

## Frequency detection of *Enterococcus faecalis* and *Enterococcus faecium* infection and antibiotic resistance pattern in diarrheic dogs in Ahvaz district

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### Abstract

Enterococci are a part of the opportunistic pathogens, which are very important in medicine. These bacteria can cause a variety of diseases in both dogs and human. The aim of the present study was to investigate the frequency detection of *Enterococcus faecalis* and *Enterococcus faecium* in companion dogs in Ahvaz and review of risk factors including age, gender, breed and diarrheic status in animals. Also, the prevalence of virulence genes was evaluated including gelatinase (*gelE*) and (*ccf*) and antibiotic susceptibility measured in obtained samples. Sampling was performed from the rectum of the 150 dogs (36 cases diarrheic and 114 non-diarrheic). The samples were evaluated by two methods of bacterial culture and PCR. In bacterial culture, 122 isolates, suspected to *Enterococcus* species, were isolated and subsequently the detection of *SodA* gene specific to *Enterococcus faecalis* and *Enterococcus faecium* was performed by PCR. Overall forty-five positive isolates were identified, which thirty four of which were *Enterococcus faecalis* (75.5%) and 11 were *Enterococcus faecium* (24.5 %). In regard to identify virulence genes (*gelE* and *ccf*), 36 out of 45 isolates were positive for virulence genes. Twenty six isolates (57.77%) had virulence genes, 5 isolates (11.11%) *ccf* gene and 5 other isolates (11.11%) *gelE* gene. In all, nine isolates (20%) had no virulence gene. Fourteen different antibiotics were used to determine antibiotic susceptibility that indicated all isolates were resistant to azithromycin, streptomycin, ampicillin and imipenem. Thereafter, the highest resistance was related to erythromycin and cephalexin (95.5%), trimethoprim sulfamethoxazole (84.4%) and gentamicin (80%), respectively. Also, the highest sensitivity was related to nitrofurantoin (62.2%), penicillin G (60%) and enrofloxacin (55.5%), respectively. There was no significant relationship between risk factors such as age, gender, breed and diarrheic condition with the presence of *Enterococcus* in the studied dogs ( $P>0.05$ ). The results showed that the prevalence of *Enterococci* was relatively significant (30%) in dogs of Ahvaz district. Antibiotic resistance was significant in the two species of Enterococci. Finally, because of the very high importance of antibiotic resistance, appropriate administration of antibiotics is recommended.

**Key words:** *Enterococcus*, *Faecalis*, *Faecium*, Virulence gene, Antibiotic resistance

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## References

- Bang, K., An, J. U., Kim, W., Dong, H. J., Kim, J., & Cho, S. (2017). Antibiotic resistance patterns and genetic relatedness of *Enterococcus faecalis* and *Enterococcus faecium* isolated from military working dogs in Korea. *Journal of Veterinary Science*, *18*(2), 229-236.
- Ben Said, L., Dziri, R., Sassi, N., Lozano, C., Ben Slama, K., Ouzari, I., Torres, C., & Klibi, N. (2017). Species distribution, antibiotic resistance and virulence traits in canine and feline enterococci in Tunisia. *Acta Veterinaria Hungarica*, *65*(2), 173-184.
- Bertelloni, F., Salvadori, C., Lotti, G., Cerri, D., & Ebani, V. V. (2017). Antimicrobial resistance in *Enterococcus* strains isolated from healthy domestic dogs. *Acta Microbiologica et Immunologica Hungarica*, *64*(3), 301-312.
- Bunt, G. van den., Top, J., Hordijk, J., Greeff, S. C de., Mughini-Gras, L., Corander, J., Pelt, W. van., Bonten, M. J. M., Fluit, A. C., & Willems, R. J. L. (2018). Intestinal carriage of ampicillin- and vancomycin-resistant *Enterococcus faecium* in humans, dogs and cats in the Netherlands. *Journal of Antimicrobial Chemotherapy*, *73*(3), 607-614.
- Ferguson, D. M., Talavera, G. N., Hernández, L. A., Weisberg, S. B., Ambrose, R. F., & Jay, J. A. (2016). Virulence genes among *Enterococcus faecalis* and *Enterococcus faecium* isolated from coastal beaches and human and nonhuman sources in Southern California and Puerto Rico. *Journal of Pathogens*, *1*, 1-7.
- Frye, J. G., & Jackson, C. R. (2013). Genetic mechanisms of antimicrobial resistance identified in *Salmonella enterica*, *Escherichia coli*, and *Enterococcus* spp. isolated from US food animals. *Frontiers in Microbiology*, *4*, 1-22.
- Golob, M., Pate, M., Kušar, D., Dermota, U., Avberšek, J., Papić, B., & Zdovc, I. (2019). Antimicrobial Resistance and virulence genes in *Enterococcus faecium* and *Enterococcus faecalis* from humans and retail red meat. *BioMed Research International*, *1*, 1-12
- Greene, C. E., & Prescott, J. F. (2012). *Enteric bacterial infections*. In: *Infectious diseases of the Dog and Cat*. Greene. Vol. 1. (4th Edition). St. Louis Missouri, USA. Pp: 333-334.
- Gulhan, T., Boynukara, B., Ciftci, A., Sogut, M. U., & Findik, A. (2015). Characterization of *Enterococcus faecalis* isolates originating from different sources for their virulence factors and genes, antibiotic resistance patterns, genotypes and biofilm production. *Iranian Journal of Veterinary Research*, *16*(3), 261-266.
- Hammerum, A. M. (2012). Enterococci of animal origin and their significance for public health. *Clinical Microbiology and Infection*, *18*(7), 619-625.
- Iseppi, R., Messi, P., Anacarso, I., Bondi, M., Sabia, C., Condò, C., & de Niederhausern, S. (2015). Antimicrobial resistance and virulence traits in *Enterococcus* strains isolated from dogs and cats. *New Microbiologica*, *38*, 369-378.
- Jackson, C. R., Fedorka-Cray, P. J., & Barrett, J. B. (2004). Use of a genus- and species-specific multiplex PCR for identification of enterococci. *Journal of Clinical Microbiology*, *42*(8), 3558-3565.
- Jackson, C. R., Fedorka-Cray, P. J., Davis, J. A., Barrett, J. B., & Frye, J. G. (2009). Prevalence, species distribution and antimicrobial resistance of enterococci isolated from dogs and cats in the United States. *Journal of Applied Microbiology*, *107*(4), 1269-1278.
- Kajihara, T., Nakamura, S., Iwanaga, N., Oshima, K., Takazono, T., Miyazaki, T., Izumikawa, K., Yanagihara, K., Kohno, N., & Kohno, S. (2015). Clinical characteristics and risk factors of enterococcal infections in Nagasaki, Japan: a retrospective study. *BMC Infectious Diseases*, *15*(1), 1-8.
- Marra, A., Dib-Hajj, F., Lamb, L., Kaczmarek, F., Shang, W., Beckius, G., Milici, A. J., Medina, I., & Gootz, T. D. (2007). Enterococcal virulence determinants may be involved in resistance to clinical therapy. *Diagnostic Microbiology and Infectious Disease*, *58*(1), 59-65.
- Nazarian-Firouzabadi, F., & Akrami, M. J. (2019). Antibigram analysis and tracking of the virulence-related genes in *Enterococcus faecalis* isolates. *Scientific Journal of Kurdistan University of Medical Sciences*, *24*(2), 17-29.
- Oliveira, M., Tavares, M., Gomes, D., Touret, T., São Braz, B., Tavares, L., & Semedo-Lemsaddek, T. (2016). Virulence traits and antibiotic resistance among enterococci isolated from dogs with periodontal disease. *Comparative Immunology, Microbiology and Infectious Diseases*, *46*, 27-31.

- Poeta, P., Costa, D., Rodrigues, J., & Torres, C. (2006). Antimicrobial resistance and the mechanisms implicated in faecal enterococci from healthy humans, poultry and pets in Portugal. *International Journal of Antimicrobial Agents*, 27(2), 131-137.
- Sattari-Maraji, A., Jabalameli, F., Farahani, N.N., Beigverdi, R., & Emaneini, M. (2019). Antimicrobial resistance pattern, virulence determinants and molecular analysis of *Enterococcus faecium* isolated from children infections in Iran. *BMC Microbiology*, 19, 1-8.
- Sharifi, Y., Hasani, A., Ghotaslou, R., Naghili, B., Aghazadeh, M., Milani, M., & Bazmany, A. (2013). Virulence and antimicrobial resistance in enterococci isolated from urinary tract infections. *Advanced Pharmaceutical Bulletin*, 3(1), 197-201.
- Soodmand, J., Zeinali, T., Kalidari, G., Hashemitabar, G. H., & Razmyar, J. (2018). Antimicrobial susceptibility profile of *Enterococcus* species isolated from companion birds and poultry in the Northeast of Iran. *Archives of Razi Institute*, 73(3), 207-213.
- Wood, M. W., Lepold, A., Tesfamichael, D., & Lasarev, M. R. (2020). Risk factors for enterococcal bacteriuria in dogs: A retrospective study. *Journal of Veterinary Internal Medicine*, 34(6), 2447-2453.
- Zhou, L., Sun, H., Song, Sh., Liu, J., Xia, Zh., Sun, Y., & Lyu, Y. (2019). H3N2 canine influenza virus and *Enterococcus faecalis* coinfection in dog's in China. *BMC Veterinary Research*, 15, 1-6.