clinical specimen whereas E. faecium is more drug resistant than E.faecalis [4]. Enterococci resistant to all three antimicrobial agents (penicillin, aminoglycosides and glycopeptide like vancomycin) pose a serious challenge not only for clinicians but also

for health care institutions. It results in treatment failure, selection and spreading of resistant strains in the health care institution. Imprudent use of antibiotics and colonisation pressure are the important causes of the drug resistance in enterococci [5]. Resistance in enterococci has been increasingly reported especially vancomycin resistant enterococci (VRE). Newer antibiotics such as linezolid, daptomycin and tigecycline have shown good in vitro activity against VRE [7]. Quinupristin-dalfopristin (Q/D) is another agent that has potent in vitro activity against E. faecium but poor activity against E. faecalis [6].

## MATERIAL AND METHODS

This study was carried out at C.U. Shah medical college and hospital, Surendranagar, over a period Of January 2019 to December 2020. In this

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study, antimicrobial susceptibility pattern enterococci has been evaluated. All clinical samples which were received in Microbiology laboratory were included in the study. A total of 2958 samples were received. These 129 Enterococci isolates were obtained over a period of 1 year (2019 -2020) from various samples namely, urine, blood, pus, swab and fluid. Samples were processed for microscopy and culture as per the standard guidelines. For microscopy, Gram stain was performed. For culture, all samples were inoculated on MacConkey agar and Blood agar and incubated at 37°C for 24 hours on day of sample collection. Samples which showed growth on MacConkey agar and Blood agar, identification of the colonies was done by observing its macroscopic characteristics and by Gram stain. On MacConkey agar magenta pink, small, round

colonies were considered as an Enterococci. On blood agar non hemolytic translucent considered as Enterococci. Further identification was done using gram stain which showed gram positive arranged. Results were analyzed using Microsoft Excel.

The ethical clearance for the study was obtained from the Institutional Ethics Committee prior to the study.

#### RESULTS

Total 129 samples showed growth of enterococci out of 2958 samples. Out of total 129 samples, 76 samples were of males and 53 were of females.

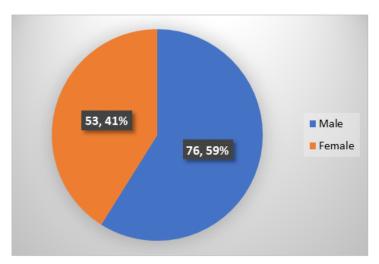


Fig 1: Gender wise distribution of patients

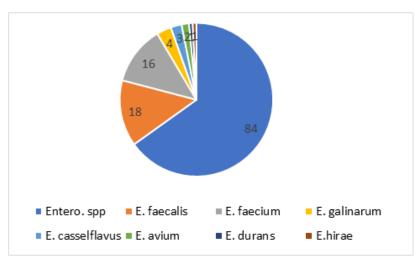


Fig 2: Species wise detection of organisms

Out of 129 isolates, 84 were Enterococcus spp, 18 were E. faecalis, 16 were E.faceium, 4 were E.

gallinarum, 3 were E.casseliflavus, 2 were E.avium, 1 was E.durans, 1 was E.hirae.

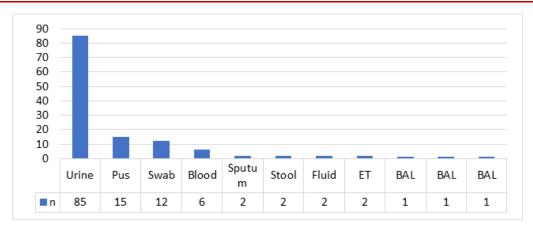


Fig 3: Clinical Sample wise Detection of organisms

Most common samples from which enterococci were isolated was urine accounting for 85

followed by Pus accounting for 15, followed by swab accounting for 12, Blood accounting for 6.

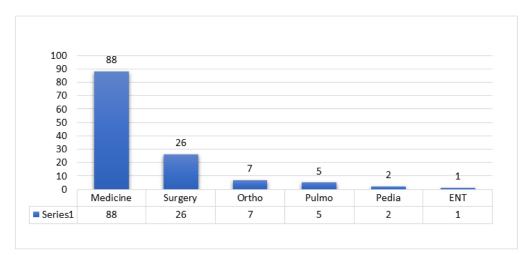


Fig 4: Department wise detection of organisms

Medicine department showed the highest isolates with 88 followed by surgery department showed 26 isolates, orthopedics department showed 7

isolates and pulmonology department showed 5 isolates.

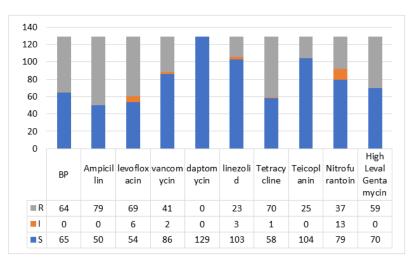


Fig 5: Antibiotic resistance pattern of Enterococci

Out of 129 Enterococci 64 showed resistance to primary line of drug like Benzyl penicillin, 79 Enterococci showed resistance to Ampicillin, 69 Enterococci showed resistance to Levofloxacin. Out of 129 enterococci 70 showed resistance to secondary line of drug like tetracycline,59 isolates showed resistance to High level Gentamycin, 41 isolates showed resistance to Vancomycin, 23 isolates showed resistance to Linezolid and 25 isolates showed resistance to Teicoplanin, 37 isolates showed resistance to Nitrofurantoin. All Enterococci were sensitive to Daptomycin, PIPTAZ, Ampicillin sulbactam.

# **DISCUSSION**

In this study Total 129 samples showed growth of Enterococci. Out of which 84 were enterococcus spp., 18 were E. faecalis, 16 were E.faecium, 4 were E. gallinarum. 3 were E. casseliflavus, 2 were E.avium, 1 was E. durans and 1 was E.hirae. Elham Jannati et al., reported 118 enterococcus spp., 235 E.faecium, 56 E.faecalis. in this study 84 Enterococcus spp. Isolated followed by E. faecalis (18) out of 409 isolates. Barreto et al. and Poeta et al., which showed that E.faecium accounting for >50% of Enterococcal isolates recovered from the patients. Most of the isolates from Urine (56) followed by Pus (15) and Swab (12). Kaarthiga at el reported that out of 310 sample, most of the isolates from Urine (68) followed by Pus (16) and Blood (14). Globally Ampicillin resistance is significantly high in clinical isolates. In our study 79 isolates showed Ampicillin resistance. Elham Jannati et al., reported a small number of isolates (10/409). Out of 129, 41 isolates showed resistance to Vancomycin and 59 isolates showed resistance to High level gentamycin. Kaarthiga et al., reported Vancomycin resistance was seen in (29/310) isolates and HLG (69/310) isolates. Jacopo Monticelli et al., reported that E.gallinarum and E.casseliflavus are intrinsically resistant Vancomycin. In Our study (3/4) E. gallinarum and (1/3) E.casseliflavus strain showed resistance Vancomycin. Out of 129, 23 isolates showed resistance to Linezolid. A. Raddaoui et al., reported 2 (out of 438) strain only were resistant to Linezolid.

#### CONCLUSION

Enterococci was isolated and identified in 129 samples and the most common sample from which enterococci was isolated was urine. The strains of Enterococci were multidrug resistant.

Multidrug resistance of Enterococci indicates that antibiotic selection should be done judiciously based on the antibiotic sensitivity pattern of the organism.

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