- 1 RESISTANCE AND VIRULENCE PROFILES AND CYTOTOXIC POTENTIAL OF Enterococcus
- 2 sp. ISOLATES FROM AQUATIC ENVIRONMENT

3

- 4 FURLANETO-MAIA, L.; MIKALOUSKI, U.; GOMES-BRAVO, G.B; GONÇALVES, A.B; FOGAÇA, M.C.M;
- 5 HOGOS, S.L. M; MORALEZ, A.T; FURLANETO, M. C.

6

- 7 Universidade Tecnológica Federal do Paraná-Campus Londrina (Av dos Pioneiros, 3131, cep
- 8 86036-370, Londrina-PR); Universidade Estadual de Londrina (Rodovia Celso Garcia Cid, PR
- 9 445, cep 86057-970, Londrina PR).

10 11

12

13

14

15

16

17

18

19

20

21

22

23

24

25

26

27

28 29

30

31

32

33

34

35

ABSTRACT:

Enterococcus spp. are ubiquitous in nature and common inhabitants of the microbiota of the gastrointestinal tract of humans and animals and are often associated with opportunistic infections. Such infections are often associated with polymicrobial biofilms and antimicrobial resistance. In recent decades unusual species of enterococcus have been described. Therefore, the understanding of the cytotoxic potential of species of the genus as well as their virulence factors are important for the identification of new strategies to control infections caused by these microorganisms. Our study aimed to evaluate the presence of vancomycin resistance genes (van A e van B), the presence of genes related to virulence factors such as gelE (production de gelatinase), ace (Enterococcus collagen adhesin), cyIA (Cytolysin-activating bacteriocins), sprE (serine endopeptidase), PAI (Island of pathogenicity), asa1 (aggregation substance), cpd (sex pheromone) e agg (aggregation substance) as well as to evaluate the cytotoxic potential of Enterococcus sp. from aquatic environments. In addition, the isolates were evaluated for the antibiotic resistance profile and the ability to form biofilms. As for the resistance genes, we observed that 8,33 % and 33,33% of the isolates had the vanA and vanB genes, respectively. As for virulence genes, frequencies of 41,66% for gelE and ace; 25 % for cylA and sprE; 33,33 % and 50% for asa1 and agg, respectively. None of the isolates analyzed showed the cpd and PAI genes. As for the phenotype of antimicrobial resistance, 66.66% of the isolates presented resistance to vancomycin and 58% to ampicillin. An isolate of E. casseliflavus (14) presented resistance to all tested antibiotics. All isolates showed metabolites with cytotoxic potential against Hela cells. The ability to form biofilm was observed in all isolates and 50% of these were classified as strongly biofilm forming. Of these, 66% had resistance to more than five of the antibiotics tested. The present study demonstrates that different species of Enterococcus sp. can present different profiles of resistance and virulence, which reinforces the importance of the studies of these profiles in *Enterococcus* sp. from the most varied environments.

36

- 37 Key words: E. casseliflavus, E. mundtii, HeLa cells, genotype, fenotype, environmental
- 38 Development Agency: CNPQ/Fundação Araucária/UTFPR