

Editorial

Special Issue—Resistant Staphylococci in Animals

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Staphylococci figure prominently among those bacteria demonstrating antimicrobial resistance (AMR) and are thus responsible for significant problems concerning the treatment of the animals and humans that they infect. In particular, livestock-associated methicillin-resistant *Staphylococcus aureus* (LA-MRSA) and methicillin-resistant *Staphylococcus pseudintermedius* (MRSP) are of growing concern. *Staphylococcus schleiferi* and coagulase-negative staphylococci occur as commensals of the skin and mucous membranes of animals. However, they have been implicated in a variety of infections and are frequently resistant to one or more antimicrobial classes. This Special Issue assembles a collection of original articles that shed further light on these fascinating bacteria and their potential impacts on both veterinary medicine and public health.

The review by Lynch and Helbig [1] provides a timely update of our knowledge regarding *Staphylococcus pseudintermedius*, the most common pathogen isolated from skin disease samples (particularly pyoderma) from dogs [2]. The high prevalence of methicillin-resistance and associated resistance to many other antimicrobials in isolates of *S. pseudintermedius* constitutes a growing concern. A range of novel potential treatments including vaccination and phage therapy are discussed.

Chanayat et al. [3] investigated the staphylococcal cassette chromosome *mec* (SCC*mec*) type and the antimicrobial susceptibility of staphylococci isolated from superficial pyoderma infections affecting dogs in Thailand. SCC*mec* type V was found in *S. aureus*, the *S. intermedius* group, *S. lentus*, *S. xylosus*, and *S. arlettae*, and although the authors do not state whether the coagulase-negative species were considered the primary bacterial pathogens in the cases from which they were isolated, the need to reduce environmental contamination and educate veterinary personnel and clients about the potential for the transmission to and from dogs of all resistant staphylococci is emphasized. The need for hygiene is supported by the results of another recent longitudinal study of MRSP-infected dogs conducted by Frosini et al. [4], in which it was reported that 45% of households were contaminated with MRSP. The data presented by Chanayat et al. [3], includes two *S. pseudintermedius* isolates that showed MICs < 2 µg/ml, an interesting finding in light of the publication by Wegener et al. [5] suggesting that such isolates showing low-level resistance may be susceptible to treatment with higher doses of beta-lactams or more frequent administration.

As well as characterizing *S. pseudintermedius* isolates causing superficial pyoderma in Taiwanese dogs, Lai et al. [6] also examined the risk factors that might lead to owners acquiring *S. pseudintermedius* from their pets. Although no significant association was found, high odds ratios were obtained for “keeping three or more dogs” and “dogs can lick the owner’s face”, suggesting support for recent publications describing the potential for *S. pseudintermedius* infections in human hosts [7,8].

Certain foods and food production systems may present a pathway for the transmission of MRSA to humans. Benrabia et al. [9] detected MRSA in 30% of poultry farms in Algeria and provided a reasonable argument for conducting interventions to reduce spread between farms and mitigate the contamination of the food chain. The lack of the molecular



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typing of isolates was a limitation of this study, as it largely precluded source attribution and comparison with MRSA isolates reported worldwide. Nevertheless, the documentation of MRSA isolation rates in food-producing animal species in different geographical regions is essential to obtain a comprehensive understanding of this public health challenge.

Staphylococcus aureus is still the leading cause of bovine mastitis in many countries. Rusenova et al. [10] compared phenotypic and genotypic methods in their evaluation of AMR among bovine mastitis isolates of *S. aureus* in Bulgaria. The discrepancies detected for some isolates are concerning and, as recommended by the authors, highlight the need for isolates to be thoroughly characterized.

Biofilm production is considered an important virulence factor for *S. aureus* [11] and coagulase-negative staphylococci (CoNS), which both cause bovine mastitis. Lee and Lee [12] showed that the prevalence of multi-drug resistance (MDR) was significantly higher in strong or moderate biofilm-producing CoNS than in those that were weak or non-formers in samples isolated from normal bulk tank milk in Korea. As biofilm production is associated with persistence of infection and antimicrobial therapy may not eliminate infection, the authors advise effective monitoring and sanitation programs to prevent the contamination of equipment with environmental CoNS, thus limiting the opportunities for cows to be infected with these organisms.

The diverse articles contained in this Special Issue constitute a valuable contribution to our understanding of staphylococcal infection in both farm and companion animals.

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References

1. Lynch, S.A.; Helbig, K.J. The complex diseases of *Staphylococcus pseudintermedius* in canines: Where to next? *Vet. Sci.* **2021**, *8*, 11. [[CrossRef](#)] [[PubMed](#)]
2. Loeffler, A.; Lloyd, D.H. What has changed in canine pyoderma? A narrative review. *Vet. J.* **2018**, *235*, 73–82. [[CrossRef](#)] [[PubMed](#)]
3. Chanayat, Y.; Akatvipat, A.; Bender, J.; Punyaapornwithaya, V.; Meeyam, T.; Anukool, U.; Pichpol, D. The SCCmec Types and antimicrobial resistance among methicillin-resistant *Staphylococcus* species isolated from dogs with superficial pyoderma. *Vet. Sci.* **2021**, *8*, 85. [[CrossRef](#)] [[PubMed](#)]
4. Frosini, S.M.; Bond, R.; King, R.; Feudi, C.; Schwarz, S.; Loeffler, A. Effect of topical antimicrobial therapy and household cleaning on methicillin-resistant *Staphylococcus pseudintermedius* carriage in dogs. *Vet. Rec.* **2022**, *190*, e937. [[CrossRef](#)] [[PubMed](#)]
5. Wegener, A.; Damborg, P.; Guardabassi, L.; Moodley, A.; Mughini-Gras, L.; Duim, B.; Wagenaar, J.A.; Broens, E.M. Specific staphylococcal cassette chromosome mec (SCCmec) types and clonal complexes are associated with low-level amoxicillin/clavulanic acid and cefalotin resistance in methicillin-resistant *Staphylococcus pseudintermedius*. *J. Antimicrob. Chemother.* **2020**, *75*, 508–511. [[CrossRef](#)] [[PubMed](#)]
6. Lai, C.-H.; Ma, Y.C.; Shia, W.-Y.; Heieh, Y.-L.; Wang, C.-M. Risk factors for antimicrobial resistance of *Staphylococcus* species isolated from dogs with superficial pyoderma and their owners. *Vet. Sci.* **2022**, *9*, 306. [[CrossRef](#)] [[PubMed](#)]
7. Somayaji, R.; Priyantha, M.A.; Rubin, J.E.; Church, D. Human infections due to *Staphylococcus pseudintermedius*, an emerging zoonosis of canine origin: Report of 24 cases. *Diagn. Microbiol. Infect. Dis.* **2016**, *85*, 471–476. [[CrossRef](#)]
8. Ference, E.H.; Danielian, A.; Kim, H.W.; Yoo, F.; Kuan, E.C.; Suh, J.D. Zoonotic *Staphylococcus pseudintermedius* sinonasal infections: Risk factors and resistance patterns. *Int. Forum. Allergy Rhinol.* **2019**, *9*, 724–729. [[CrossRef](#)]
9. Benrabia, I.; Hamdi, T.; Shehata, A.; Neubauber, H.; Wareth, G. Methicillin-resistant *Staphylococcus aureus* (MRSA) in poultry species in Algeria: Long term study of prevalence and antimicrobial resistance. *Vet. Sci.* **2020**, *7*, 54. [[CrossRef](#)] [[PubMed](#)]
10. Rusenova, N.; Vasilev, N.; Rusenov, A.; Milanova, A.; Sirakov, I. Comparison between some phenotypic and genotypic methods for assessment of antimicrobial resistance trend of bovine mastitis *Staphylococcus aureus* isolates from Bulgaria. *Vet. Sci.* **2022**, *9*, 401. [[CrossRef](#)] [[PubMed](#)]

11. Campos, B.; Pickering, A.C.; Rocha, L.S.; Aguilar, A.P.; Fabres-Klein, M.H.; de Oliveira Mendes, T.A.; Fitzgerald, J.R.; de Oliveira Barros Ribon, A. Diversity and pathogenesis of *Staphylococcus aureus* from bovine mastitis: Current understanding and future perspectives. *BMC Vet. Res.* **2022**, *18*, 115.
12. Lee, Y.J.; Lee, Y.J. Characterization of biofilm producing coagulase-negative staphylococci isolated from bult tank milk. *Vet. Sci.* **2022**, *9*, 430. [[CrossRef](#)] [[PubMed](#)]

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