

Vancomycin Heteroresistance in Bloodstream Isolates of *Staphylococcus capitis*[∇]

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Nine *Staphylococcus capitis* isolates from blood cultures of newborns were examined for resistance to vancomycin. MICs were within the susceptible range, but population profiling revealed a resistant subpopulation. Only isolates with the largest subpopulation were identified as heteroresistant to vancomycin by Etest. This finding may have therapeutic implications.

Recent reports indicate the possible emergence of *Staphylococcus capitis* as a significant pathogen causing late-onset sepsis in very-low-birth-weight (VLBW) infants (<1,500 g) (6, 16, 18, 20). Reduced susceptibility to vancomycin has been reported in several species of coagulase-negative staphylococci (1, 2, 4, 9, 10, 17); however, there is very little information on the levels of such resistance in *S. capitis*. Reduced susceptibility to vancomycin occurring in methicillin-resistant staphylococci translates into limited treatment options, particularly in newborn infants.

Heteroresistant *S. capitis* isolates may escape detection because MICs of vancomycin of ≤ 4 $\mu\text{g/ml}$ are generally interpreted as susceptible (5). Agar-based screening tests for detecting heteroresistance in staphylococci are simple to perform (14, 15, 19); however, the sensitivity and specificity of Etest strips are superior (22). Population analysis profiling (PAP) is the most reliable method for detecting heteroresistance, but it is time-consuming and fails to provide results in a clinically relevant time frame (19). Data on the prevalence and level of vancomycin resistance are essential for assessing clinical relevance and treatment options for infants infected with *S. capitis*. This study examines a collection of nine *S. capitis* isolates obtained from blood cultures of VLBW infants in the Neonatal Intensive Care Unit (NICU) at the Royal Women's Hospital, Melbourne, Australia between 1998 and 2002 (3). Pulsed-field gel electrophoresis combined with Southern blot analysis and probing with insertion element IS256 showed they were closely related yet unique, except for two isolates from the same infant (3). The reference strains used were *S. capitis* ATCC 27840 and *Staphylococcus aureus* Mu3 (ATCC 700698) and Mu50 (ATCC 700699) (11–13).

Isolates were screened for vancomycin heteroresistance on brain heart infusion agar (BHIA) (Oxoid Ltd., Hampshire, England) containing 4 $\mu\text{g/ml}$ of vancomycin (VAN 4) (Sigma-

Aldrich Pty. Ltd., Sydney, Australia) (12). MICs were determined by conventional methods and by vancomycin and teicoplanin Etest strips (AB Biodisk, Solna, Sweden) (8), taking care to include small colonies within the clear zone. The PAP profiles were interpreted by calculating area under the concentration-time curve for test and Mu3 ($\text{AUC}_{\text{test}}/\text{AUC}_{\text{Mu3}}$) ratios with the aid of GraphPad Prism 5 software (San Diego, CA) (19, 21). Since the AUC is affected by the size of the initial inoculum, CFUs were standardized to match the initial inoculum of Mu3 for each replicate. All tests were performed on at least three separate occasions.

The various methods differed in their ability to detect resistant subpopulations of *S. capitis*: broth microdilution (CLSI) was the least sensitive method, detecting only one resistant isolate, and the Etest detected three resistant isolates, while VAN 4 screening and PAP analysis detected resistant subpopulations in all isolates (Table 1). Three isolates produced variable results on the VAN 4 screening plates, indicating instability of their heteroresistant phenotype. Visual examination of the PAP graphs showed heterogeneous resistance with strain-dependent differences in the sizes of the resistant subpopulation (Fig. 1). For the three most resistant strains (Mu50 and isolates 15 and 22), there was complete agreement between the results of PAP-AUC analysis, VAN 4 screens, and the Etest. Isolate 6 had a very high PAP-AUC value, produced variable screening results, and was interpreted as nonheteroresistant by the Etest. The discrepant PAP-AUC value could be explained by the unusual shape of the PAP graph, reflecting high colony counts on VAN 2 followed by a sharp drop on VAN 3 plates (Fig. 1). All other isolates and reference strains showed PAP-AUC values close to those of Mu3, but were not heteroresistant according to the Etest. Rankings of isolates according to the size of the resistant subpopulation were generally similar by PAP-AUC analysis and MICs generated by Etests.

These results suggest that a vancomycin-heteroresistant subpopulation is present in all isolates of *S. capitis*. They confirm the unreliability of conventional MICs except for the most resistant isolates and show that the Etest and the VAN 4 screening tests detect only the most resistant isolates but fail to

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TABLE 1. Vancomycin resistance of *S. capitis* isolates

Isolate or strain no.	Result by ^a :								
	PAP-AUC ^b		Vancomycin screen (4 µg/ml) ^c		CLSI standard		Etest		Interpretation
	Mean (SE) AUC _{test/Mu3} ratio	Interpretation	No. of colonies	Interpretation	MIC µg/ml	Interpretation	Vancomycin	Teicoplanin	
<i>S. capitis</i> 6	1.7 (0.15)	I	7, 86, CG	V	2	S	6	2	S
<i>S. capitis</i> 22	1.7 (0.12)	I	CG	R	4	S	12	16	H
<i>S. capitis</i> 15	1.3 (0.08)	I	CG	R	4	S	12	16	H
<i>S. aureus</i> Mu50	1.6 (0.05)	I	CG	R	8	R	16	12	H
<i>S. capitis</i> 17	1.2 (0.04)	H	4	H	2	S	3	4	S
<i>S. capitis</i> 8a ^d	1.1 (0.03)	H	2	H	1	S	4	3	S
<i>S. capitis</i> 9	1.1 (0.03)	H	0, 2, CG	V	1	S	4	2	S
<i>S. capitis</i> 16	1.1 (0.03)	H	1	H	1	S	3	2	S
<i>S. capitis</i> 8b ^d	1.0 (0.04)	H	12	H	1	S	8	3	S
<i>S. capitis</i> 18	1.0 (0.05)	H	0, 36, CG	V	2	S	4	4	S
<i>S. capitis</i> ATCC 27840	1.0 (0.01)	H	1	H	1	S	2	0.2	S
<i>S. aureus</i> Mu3	1	H	14	H	2		8	16	H

^a I, intermediate; S, susceptible; H, possibly heteroresistant; R, potentially resistant; V, variable results; CG, confluent growth on at least three of four replicates.
^b Mean (standard error) AUC determined by PAP. An AUC_{test/Mu3} ratio of ≤0.90 was interpreted as susceptible, a ratio of 0.90 to 1.3 was interpreted as heteroresistant, and a ratio of ≥1.3 was interpreted as intermediate.
^c Number of colonies on BHIA containing 4 µg/ml vancomycin after 48 h.
^d Isolates 8a and 8b were collected on different occasions from the same infant.

detect isolates with smaller resistant subpopulations, which could be enriched during vancomycin therapy.

This report is, to the best of our knowledge, only the second to describe a cluster of vancomycin-heteroresistant *S. capitis* strains among VLBW infants in an NICU. Van Der Zwet et al. (18) demonstrated variable proportions of resistant subpopulations in *S. capitis* isolates with closely related or identical genetic profiles. These data suggest that heteroresistant *S. capitis* strains, which are undetectable by standard MIC measurement, could be emerging pathogens in NICUs.

The origin of these nine closely related vancomycin-heteroresistant isolates, present in the NICU for at least 5 years, is enigmatic. It is possible that frequent vancomycin use in the

unit led to enrichment of resistant cells present in a common ancestor, resulting in subpopulations of variable size. Although the outcome for *S. capitis* septicemia in VLBW infants is generally good in our unit, it is of concern that further increases in vancomycin resistance could occur. Our observation of a resistant subpopulation in all *S. capitis* strains examined, including ATCC 27840, which was deposited in 1975, suggests that heteroresistance to vancomycin could be an intrinsic property of *S. capitis*. Although more isolates of *S. capitis* need to be examined, given that the relationship between vancomycin heteroresistance and treatment failure is still uncertain (7), it would be wise to consider all isolates as potentially resistant and to monitor clinical responses to vancomycin very carefully,

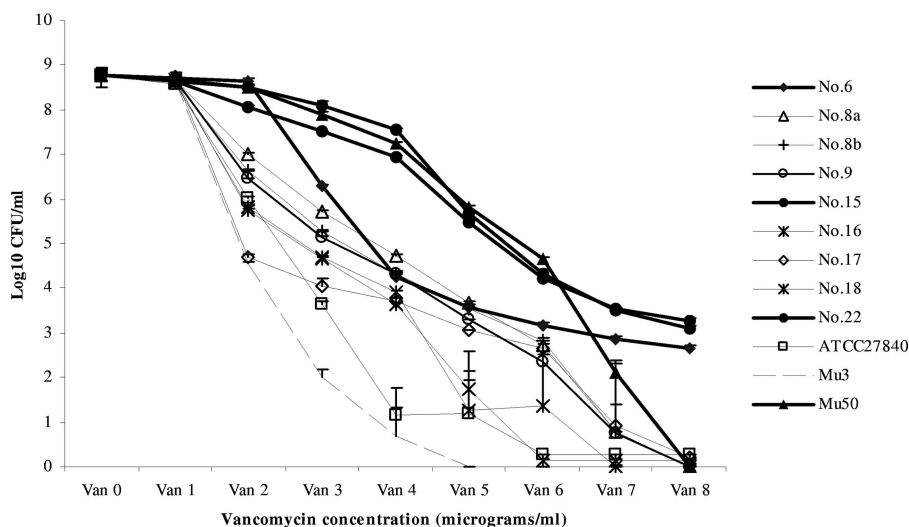


FIG. 1. Population analysis profiling of *S. capitis* strains isolated from blood cultures of infants. Heavy lines, intermediate. The means of three separate investigations and standard errors are presented.

particularly with more deep-seated infections such as osteomyelitis or infections where antibiotic penetration is an issue, such as endocarditis and meningitis. There is an urgent need for more data on the clinical relevance of vancomycin heteroresistance in staphylococci, in particular *S. capitis*, and for the development of reliable, rapid, and inexpensive methods to detect such resistance (7).

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