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# Revealing the clinical relevance of Staphylococcus borealis

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ABSTRACT Staphylococcus borealis, previously misidentified as Staphylococcus haemolyticus, was first described as a new species in 2020. In this study, we aimed to describe the clinical relevance of S. borealis by combining clinical data, antibiotic susceptibility profiles, and biofilm formation in isolates obtained from hospitalized and non-hospitalized patients. We established a collection of 129 S. borealis isolates from 129 adult patients from seven Norwegian hospitals. We describe clinical data at the time of microbiological specimen collection. Antibiotic susceptibility and biofilm formation were tested using established methods. Of 129 isolates (37%), 48 were from patients admitted to a hospital, the remaining from outpatients. The median (IQR) age was 62 (51-78) years, and 85/129 (66%) of the isolates were from male patients. The majority (81/129, 63%) of the S. borealis isolates were isolated from urine cultures, followed by isolation from skin and soft tissue cultures (35/129, 27%), blood cultures (8/129, 6%), and two implant-associated infections (2/129, 2%). Resistance to ≥3 antibiotic classes was observed in 43/129 (33%) of the isolates. All isolates formed a biofilm under the conditions tested; 59/129 (46%) weak, 40/129 (31%) medium, and 29/129 (23%) were strong biofilm producers. S. borealis clinical samples were predominately obtained from elderly male patients, and the majority of samples were from patients with suspected urinary tract or skin and soft tissue infections. The level of multidrug resistance was comparable to other coagulase-negative staphylococcal species, but resistance toward methicillin and penicillin was lower than in clinical S. haemolyticus isolates.

**IMPORTANCE** This study contributes novel knowledge on the clinical relevance of *Staphylococcus borealis*; this is of importance when clinical microbiologists encounter *S. borealis* identified in patient samples. *S. borealis* was mainly identified in microbiological specimens from middle-aged to elderly patients, predominantly males. Hospitalized patients were also frequently immunocompromised and had other underlying conditions accompanying a suspected *S. borealis* infection.

**KEYWORDS** *Staphylococcus borealis*, urinary tract infections, skin and soft tissue infections, multidrug resistance, biofilm formation, clinical relevance

oagulase-negative staphylococci (CoNS) is a heterogeneous group of species within the genus *Staphylococcus*. They are a major cause of opportunistic infection in immunocompromised individuals and are associated with biofilm formation on indwelling medical devices. However, there are differences in both virulence and antimicrobial resistance (AMR) patterns within the CoNS group (1). *Staphylococcus borealis* was previously misidentified as *Staphylococcus haemolyticus*. However, based on 16S rRNA gene sequencing and matrix-assisted laser desorption ionization time-of-flight mass spectrometry (MALDI-TOF MS) analyses, marked differences between *S. haemolyticus* and *S. borealis* isolates were detected. Subsequently, following whole genome sequence analyses, *S. borealis* was described as a new member of the CoNS group

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in 2020. *S. borealis* is genotypically distinct from other CoNS, has a yellow-pigmented phenotype, and produces urease (2).

Some CoNS are associated with specific infection sites, like the urinary tract (*Staphylococcus saprophyticus*), or invasive infections, like endocarditis (*Staphylococcus lugdunensis*). Two recent publications have described that *S. borealis* may cause bovine mastitis (3, 4). Fecal carriage of *S. borealis* has also been reported in pigs (5) and in samples from healthy children (6). However, there is virtually no data on the clinical relevance of this new staphylococcal species in humans.

*S. haemolyticus* is ranked as the most antibiotic-resistant member of the CoNS group, and nosocomial infections are "difficult to treat" (7). Very little is known about AMR in *S. borealis*, but the first genomic analyses identified the presence of genes encoding resistance to multiple antimicrobial agents (2). Fecal isolates from humans (6) and pigs (5) were reported as multi-drug resistant (MDR), and the porcine isolates carried both SCCmec type V and the unusual macrolide resistance genes *ermT* and *erm 43* (5). The findings of *S. borealis* both in porcine and bovine samples suggest a zoonotic potential. However, the level of AMR in *S. borealis* isolates from humans is still unknown.

In 2021, the MALDI-TOF spectrum for *S. borealis* was incorporated in the updated Bruker database used by the MALDI biotyper. Clinical microbiologists will therefore now encounter *S. borealis* identified in patient samples. Without clinical background information, it is challenging to interpret the significance of finding *S. borealis*. In this study, we present data from a large (n = 129) collection of clinical *S. borealis* isolates identified from seven Norwegian hospitals. Our aim was to describe clinical data collected during infection episodes where *S. borealis* was detected and the results of detailed phenotypic studies of the antibiotic susceptibility and the biofilm-forming ability of *S. borealis*.

## **MATERIALS AND METHODS**

## Study design

This is a national multicenter study including seven clinical microbiology departments in Norwegian hospitals. We established a collection of 129 *S. borealis* isolates using both a retrospective and prospective approach. All positive blood culture isolates are routinely stored at the microbiological departments. In the retrospective part of the study, all clinical *S. haemolyticus* blood culture isolates that had been collected and stored between 2014 and 2021 in the participating hospitals were re-analyzed by MALDI-TOF MS (Bruker Daltonics).

We searched for isolates that, with the latest MALDI-TOF Bruker database update, 2021, would now be classified as *S. borealis* (score ≥2.0). In the prospective part, we searched for all *S. borealis* isolates identified between February 2021 and November 2022. All identified *S. borealis* isolates were transported to one central laboratory for further investigation. All *S. borealis* isolates, independent of them being considered as the main cause of infection or not, were stored for the purpose of this study.

#### Clinical data

Clinical data were collected from the medical information accompanying the specimen sent to the clinical microbiology laboratories. These data included age, sex, source of specimen, hospital ward for admitted patients or whether they were treated as outpatient, clinically suspected type of infection, and if antibiotic therapy had been initiated (Table S1). We included only patients  $\geq$  18 years of age, and all patient data were anonymized.

# Antimicrobial susceptibility testing (AST)

AST was performed for this study by the disk diffusion method according to the EUCAST guidelines v.14.0 (8, 9). The following panel of antimicrobial agents was tested:

fusidic acid (FD10), cefoxitin (FOX30), penicillin G (P1), erythromycin (E15), clindamycin (DA2), trimethoprim—sulfamethoxazole (SXT25), trimethoprim (TMP 5), nitrofurantoin (NFT 100), gentamicin (CN10), linezolid (LZD 10), ciprofloxacin (CIP 5), tetracycline (TE 30), and rifampicin (RD5) (Oxoid, Ireland). Vancomycin (0.016–256  $\mu$ g/mL) was tested using the broth microdilution method (Liofilchem, Italy), according to the manufacturer's guidelines. A cefoxitin inhibition zone <22 mm was interpreted as methicillin resistance, in line with EUCAST recommendations (10, 11). Multidrug resistance (MDR) was defined as resistance to  $\geq$ 3 categories of antimicrobial agents. We interpreted AST data for *S. borealis* using breakpoints for other CoNS, as suggested by EUCAST (10).

## **Biofilm formation**

The biofilm-forming ability of the isolates was examined using the modified Christensen method (12). Briefly, bacterial overnight cultures were diluted 1:100 and grown in Tryptic Soy Broth (Bacto Tryptic Soy Broth, art. nr. 211825, BD, USA) with 1% glucose overnight in 96-well plates (Nunc Microwell, Thermo Scientific, Denmark) before staining with 0.1% crystal violet solution (Sigma Aldrich, Germany). After staining of the biofilm, the crystal violet was dissolved using 70% ethanol, and the optical density (OD) was read at 570 nm using ClariostarPlus (BMG Labtech, Germany). Three biological replicates using eight wells for each replicate were performed for all isolates. Biofilm formation of six urine catheter isolates and six skin and soft tissue isolates was tested using 60% artificial urine. The artificial urine had a pH of  $6 \pm 0.08$  and was made according to Sarigul et al. (13). The strong biofilm producer *Staphylococcus epidermidis* RP62A and the poor biofilm producer *S. haemolyticus* 51-03 were used as positive and negative controls, respectively. After the removal of the highest and lowest values of eight replicates/parallels, the average OD values of three replicates were used.

# Phenotypic description

For phenotypic analyses of pigmentation and hemolysis, the isolates were streaked on horse blood agar plates (Thermofisher, Oxoid, UK) and on chocolate agar plates (Thermofisher Oxoid, UK) before incubation at 37°C for 12–18 h. Beta hemolysis was observed as a clear zone around the colonies.

#### Statistical analysis

Data were analyzed using SPSS Statistics 29. Continuous variables were analyzed with a non-parametric test and proportions with a  $\chi$  test. P-values less than 0.05 were considered significant.

## **RESULTS**

## Establishment of the S. borealis strain collection

Using the updated MALDI-TOF database for re-classification, we found 12 (3.9%) *S. borealis* isolates among 308 *S. haemolyticus* blood culture isolates that were re-tested in the retrospective part collected between 2014 and 2021. Seven of these 12 isolates were included in the study. The remaining 122 isolates in this study, from different body sites/sources, were collected in the prospective part of the study during 2021–2022.

## Clinical data

Table 1 shows baseline epidemiological and clinical data, including source of specimen and suspected type of infection, presented separately for hospitalized and non-hospitalized patients. The median (IQR) age of all patients was 62 (51-78) years, and 97/129 (75%) were ≥50 years of age. Overall, 85/129 (66%) isolates were from male patients. Most of the isolates, 81/129 (63%), were from patients receiving outpatient care. The majority of the *S. borealis* isolates were isolated from urine cultures 81/129 (62.8%), followed by isolates from skin and soft tissue specimens (35/129, 27.1%), blood cultures (8/129, 6.2%),

**TABLE 1** Source of specimen, infection type, and underlying conditions of patients with a positive identification of *S. borealis* 

	All patients	Non-hospitalized	Hospitalized
	N = 129	N = 81	N = 48
Age, median (IQR), years	62 (51–78)	59 (40–75)	73 (58–81)
Male	85 (65.9%)	48 (59.3%)	37 (77%)
Specimen			
Urine	81 (62.8%)	57 (70.4%)	24 (50%)
Skin and soft tissue	35 (27.1%)	22 (27.2%)	13 (27.1%)
Blood culture	8 (6.2%)	0	8 (16.7%)
Other <sup>a</sup>	5 (3.9%)	2 (2.5%)	3 (6.3%)
Suggested type/focus infection			
Urinary tract infections	50 (38.9%)	36 (44.4%)	14 (29.1%)
Skin and soft tissue infections	15 (11.6%)	13 (19.8%)	2 (4.2%)
Sepsis	7 (5.4%)	0	7 (14.5%)
Implant-associated infection <sup>b</sup>	2 (1.6%)	0	2 (4.2%)
Airway infections	2 (1.6%)	0	2 (4.2%)
Other	11 (8.5%)	7 (8.6%)	4 (8.3%)
No verified infection	43 (33.3%)	24 (29.6%)	19 (39.6%)
Immunocompromised			
Not reported	29 (22.5%)	21 (25.9%)	8 (16.7%)
No	89 (68.9%)	59 (72.8%)	30 (62.5%)
Yes	11 (8.5%)	1 (1.2%)	10 (20.8%)

<sup>&</sup>lt;sup>a</sup>Ear secretion, expectorate, abortion.

and specimens from implant-associated infections 2 (1.6%). All isolates (n = 7) included from the prospective study were blood culture isolates.

In total, 9/35 (25.7%) of samples from suspected skin and soft tissue infections (SSTIs) were from genital wounds. Suspected urinary tract infections (UTIs) and SSTIs were more common in non-hospitalized patients, in contrast to sepsis and implant-associated infections, which only occurred in hospitalized patients. Three of the patients with a UTI were reported to have a urinary catheter, and in total, six patients had some type of indwelling catheter. Of the hospitalized patients, 10/48 (20.8%) were reported to be immunocompromised, and two of the patients reported to be immunocompromised also had growth of *S. borealis* in the blood culture.

# Antimicrobial susceptibility profile

The AST profiles are presented in Table 2 and Table S2. All isolates were susceptible to vancomycin and linezolid. Rates of resistance to nitrofurantoin, tetracycline, and rifampicin were below 10% in both hospitalized and non-hospitalized patients. The highest rates of resistance were seen towards penicillin, erythromycin, and ciprofloxacin. Overall, 27/129 (20.9%) isolates were identified as methicillin-resistant. Inducible resistance to clindamycin was observed in 7/129 (5.4%) isolates. In total, 43/129 (33.3%) of the isolates were classified as MDR, with a higher proportion of MDR isolates from hospitalized (50%) versus non-hospitalized (23.5%) patients. MDR was not associated with any particular type of infection.

## **Biofilm formation**

Biofilm formation in TSB containing 1% glucose was observed for all isolates, apart from one isolate that did not grow in broth. The strong biofilm producer *S. epidermidis* RP62A formed biofilm yielding an average OD value of 1.7, while the poor biofilm producer *S. haemolyticus* 51-03 formed biofilm with an average OD value of 0.8. Based on the OD values of the reference strains, the *S. borealis* strains were classified as weak biofilm

<sup>&</sup>lt;sup>b</sup>Out of in total six reported patients with implants, only two had an implant-associated infection.

External otitis, abdominal pain.

**TABLE 2** Antimicrobial susceptibility, tested by the disk diffusion method, of 129 *S. borealis* isolated from non-hospitalized and hospitalized patients. The table is showing the number of resistant isolates

Antimicrobial agent	Non-hospitalized	Hospitalized	<i>P</i> -value
	N = 81	N = 48	
Fusidic acid	21 (25.9%)	12 (25%)	0.90
Cefoxitin <sup>a</sup>	10 (12.3%)	17 (35.4%)	0.0018 <sup>d</sup>
Penicillin	20 (24.6%)	25 (52.1%)	0.0016
Erythromycin	35 (43.2%)	16 (33.3%)	0.26
Clindamycin <sup>b</sup>	26 (32.1%)	17 (35.4%)	0.69
Trimethoprim-sulfamethoxazole	0	2 (4.2%)	0.28
Gentamicin	4 (4.9%)	6 (12.5%)	0.12
Linezolid	0	0	
Ciprofloxacin	14 (7.3%)	23 (47.9%)	0.0002
Tetracycline	6 (7.4%)	1 (2.1%)	0.19
Rifampicin	3 (3.7%)	3 (6.3%)	0.50
Nitrofurantoin	1 (1.3%)	1 (2.1%)	0.70
Trimethoprim	7 (8.6%)	10 (20.8%)	0.047
Vancomycin	0	0	
MDR	19 (23.5%)	24 (50%)	0.17

<sup>&</sup>lt;sup>a</sup>Cefoxitin used as indicator of methicillin resistance.

producers (OD 0.1–0.7), medium biofilm producers (OD 0.8–1.7), and strong biofilm producers (OD >1.7). Of the 128 tested *S. borealis* isolates, 59 (46.1%) were weak biofilm producers, 40 (31.3%) were medium biofilm producers, and 29 (22.3%) were strong biofilm producers under the experimental conditions used (Fig. S1). When tested under conditions that included artificial urine, biofilm formation was less pronounced for all 12 isolates, including the positive control, compared with results from the modified Christensen method. Only one urine culture isolate remained a strong biofilm producer when tested in artificial urine.

## Phenotypic description

Phenotypically, *S. borealis* isolates displayed a grayish white to yellow pigmentation on blood agar plates, with a more pronounced yellow pigmentation on chocolate agar media. Of the 129 isolates, 98 (76%) displayed yellow colonies on chocolate agar, while 31 (24%) displayed white colonies on chocolate agar (Fig. 1). There was no correlation between pigmentation and isolation site. On blood agar plates, a clear  $\beta$ -hemolytic zone was seen in all isolates after incubation for 24 h at 37°C. One clinical isolate had a mucoid colony morphology (Fig. 1). This isolate was from a post-operative bacteremia episode following resection of a bladder tumor. The mucoid colony was re-analyzed by MALDI-TOF and confirmed as *S. borealis*.

#### **DISCUSSION**

In this study, *S. borealis* was mainly identified in microbiological specimens from middle-aged to elderly patients, predominantly males. Hospitalized patients were also frequently immunocompromised and/or had different underlying conditions, such as cancer, kidney transplant, or wounds accompanying a suspected *S. borealis* infection. These findings are well in line with the "high-risk population" of patients also seen in other opportunistic CoNS infections (1).

The main clinical finding is that *S. borealis* was most frequently identified in urine cultures and possible UTIs. However, the clinicians interpreting the microbiology results were often not yet aware of the potential clinical relevance of identifying *S. borealis*. As such, they may have considered the results as contamination and not an infection.

<sup>&</sup>lt;sup>b</sup>Inducible resistance to clindamycin was observed in 7/129 (5.4%) of the isolates.

<sup>&</sup>lt;sup>c</sup>MDR: Multidrug resistance, defined as resistance to ≥3 categories of antimicrobial agents.

<sup>&</sup>lt;sup>d</sup>Bold represents statistically significant values.

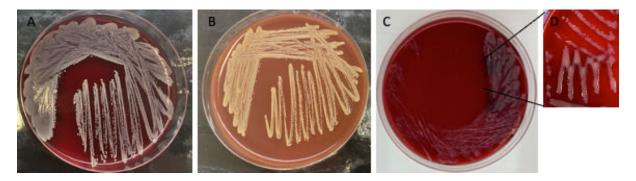


FIG 1 Phenotypic variation of *S. borealis* colonies. (A) *S. borealis* grown on blood agar plates. (B) *S. borealis* colonies on chocolate agar plates. (C and D) The mucoid phenotype observed.

Moreover, information on antibiotic treatment was not given for all possible UTI cases, making a further interpretation of the severity of infection difficult. The positive urine cultures were observed in individuals with serious underlying conditions, such as kidney transplant and kidney failure, but also in individuals with no underlying condition and in pregnant women. *S. saprophyticus* is known to cause uncomplicated UTIs, predominantly in young and middle-aged female patients (14, 15), but other CoNS have usually not been perceived as uropathogens (16). However, in a study from Tanzania, *S. haemolyticus* and *S. epidermidis* were also frequently identified in UTIs (17). *S. saprophyticus* has few virulence factors but produces urease, adhesins to uroepithelial cells, and transport systems involved in osmotolerance (15, 18). *S. borealis* also produces urease, a virulence factor associated with UTIs not only in *S. saprophyticus*, but also in other uropathogenic species, such as *Proteus mirabilis* (19) and *Klebsiella pneumoniae* (20, 21).

In our study, ~30% *S. borealis* isolates were from skin and soft tissue samples. CoNS colonize skin and mucosal surfaces (22). However, they rarely cause SSTIs, except in elderly or immunosuppressed patients, who may present with abscesses and sometimes vulval infections (23). In a recent study from Nigeria, *Staphylococcus cohnii*, *Staphylococcus condimentii*, *Staphylococcus sciuri*, and *Staphylococcus saprophyticus* were identified in 91/265 (34%) of skin and soft tissue samples (24). The high number of *S. borealis* isolates identified from skin and soft tissue samples in our study, and in particular associated with genital wounds, is interesting. However, we lacked detailed data on other bacteria detected during the same infection episode, and we did not have pathology reports that could elucidate on invasiveness. We could therefore not clearly resolve whether *S. borealis* was a contaminant, part of a mixed infection, or the main cause of the SSTI.

MDR was observed in 1/3 of all clinical *S. borealis* isolates. Even higher MDR rates were found in samples from hospitalized patients, in line with data on *S. haemolyticus* (2, 25) and other CoNS species (26). The high AMR rates against erythromycin and fusidic acid were comparable to results when mapping AMR in commensal CoNS isolates in Germany (27). However, when compared with a study mapping MDR (5.2%) in commensal CoNS isolates in Norway, the MDR rates (23.5%) were higher in *S. borealis* isolated from non-hospitalized patients (22). Methicillin resistance was only observed in 21% of *S. borealis* isolates from hospitalized patients compared with 70%–100% in clinical *S. haemolyticus* isolates (28, 29). The presence of MDR in *S. borealis* potentially adds on to the role of CoNS as resistance gene reservoirs for other CoNS and the more pathogenic *S. aureus* (30–32). The AST results should be interpreted with caution, as there is limited data on AST and clinical effects of antimicrobial agents for infections caused by *S. borealis*. Depending on the frequency of identification, clinical significance, and perceived clinical need, breakpoints for *S. borealis* may be established by EUCAST in the future (33).

All *S. borealis* isolates formed biofilm *in vitro* using a standard biofilm assay, and around 20% formed more biofilm than the strong biofilm producer *S. epidermidis* RP62a. Biofilm formation, primarily on medical implants, is one of the hallmarks of "difficult

to treat" CoNS infections (1, 34), but we identified only two patients with a suspected *S. borealis*-associated medical implant infection and three patients with urinary tract catheters. However, biofilm formation is also a well-known virulence factor for many uropathogens and considered an important mechanism behind high rates of recurrent UTIs in some patients (35). Further testing of *S. borealis* biofilm formation using artificial urine did not reveal an improved biofilm forming ability; however, conditions could have been further optimized to also include catheter material.

This study has several strengths. It is the largest collection to date of S. borealis isolates presented with clinical data. The broad AST and biofilm testing were performed with established methods and give updated information on this newly described species. There are also several limitations. We used clinical data from the medical information accompanying the specimen sent to the clinical microbiology laboratories. However, these data often lacked the granularity required to clearly establish a causal relationship between the culture result and a clinical infection. Moreover, the study was designed as an observational study, not designed to determine causality. Additionally, since S. borealis was not a known pathogen during the study, the potential clinical relevance of identifying S. borealis was unclear to clinicians. This hampers the interpretation of some of our clinical results. However, our study indicates that S. borealis is a novel CoNS uropathogen. The clinical relevance of S. borealis in other types of infections needs to be further explored in large clinical studies with more detailed clinical descriptions. However, it seems to have a role in SSTIs and potentially invasive bacteremia in immunocompromised patients (36, 37). Ethical approval for our study included an opt-out approach, and we decided only to contact adults who were able to respond in person. Thus, lack of information on the relevance of S. borealis in children is also a limitation.

#### Conclusion

In this first study describing the potential clinical relevance of *S. borealis*, we have described that this new staphylococcal species was mainly found in clinical samples from elderly males in specimens obtained due to suspected UTIs and SSTIs. The level of MDR in *S. borealis* is comparable to other CoNS, but resistance toward methicillin and penicillin is notably lower than what is observed in clinical *S. haemolyticus* isolates.

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Claus Klingenberg, Conceptualization, Data curation, Formal analysis, Methodology, Writing – review and editing | Hermoine Jean Venter, Methodology, Writing – review and editing | Jan Egil Afset, Methodology, Writing – review and editing | Olaf Stromme, Methodology, Writing – review and editing | Therese Johansen, Methodology, Writing – review and editing | Kyriakos Zaragkoulias, Data curation, Validation, Writing – review and editing | Ståle Tofteland, Methodology, Writing – review and editing | Pia Littauer, Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Project administration, Validation, Writing – original draft.

## **ETHICS APPROVAL**

All patients from whom an *S. borealis* isolate was identified in a microbiological sample received written information about the study and the possibility to opt-out of allowing researchers to collect clinical data. If the patients opted not to participate in the study, they had to notify the principal investigator within four weeks after receiving this letter. The study was approved by the Regional Committee for Medical and Health Research Ethics (REK-Nord 2021/216491).

#### **ADDITIONAL FILES**

The following material is available online.

## Supplemental Material

Fig. S1 (Spectrum01988-24-S0001.tiff). Biofilm formation of *S. borealis* isolates.

**Supplemental material (Spectrum01988-24-S0002.docx).** Supplemental figure and table legends.

Table S1 (Spectrum01988-24-S0003.docx). Clinical questions asked.

**Table S2 (Spectrum01988-24-S0004.xlsx).** Results from antimicrobial susceptibility testing by disc diffusion.

#### **REFERENCES**

- Becker K, Heilmann C, Peters G. 2014. Coagulase-negative staphylococci. Clin Microbiol Rev 27:870–926. https://doi.org/10.1128/CMR.00109-13
- Pain M, Wolden R, Jaén-Luchoro D, Salvà-Serra F, Iglesias BP, Karlsson R, Klingenberg C, Cavanagh JP. 2020. Staphylococcus borealis sp. nov., isolated from human skin and blood. Int J Syst Evol Microbiol 70:6067–6078. https://doi.org/10.1099/ijsem.0.004499
- Król J, Wanecka A, Twardoń J, Florek M, Marynowska M, Banaszkiewicz S, Kaczmarek-Pieńczewska A, Pląskowska E, Brodala M, Chwirot W,
- Korzeniowska-Kowal A, De Buck J. 2023. *Staphylococcus borealis* a newly identified pathogen of bovine mammary glands. Vet Microbiol 286:109876. https://doi.org/10.1016/j.vetmic.2023.109876
- Freu G, Gioia G, Gross B, Biscarini F, Virkler P, Watters R, Addis MF, Franklin-Guild RJ, Runyan J, Masroure AJ, Bronzo V, Dos Santos MV,

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Moroni P. 2024. Frequency of non-aureus staphylococci and mammalii-cocci species isolated from quarter clinical mastitis: a 6-year retrospective study. J Dairy Sci 107:3813–3823. https://doi.org/10.3168/jds.2023-24086

- Abdullahi IN, Lozano C, Simón C, Zarazaga M, Torres C. 2023. Within-host diversity of coagulase-negative staphylococci resistome from healthy pigs and pig farmers, with the detection of cfr-carrying strains and MDR-S. borealis. Antibiotics (Basel) 12:10. https://doi.org/10.3390/antibiotics1 2101505
- Ocloo R, Newton-Foot M, Ziebuhr W, Whitelaw AC. 2023. Molecular epidemiology and antibiotic resistance of staphylococci other than Staphylococcus aureus in children in Cape Town, South Africa. Front Microbiol 14:1239666. https://doi.org/10.3389/fmicb.2023.1239666
- Becker K, Both A, Weißelberg S, Heilmann C, Rohde H. 2020. Emergence of coagulase-negative staphylococci. Expert Rev Anti Infect Ther 18:349– 366. https://doi.org/10.1080/14787210.2020.1730813
- Matuschek E, Brown DFJ, Kahlmeter G. 2014. Development of the EUCAST disk diffusion antimicrobial susceptibility testing method and its implementation in routine microbiology laboratories. Clin Microbiol Infect 20:0255–66. https://doi.org/10.1111/1469-0691.12373
- EUCAST. 2024. The European committee on antimicrobial susceptibility testing. breakpoint tables for interpretation of MICs and zone diameters
- EUCAST. 2023. Breakpoint tables for interpretation of MICs and zone diameters v 13. The European Committee on Antimicrobial Susceptibility Testing. Available from: http://www.eucast.org/clinical\_breakpoints
- Skov R, Larsen AR, Kearns A, Holmes M, Teale C, Edwards G, Hill R. 2014. Phenotypic detection of mecC-MRSA: cefoxitin is more reliable than oxacillin. J Antimicrob Chemother 69:133–135. https://doi.org/10.1093/j ac/dkt341
- Christensen GD, Simpson WA, Younger JJ, Baddour LM, Barrett FF, Melton DM, Beachey EH. 1985. Adherence of coagulase-negative staphylococci to plastic tissue culture plates: a quantitative model for the adherence of staphylococci to medical devices. J Clin Microbiol 22:996–1006. https://doi.org/10.1128/jcm.22.6.996-1006.1985
- Sarigul N, Korkmaz F, Kurultak İ. 2019. A new artificial urine protocol to better imitate human urine. Sci Rep 9:20159. https://doi.org/10.1038/s41 598-019-56693-4
- Schleifer KH, Kloos WE. 1975. Isolation and characterization of staphylococci from human skin I. amended descriptions of Staphylococcus epidermidis and Staphylococcus saprophyticus and descriptions of three new species: Staphylococcus cohnii, Staphylococcus haemolyticus, and Staphylococcus xylosus. Int J Syst Bacteriol 25:50–61. https://doi.org/ 10.1099/00207713-25-1-50
- Kuroda M, Yamashita A, Hirakawa H, Kumano M, Morikawa K, Higashide M, Maruyama A, Inose Y, Matoba K, Toh H, Kuhara S, Hattori M, Ohta T. 2005. Whole genome sequence of *Staphylococcus saprophyticus* reveals the pathogenesis of uncomplicated urinary tract infection. Proc Natl Acad Sci U S A 102:13272–13277. https://doi.org/10.1073/pnas.0502950 102
- Megged O. 2022. Coagulase-negative staphylococci: a rare cause of urinary tract infections in children with consequences on clinical practice. Eur J Pediatr 181:1099–1104. https://doi.org/10.1007/s00431-0 21-04308-4
- Phillip S, Mushi MF, Decano AG, Seni J, Mmbaga BT, Kumburu H, Konje ET, Mwanga JR, Kidenya BR, Msemwa B, Gillespie S, Maldonado-Barragan A, Sandeman A, Sabiti W, Holden MTG, Mshana SE. 2023. Molecular characterizations of the coagulase-negative staphylococci species causing urinary tract infection in tanzania: a laboratory-based crosssectional study. Pathogens 12:180. https://doi.org/10.3390/pathogens12 020180
- Heilbronner S, Foster TJ. 2021. Staphylococcus lugdunensis: a skin commensal with invasive pathogenic potential. Clin Microbiol Rev 34:e00205-20. https://doi.org/10.1128/CMR.00205-20
- Gatermann S, John J, Marre R. 1989. Staphylococcus saprophyticus urease: characterization and contribution to uropathogenicity in unobstructed urinary tract infection of rats. Infect Immun 57:110–116. https://doi.org/ 10.1128/iai.57.1.110-116.1989

- Jones BD, Lockatell CV, Johnson DE, Warren JW, Mobley HL. 1990. Construction of a urease-negative mutant of *Proteus mirabilis*: analysis of virulence in a mouse model of ascending urinary tract infection. Infect Immun 58:1120–1123. https://doi.org/10.1128/iai.58.4.1120-1123.1990
- Podschun R, Sievers D, Fischer A, Ullmann U. 1993. Serotypes, hemagglutinins, siderophore synthesis, and serum resistance of *Klebsiella* isolates causing human urinary tract infections. J Infect Dis 168:1415–1421. https://doi.org/10.1093/infdis/168.6.1415
- Cavanagh JP, Wolden R, Heise P, Esaiassen E, Klingenberg C, Aarag Fredheim EG. 2016. Antimicrobial susceptibility and body site distribution of community isolates of coagulase-negative staphylococci. APMIS 124:973–978. https://doi.org/10.1111/apm.12591
- Natsis NE, Cohen PR. 2018. Coagulase-negative Staphylococcus skin and soft tissue infections. Am J Clin Dermatol 19:671–677. https://doi.org/10. 1007/s40257-018-0362-9
- Akinduti PA, Obafemi YD, Ugboko H, El-Ashker M, Akinnola O, Agunsoye CJ, Oladotun A, Phiri BSJ, Oranusi SU. 2022. Emerging vancomycin-non susceptible coagulase negative staphylococci associated with skin and soft tissue infections. Ann Clin Microbiol Antimicrob 21:31. https://doi.or g/10.1186/s12941-022-00516-4
- Pain M, Hjerde E, Klingenberg C, Cavanagh JP. 2019. Comparative genomic analysis of *Staphylococcus haemolyticus* reveals key to hospital adaptation and pathogenicity. Front Microbiol 10:2096. https://doi.org/1 0.3389/fmicb.2019.02096
- Felgate H, Crossman LC, Gray E, Clifford R, Correia A, Dean R, Wain J, Langridge GC. 2023. Known mechanisms cannot account for a third of reduced susceptibility in non-aureus staphylococci. NPJ Antimicrob Resist 1:15. https://doi.org/10.1038/s44259-023-00008-1
- Marincola G, Liong O, Schoen C, Abouelfetouh A, Hamdy A, Wencker FDR, Marciniak T, Becker K, Köck R, Ziebuhr W. 2021. Antimicrobial resistance profiles of coagulase-negative staphylococci in communitybased healthy individuals in germany. Front Public Health 9:684456. http s://doi.org/10.3389/fpubh.2021.684456
- Cavanagh JP, Hjerde E, Holden MTG, Kahlke T, Klingenberg C, Flægstad T, Parkhill J, Bentley SD, Sollid JUE. 2014. Whole-genome sequencing reveals clonal expansion of multiresistant Staphylococcus haemolyticus in European hospitals. J Antimicrob Chemother 69:2920–2927. https://doi. org/10.1093/jac/dku271
- Rossi CC, Ahmad F, Giambiagi-deMarval M. 2024. Staphylococcus haemolyticus: an updated review on nosocomial infections, antimicrobial resistance, virulence, genetic traits, and strategies for combating this emerging opportunistic pathogen. Microbiol Res 282:127652. https://do i.org/10.1016/j.micres.2024.127652
- John J, George S, Nori SRC, Nelson-Sathi S. 2019. Phylogenomic analysis reveals the evolutionary route of resistant genes in *Staphylococcus* aureus. Genome Biol Evol 11:2917–2926. https://doi.org/10.1093/gbe/ev z213
- Fišarová L, Pantůček R, Botka T, Doškař J. 2019. Variability of resistance plasmids in coagulase-negative staphylococci and their importance as a reservoir of antimicrobial resistance. Res Microbiol 170:105–111. https:// doi.org/10.1016/j.resmic.2018.11.004
- Smith JT, Andam CP. 2021. Extensive horizontal gene transfer within and between species of coagulase-negative Staphylococcus. Genome Biol Evol 13:evab206. https://doi.org/10.1093/gbe/evab206
- 33. EUCAST. 2024. EUCAST guidance on When there are no breakpoints in breakpoint tables
- Otto M. 2018. Staphylococcal biofilms. Microbiol Spectr 6. https://doi.org/10.1128/microbiolspec.gpp3-0023-2018
- Lila ASA, Rajab AAH, Abdallah MH, Rizvi SMD, Moin A, Khafagy E-S, Tabrez S, Hegazy WAH. 2023. Biofilm lifestyle in recurrent urinary tract infections. Life (Basel) 13:148. https://doi.org/10.3390/life13010148
- Argemi X, Hansmann Y, Prola K, Prévost G. 2019. Coagulase-negative staphylococci pathogenomics. Int J Mol Sci 20:1215. https://doi.org/10.3 390/iims20051215
- von Eiff C, Peters G, Heilmann C. 2002. Pathogenesis of infections due to coagulase-negative staphylococci. Lancet Infect Dis 2:677–685. https://d oi.org/10.1016/s1473-3099(02)00438-3