Sequential Emergence of Antibiotic Resistance in Enterococcal Bloodstream Isolates over 25 Years

David R. Murdoch,^{1*} Stanley Mirrett,¹ Lizzie J. Harrell,¹ Janet S. Monahan,² and L. Barth $Reller¹$

*Clinical Microbiology Laboratories, Duke University Medical Center, Durham, North Carolina 27710,*¹ *and University of Colorado Hospital, Denver, Colorado*²

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We determined the antibiotic susceptibilities of 1,785 enterococcal bloodstream isolates collected over 25 years. Antibiotic resistance emerged at a greater rate in *Enterococcus faecium* **than in other enterococcal species, and** *E. faecium* **isolates became proportionally more common over time. Our findings confirm the pattern of emerging antibiotic resistance among enterococci and highlight the increasing importance of** *E. faecium* **as a cause of bloodstream infection.**

Large surveillance programs have provided considerable data about the antimicrobial susceptibilities of enterococci over the past decade (4, 5). However, few studies have reviewed enterococcal isolates collected prior to 1990 by using current criteria for species identification and susceptibility testing. The principal objective of this study was to document the sequential emergence of antimicrobial resistance among enterococci isolated over the past 25 years, as important acquired resistance mechanisms have emerged. Therefore, we tested enterococcal bloodstream isolates collected from 1975 to 1999 by using current antimicrobial susceptibility testing methodology and interpretative criteria.

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Sequential isolates were collected at the University of Colorado Hospital, Denver, Colo., from 1975 to 1988 and at Duke University Medical Center, Durham, N.C., from 1989 to 1999. Only one isolate per patient for each bacteremic episode was included (i.e., there was >7 days between episodes unless different species were isolated), and all had been stored at -70° C. Isolates from before 1993 were tested retrospectively, while most other isolates were tested immediately after initial isolation. An additional 100 enterococcal bloodstream isolates collected at the University of Colorado Hospital during the year 2000 by using the same clinical criteria were also tested to determine the present antimicrobial susceptibilities and species profiles of enterococci at this site.

Enterococci were initially identified by their appearance when stained with Gram stain and by their colonial morphology, plus their ability to grow in 6.5% NaCl and to hydrolyze esculin in the presence of 40% bile salts and/or their ability to hydrolyze pyrrolidonyl- β -naphthylamide. Species-level identification was made with the MicroScan Pos Combo Panel type 10 or 13 (Dade Behring, Inc., West Sacramento, Calif.) for all isolates collected before 1993 and the API 20 Strep identification system (bioMérieux Vitek, Inc., Hazelwood, Mo.) for isolates collected since 1993. These methods were supplemented by conventional test procedures (1).

All susceptibility testing was performed according to the methods and interpretative criteria recommended by the National Committee for Clinical Laboratory Standards (8). For isolates collected before 1993, the broth microdilution method (MicroScan Pos Combo Panel type 10 or 13) was used to determine susceptibilities to ampicillin, vancomycin, gentamicin, and streptomycin. For the remaining isolates, the disk diffusion method was used to determine susceptibilities to ampicillin and vancomycin and an agar dilution method (Synergy Quad; Remel, Lenexa, Kans.) was used to detect high-level gentamicin and streptomycin resistance and vancomycin resistance. All isolates were tested for β -lactamase production by using nitrocefin disks (Cefinase; Becton Dickinson Diagnostic Systems, Sparks, Md.). Ampicillin-resistant isolates were further tested by the broth microdilution method to determine precise MICs. Custom-made frozen reference microdilution panels were used for this purpose (Dade Behring, Inc.).

A total of 1,785 enterococcal bloodstream isolates collected from 1975 to 1999 were tested. There were 1,311 *Enterococcus faecalis*, 410 *E. faecium*, 17 *E. casseliflavus*, 12 *E. avium*, 10 *E. gallinarum*, 6 *E. durans*, and 4 *E. raffinosus* isolates. The other 15 isolates could not be identified to the species level, could not be found, or could not be recovered for testing. Overall, *E. faecium* isolates became proportionally more common after 1979. The ratios of *E. faecalis* to *E. faecium* isolates were 3:1 from 1975 to 1979, 7:1 from 1980 to 1984, 7:1 from 1985 to 1989, 4:1 from 1990 to 1994, and 3:1 from 1995 to 1999. For 1999 alone, the ratio was 2.2:1.

The antimicrobial susceptibilities of the isolates are shown in Table 1. Ampicillin resistance was first detected in *E. faecium* in 1989, with resistance rates steadily increasing to 80% of *E. faecium* isolates during 1999. Over this time period there was a gradual temporal rise in MICs for ampicillin-resistant *E. faecium* isolates, although the median MIC for these isolates remained at 64 μ g/ml for most years (Fig. 1). β -Lactamase production was detected in only five *E. faecalis* isolates, all during an 11-month period (February 1990 to January 1991);

^{*} Corresponding author. Present address: Microbiology Unit, Canterbury Health Laboratories, P.O. Box 151, Christchurch, New Zealand. Phone: 64 3 364 1530. Fax: 64 3 364 0238. E-mail: david. murdoch@cdhb.govt.nz.

Yr	E. faecalis isolates					E. faecium isolates					All isolates				
	No.	No. $(\%)$ resistant to:				No.	No. $(\%)$ resistant to:				No.	No. $(\%)$ resistant to:			
		AMP	VAN	GEN	STR		AMP	VAN	GEN	STR		AMP	VAN	GEN	STR
1975-1979	57	θ	0		16(28)	20				9(45)	83				28(34)
1980-1984	86	θ	0		24(28)	12				(8)	99				25(25)
1985-1989	83	Ω	θ	9(11)	28(34)	12	(8)		1(8)	(8)	98	2(2)		9(9)	30(31)
1990-1994	448		(0.2)	101(23)	135 (30)	112	63 (56)	18(16)	37(33)	54 (48)	594	70(12)	19(3)	(24) 141	195 (33)
1995-1999	637	Ш	16(3)	161(25)	192(30)	254	(74) 189	67 (26)	72(28)	(47) 20	911	194 (21)	83(9)	234(26)	312(34)
2000^b	75	θ	(1,	13 (17)	23(31)	22	14 (64)	8(36)	(32)	11(50)	100	14 (14)	9 (9)	20(20)	34(34)

TABLE 1. Patterns of acquired resistance in enterococcal bloodstream isolates to ampicillin, vancomycin, gentamicin, and streptomycin*^a*

^a AMP, ampicillin; VAN, vancomycin; GEN, high-level gentamicin; STR, high-level streptomycin.

^b Colorado isolates only.

at the time, these isolates were found to be part of the large "mid Atlantic" clone (7). The first *E. faecalis* isolate with ampicillin resistance not due to β -lactamase production was detected in 1997. Vancomycin resistance was first detected in *E. faecium* in 1992 and in *E. faecalis* in 1993. High-level gentamicin resistance first appeared in *E. faecalis* during 1985 and in *E. faecium* in 1989. Of the 64 enterococcal isolates of species other than *E. faecalis* and *E. faecium*, 4 were resistant to ampicillin (3 *E. raffinosus* isolates and 1 isolate not identified to the species level), 4 had high-level gentamicin resistance (none identified to the species level), and 11 had high-level streptomycin resistance (5 isolates not identified to the species level, 3 *E. avium* isolates, 2 *E. gallinarum* isolates, and 1 *E. casseliflavus* isolate). Reduced susceptibility to vancomycin was detected only in *E. casseliflavus* and *E. gallinarum* and was presumed to represent intrinsic vancomycin resistance.

The first multidrug-resistant isolate (i.e., with significant resistance to two or more antibiotics) was an *E. faecalis* strain from 1985 that exhibited high-level resistance to both gentamicin and streptomycin. In 1999, 59 (35%) of 168 isolates were multidrug resistant. Of these, 33 were identified as *E. faecium* isolates, 30 were resistant to vancomycin, and 8 (all of which were *E. faecium* isolates) were resistant to all four antibiotics tested.

The 100 Colorado isolates collected in 2000 comprised 75 *E. faecalis* isolates, 22 *E. faecium* isolates, and 1 isolate each of *E. casseliflavus*, *E. gallinarum*, and *E. durans*. The antimicrobial susceptibility patterns of these isolates were similar to those of the most recent isolates from North Carolina (Table 1), indicating similar courses of emergence of antimicrobial resistance in the two centers.

Overall, the sequential appearance of antibiotic resistance in enterococci and the resistance rates determined by our study are similar to the findings from other United States studies during the same time period (4, 5,10, 11). In general, antibiotic resistance has emerged at a greater rate in *E. faecium* than in other enterococcal species. For the *E. faecium* isolates that were resistant to ampicillin, we documented a gradual increase

FIG. 1. MICs for ampicillin-resistant *E. faecium* isolates by year of isolation.

in MICs over time, although this was not as marked in our study as in those reported by other investigators (3a).

As well as becoming more resistant, *E. faecium* isolates became proportionally more common during the study period. Data from other studies support this trend. In United States studies before 1990, the ratio of *E. faecalis* to *E. faecium* was 10:1 to 13:1 for both bloodstream isolates and isolates from other sites (3, 6, 11). Thereafter, the ratio fell to 5:1 in 1992 (4) and to 3:1 from 1997 to 1999 (5). The increasing predominance of *E. faecium* as a cause of bloodstream and other infections may partly reflect the high rate of antimicrobial resistance in this species and the fact that antibiotic pressure in a hospital environment allows for its selection. The emergence of *E. faecium* as a cause of bacteremia is of particular concern given that this species is associated with a higher mortality than *E. faecalis* is (9), independent of the presence or absence of vancomycin resistance (2). Moreover, the large numbers of strains that are resistant to both ampicillin and vancomycin leave few therapeutic options and none with documented synergy (10).

A major limitation of the present study is the fact that the isolates were collected sequentially, rather than concurrently, from the two different medical centers. However, the distribution of species and antimicrobial susceptibility patterns of the enterococcal bloodstream isolates from the University of Colorado Hospital during 2000 were similar to those of recent isolates from Duke University Medical Center, suggesting that resistance has emerged in a similar fashion in both centers.

The results of this study confirm the pattern of emerging antibiotic resistance among enterococci over recent years and highlight the increasing importance of *E. faecium* as a cause of bloodstream infection.

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