Emerging Incidence of *Enterococcus faecium* among Hospital Isolates (1993 to 2002)

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Historically, most clinical microbiology laboratories report that 80 to 90% of enterococci are *Enterococcus faecalis*, whereas *E. faecium* accounts for 5 to 10% of isolates. At our medical center from 1993 to 2002, we evaluated the percentages of *E. faecium* among all enterococcal isolates and the percentages of *E. faecium* isolates that were vancomycin resistant. Over this 10-year period, the percentage of enterococci that were identified as *E. faecium* increased from 12.7 to 22.2% (*P* < 0.001) and the proportion of *E. faecium* that was vancomycin resistant increased from 28.9 to 72.4% (*P* < 0.001). Both the percentage of *E. faecium* among the enterococci and the proportion of vancomycin-resistant *E. faecium* increased significantly over this 10-year period.

Enterococci are normal inhabitants of the gastrointestinal tract, but they may be associated with invasive disease (7). Common sites of clinical infection include the urinary tract, the bloodstream, and intra-abdominal or pelvic wounds. Previously, due to the low pathogenicity of enterococci, some investigators advocated not treating this organism, but targeted antimicrobial therapy is now common (3). However, with the emergence of antimicrobial resistance among enterococci, treatment options present ongoing challenges.

*Enterococcus faecalis* and *Enterococcus faecium* are the predominant enterococcal species identified in clinical microbiology laboratories. Historically, these laboratories report that 80 to 90% of enterococci are *E. faecalis*, whereas *E. faecium* accounts for 5 to 10% of enterococci (2, 6). The enterococci *E. durans*, *E. avium*, *E. raffinosus*, *E. gallinarum*, and *E. casseliflavus* are reported much less frequently than *E. faecalis* and *E. faecium* (6). More recently, the SENTRY Antimicrobial Resistance Surveillance Program reported that in 2001, *E. faecium* accounted for 20% of clinical enterococcal infections in the United States (5). This finding is of potential concern, as *E. faecium* is more commonly associated with vancomycin resistance than are the other enterococci (14).

*E. faecium* strains expressing high levels of ampicillin resistance (MIC > 128 [micrograms]/ml) emerged in medical centers in the United States in the late 1980s (14). Subsequently, vancomycin-resistant enterococci (VRE) were reported in France in 1985 and in the United States in 1989 (6). Since that time, the incidence of VRE has continued to increase, and VRE make up in excess of 25% of enterococci in some intensive care units (11). Although any of the enterococcal species may be associated with ampicillin or glycopeptide resistance, *E. faecium* more commonly acquires resistance relative to the other enterococcal species (14).

Clinical microbiology laboratories often identify enterococci to the species level for clinical as well as surveillance isolates.

Over the past decade, we have observed a significant increase in the percentage of *E. faecium* isolates, a result which suggests a changing epidemiology of this organism at our medical center. Despite our adherence to standard infection control practices, the percentage of *E. faecium* isolates identified and the concomitant vancomycin resistance continued to increase. Therefore, we reviewed the changing incidence of enterococci at our hospital.

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Data from the clinical microbiology laboratory at Northwestern Memorial Hospital (Chicago, Ill.) were extracted to include all enterococcal isolates found from 1993 to 2002. These data were evaluated by a search of the Sunquest Information Systems (Tucson, Ariz.) database for the following bacterial organisms: *E. faecalis*, *E. faecium*, *E. avium*, *E. durans*, *E. casseliflavus*, and *E. gallinarum*. All enterococci were identified to the species level according to standard methods. Each positive culture result was reviewed to determine whether the isolate was a clinical or surveillance (rectal swab) specimen and was categorized by the year of identification. The data are for one isolate per patient per month (duplicates and surveillance cultures were removed). Vancomycin resistance was determined according to NCCLS standards (10), recorded for each isolate, and categorized by the year of identification. The isolates were categorized and graphed by using Microsoft Excel (Seattle, Wash.). Only clinical isolates were included in the data analysis. The percentage of *E. faecium* among total enterococci and the rate of VRE over time were analyzed by linear regression analysis. A *P* value of ≤0.05 was considered statistically significant.

From 1993 to 2002, 18,856 enterococci were identified in our clinical microbiology laboratory. The percentages of enterococci that were identified as *E. faecium* increased from 12.7% in 1993 to 22.2% in 2002 (*P* < 0.001) (Fig. 1). The percentages of *E. faecium* that were vancomycin resistant increased from 28.9% in 1993 to 72.4% in 2002 (*P* < 0.001). The percentages of *E. faecalis* that were vancomycin resistant increased from 0.2% in 1993 to 0.5% in 2002, but this increase was not statis-
rates of vancomycin resistance during this 10-year time period.

Since the total numbers of \textit{E. faecalis} remained stable during the 10-year period, the increasing numbers of \textit{E. faecium} were responsible for the increase in total enterococci. We observed significant increases in the rates of incidence of \textit{E. faecium} and VRE during the study period. Both of these trends are potentially problematic, as increasing rates of these organisms limit treatment options. While new antimicrobials, such as linezolid, daptomycin, and quinupristin-dalfopristin, have recently been developed to treat serious enterococcal infections, resistance to these agents has already emerged (4, 13, 16).

Addressing this trend for enterococci requires a combined infection control and antimicrobial utilization approach (12). An active surveillance program to detect colonization with VRE has been in place at our medical center for 8 years and is associated with a high rate of compliance. Despite utilization of the Centers for Disease Control Hospital Infection Control Practice Advisory Committee (HICPAC) (1) and Society for Healthcare Epidemiology of America (SHEA) (9) guidelines regarding control of this organism, we continue to observe increasing rates of VRE. Antibiotic prescribing practices also contribute to increasing rates of VRE (8, 15). While there is some controversy regarding which antimicrobial agents are most likely responsible, cephalosporins and vancomycin are consistent risk factors (8, 15).

Our study has several limitations. Since we reviewed only microbiological records to determine the numbers of enterococci, we were not able to determine the role infection control practices or antimicrobial utilization played in this changing epidemiology. We also did not determine whether these organisms represented community pathogens or were healthcare associated. Finally, since this was a clinical microbiology laboratory-based project, we were unable to distinguish colonization from infection.

In summary, there has been a significant increase in the numbers of enterococci that are \textit{E. faecium} and the proportion that are vancomycin resistant over a 10-year period at our medical center. Nearly three-quarters of our \textit{E. faecium} are vancomycin resistant, and this changing epidemiology has important clinical and infection control implications.

REFERENCES