Epidemiology and Infection

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Enterococcal bacteraemia: predictive and prognostic risk factors for ampicillin resistance

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Original Paper

Cite this article: Matsumura T, Nagao M, Nakano S, Yamamoto M, Matsumura Y, Ichiyama S (2018). Enterococcal bacteraemia: predictive and prognostic risk factors for ampicillin resistance. *Epidemiology and Infection* **146**, 2028–2035. https://doi.org/10.1017/S0950268818002479

Received: 1 September 2017
Revised: 3 August 2018
Accepted: 3 August 2018
First published online: 31 August 2018

Key words:
Ampicillin-resistance; bloodstream infections; enterococcus; epidemiology; mortality

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Abstract

To identify the predictive and prognostic factors associated with ampicillin-resistant enterococcal bacteraemia, we retrospectively reviewed demographic, microbiological and clinical data of patients attending the Kyoto University Hospital, Japan, between 2009 and 2015. Logistic regression and Cox regression analyses were performed to determine the predictive and prognostic factors, respectively. In total, 235 episodes of enterococcal bacteraemia were identified. As ampicillin susceptibility was uniform for *Enterococcus faecalis* isolates and almost all ampicillin-resistant isolates were *E. faecium*, bacteraemia due to these species was investigated separately. *E. faecalis* and *E. faecium* accounted for 41.7% (98/235) and 48.1% (113/235) of the isolates, respectively and 91.2% of all *E. faecium* were ampicillin resistant. Nosocomial *E. faecium* bacteraemia acquisition (odds ratio (OR), 13.6; 95% confidence intervals, 3.16–58.3) was associated with ampicillin-resistant isolates. Bacteraemia from an unknown source (hazard ratio (HR), 2.91; 95% CI 1.36–6.21) and an increased Pitt bacteraemia score (PBS) (HR, 1.36; 95% CI 1.21–1.52) were associated with 30-day mortality in *E. faecium* infections. Likewise, bacteraemia from an unknown source (HR, 4.17; 95% CI 1.25–13.9) and increased PBS (HR, 1.27; 95% CI 1.09–1.48) were associated with 30-day mortality in patients with *E. faecalis* bacteraemia. The empirical therapeutic administration of glycopeptides is recommended for patients with bacteraemia from an unknown source in whom severe *E. faecium* bacteraemia is suspected.

Introduction

Enterococcal species are significant pathogens in bloodstream infections (BSIs) and account in the USA, for approximately 4% and 9% of community-acquired and nosocomial bacteraemia, respectively, and are the second most common cause of nosocomial BSIs [1–4]. Similarly, in Europe, the prevalence of enterococcal BSIs appears to be increasing [5]. In Japan, enterococci are the third most common pathogen in nosocomial BSIs and the crude mortality rate of enterococcal bacteraemia is the second worst in Japanese university hospitals [6]. Worldwide, reported mortality rates for enterococcal BSI range from 14% to 48% [7–11]. The treatment of enterococcal bacteraemia is complicated by antimicrobial resistance as all *Enterococcus* species are inherently resistant to cephalosporins and acquired resistance to penicillins, aminoglycosides and glycopeptides is increasingly common [12]. Although the prevalence of vancomycin-resistant enterococci (VRE) is gradually increasing in North America and parts of Europe [5, 12], the prevalence of vancomycin-susceptible isolates remains high in many European countries and Japan [13]. Indeed, survey reports suggest that VRE isolates are relatively less common in enterococcal bacteraemia [13–18], while the ampicillin-susceptible species, e.g. *E. faecalis* [19], are more frequently isolated from blood cultures than their ampicillin-resistant counterparts, such as *E. faecium*. Ampicillin is not recommended as the first choice for empirical therapy to treat enterococcal bacteraemia in a setting with the frequent isolation of *E. faecium*. In regions where the prevalence of VRE is low, the susceptibility or resistance to ampicillin of isolates is the major concern when selecting initial antibiotics for treatment. In such a context, the aim of this study was to identify the predictive and prognostic factors associated with ampicillin-resistant enterococcal bacteraemia.

Methods

Study design and population

All patients aged 16 years or older with a first episode of enterococcal bacteraemia attending Kyoto University Hospital, a 1121-bed tertiary care hospital in Kyoto, Japan, were identified from a laboratory database of all positive blood cultures from 1 January 2009 to 31 December 2015. Patients were excluded who had more than one enterococcal species in

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blood cultures and if they had another bacteraemic episode with an enterococcal species different from that in the first episode during the study period. Study approval was obtained from the Institutional Review Board at Kyoto University and informed consent was waived due to the anonymous nature of the data.

Patient data

Electronic medical charts of all patients with enterococcal bacteraemia were reviewed for demographic (age and sex), microbiological (enterococcal species isolated from blood, antimicrobial susceptibility, the presence of mixed flora) and clinical data. Minimum inhibitory concentrations were determined by a broth microdilution test and interpreted according to standard criteria [20].

Clinical data included the acquisition site (nosocomial, community or intensive care unit (ICU)-acquired), underlying diseases, a prior operation within 6 months, presence of indwelling devices, immunosuppression, prior antibiotic exposure within 30 days, number of days from admission to onset, prior hospitalisation or ICU admission within 1 year, the isolation of enterococci from body sites other than blood within 1 year of the index culture, the source of bacteraemia and exclusion of contamination and 30-day mortality. An infection was designated as nosocomial if onset occurred more than 48 h after admission to hospital [21]. Likewise, an infection was considered as healthcare-associated community-onset if onset of bacteraemia occurred within 48 h and patients fulfilled any of the following criteria prior to the onset of bacteraemia: (1) received intravenous treatment at home, wound care, home-based nursing care or self-administered intravenous medical therapy within the previous 30 days; (2) visited a hospital or haemodialysis clinic or received intravenous chemotherapy within the prior 30 days (3) had been admitted to an acute care hospital for at least 2 days within the prior 90 days; or (4) had been admitted to a nursing home or long-term care facility [22]. A community-acquired infection was designated if onset was within 48 h of admission but did not meet the criteria for healthcare-associated community-onset infection. Likewise, an ICU-acquired infection was one where bacteraemia was diagnosed after 48 h following admission to ICU. A prior operation included any skin incisions and excluding puncture procedures. Neutropaenia was defined as an absolute neutrophil count <500/μl and severity of bacteraemia was based on the Pitt bacteraemia score (PBS) [23]. The degree of comorbidity was evaluated by the Charlson comorbidity index (CCI) [24].

Epidemiological characteristics

The proportions of healthcare-associated community-onset and nosocomial enterococcal bacteraemias each year were determined as well as the trend of the rate for ampicillin-resistant strains in all enterococcal bacteraemias. Similarly, we examined the trend of the antimicrobial use density (AUD) for penicillins, carbapenems, cephalosporins, aminoglycosides, fluoroquinolones and anti-methicillin-resistant *Staphylococcus aureus* (MRSA) antibiotics in the wider hospital and compared the results with the detection rate of ampicillin-resistant enterococcal strains in nosocomial bacteraemia.

Statistical analysis

The Cochran-Armitage test was used to determine trends in the proportion of healthcare-associated community-onset enterococcal bacteraemia cases and the isolation rate of ampicillin-resistant strains. Spearman's rank correlation analysis was used to test for association between antibiotic consumption and the rate of ampicillin-resistant strains. In the comparison of ampicillin-resistant and ampicillin-sensitive cases, categorical variables were compared using the Fisher exact test and continuous variables by the Mann–Whitney U test. The threshold for significance was a P-value <0.05. Factors with a P-value <0.10 in a single-variable analysis were entered into a logistic regression analysis. We conducted a backward stepwise procedure so that only significant variables were left in the model. In the comparison of 30-day survivors and non-survivors cases, categorical variables were compared using the Kaplan–Meier method and compared with a log-rank test and continuous variables by the Cox Proportional Hazards analysis. The threshold for significance was a P-value <0.05. For 30-day mortality, the CCI and factors with a P-value <0.10 in a single-variable analysis were entered into the Cox Proportional Hazards analysis. We also conducted a backward stepwise procedure so that only significant variables were left in the model. All statistical analyses were conducted using EZR version 1.33 (Saitama Medical Center, Jichi Medical University, Saitama, Japan) [25].

Results

A total of 255 patients with 272 episodes of positive blood cultures for enterococci were identified during the study period; five cases were considered as pseudobacteraemia on clinical review. There were 32 episodes where the same blood culture harboured different enterococcal species or multiple blood cultures with different enterococci in 15 patients. Therefore, 235 patients with single bacteraemic episodes were included in the study.

In total, 113 (48.1%), 98 (41.7%) and 24 (10.2%) episodes were caused by *E. faecium*, *E. faecalis* and other enterococci, respectively. Their antibiotic susceptibilities are shown in Supplemental Table S1. There were 124 (52.8%) ampicillin-susceptible and 111 (47.2%) ampicillin-resistant episodes of enterococcal bacteraemia.

All *E. faecalis* isolates were susceptible to ampicillin and as very few ampicillin-resistant non-*faecium* isolates were identified, we focused only on *E. faecium* cases for further analysis. The great majority of *E. faecium* bacteraemias (101; 89.4%) were designated as nosocomially acquired and the remainder as healthcare-associated community-onset cases. Figure 1 shows that over the study period the number of nosocomial *E. faecium* bacteraemias decreased from 25 to 6 episodes and the proportion of healthcare-associated community-onset episodes increased from 3.8% to 25%. The detection rate for ampicillin-resistant strains among all cases and nosocomial bacteraemias exceeded 81% and 87%, respectively. Healthcare-associated community-onset cases due to ampicillin-resistant strains were first identified at our hospital in 2012 and since 2014, all such isolates have been ampicillin resistant; significant increases were evident in the proportion of healthcare-associated community-onset cases ($P = 0.009$) and the isolation rate for ampicillin-resistant strains from bacteraemias ($P = 0.005$). The AUD trajectory for each antibiotic is shown in Supplemental Figure S1. Spearman's rank correlation coefficients for penicillins, carbapenems, cephalosporins, aminoglycosides, fluoroquinolones and anti-MRSA agents were 0.185 ($P = 0.691$), 0.482 ($P = 0.274$), −0.074 ($P = 0.875$), 0.259 ($P = 0.574$), −0.185 ($P = 0.691$) and −0.445 ($P = 0.317$), respectively. No correlation was observed between the amounts of β-lactams, aminoglycosides,

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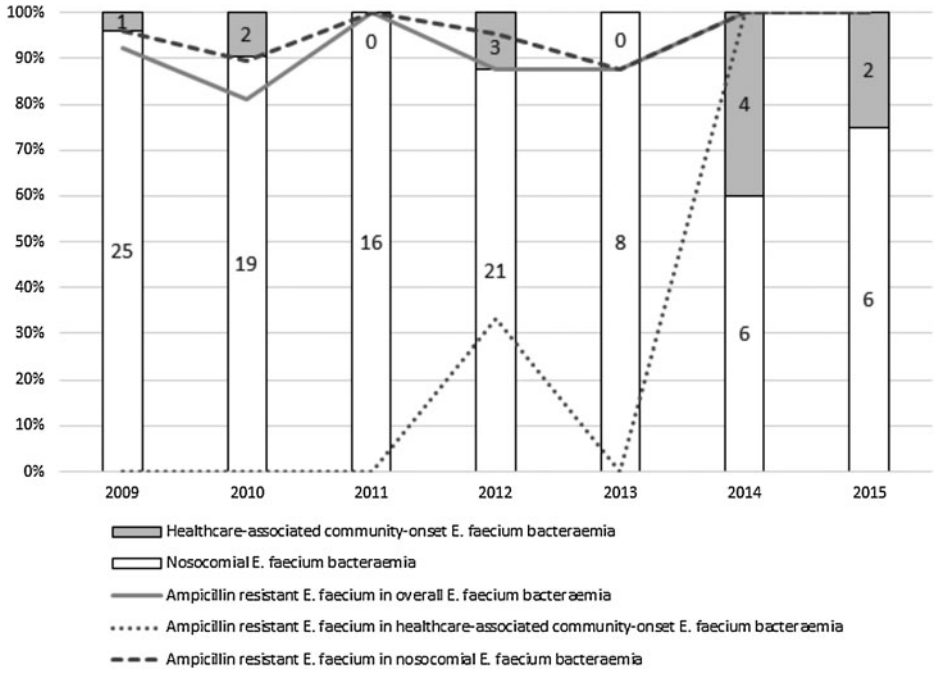


Fig. 1. Proportion of nosocomial and community-acquired *E. faecium* bacteraemia cases and the detection rate of ampicillin-resistant isolates.

fluoroquinolones and anti-MRSA agents prescribed and the detection rate for ampicillin-resistant *E. faecium*.

Table 1 compares the patients from whom ampicillin-resistant and ampicillin-susceptible *E. faecium* were isolated and shows that the factors most associated with ampicillin resistant cases in a single-variable analysis were nosocomial acquisition, comorbid liver disease, prior surgery, solid organ transplantation, administration of immunosuppressants, prior exposure to penicillins, carbapenems and sulfamethoxazole/trimethoprim and the duration from admission to onset as well as prior ICU hospitalisation. Conversely, only patient age and prior hospitalisation were significantly associated with ampicillin-susceptible *E. faecium* bacteraemia cases. By multivariable logistic regression analysis only nosocomial acquisition (odds ratio (OR), 13.6; 95% confidence interval (CI) 3.16–58.3, $P < 0.001$) was significantly associated with ampicillin-resistant *E. faecium*.

There were substantial differences between patient characteristics with *E. faecalis* and *E. faecium* bacteraemia (Supplemental Table S2) and therefore prognostic factors for both species were investigated separately. Table 2 shows that by single-variable analysis, the factor significantly associated with 30-day mortality in patients with *E. faecalis* was bacteraemia from an unknown source. By multivariable analysis, BSI from an unknown source remained significantly associated with 30-day mortality in patients with *E. faecalis* bacteraemia (HR, 4.17; 95% CI 1.25–13.9, $P = 0.020$) as well as a higher PBS (HR, 1.27; 95% CI 1.09–1.48, $P = 0.002$) (per 1-point increase in score).

A comparison of the clinical characteristics of patients who died within or survived 30 days with *E. faecium* bacteraemia are presented in Supplemental Table S3. By single-variable analysis, a number of factors were significantly associated with earlier death in patients with *E. faecium* bacteraemia including an indwelling central venous line and arterial line, mechanical ventilation, bone marrow transplantation/haematopoietic stem cell transplantation, neutropenia, prior exposure to sulfamethoxazole/trimethoprim, bacteraemia from an unknown source and increased PBS. Conversely, intra-abdominal infection was the sole factor significantly associated with patient survival for more than 30 days. Moreover, a Kaplan–Meier plot of survival time showed that the isolation of ampicillin-resistant *E. faecium* strains was not significantly associated with 30-day mortality ($P = 0.630$) (Fig. 2). Finally, by multivariable analysis, the factors significantly associated with 30-day mortality in patients with *E. faecium* bacteraemia were bacteraemia from an unknown source (HR, 2.91; 95% CI 1.36–6.21, $P = 0.006$) and higher PBS (HR, 1.36; 95% CI 1.21–1.52, $P < 0.001$) (per 1-point increase in score).

Discussion

This study investigated the predictive and prognostic factors for ampicillin-resistant enterococcal bacteraemia. As all the *E. faecalis* isolates were susceptible to ampicillin, we chose to focus on factors predictive of ampicillin-resistant strains in *E. faecium* bacteraemia alone and the key finding was that nosocomial acquisition was the factor most predictive in these cases. By contrast, prognostic factors for both *E. faecalis* and *E. faecium* bacteraemia were found to be unknown acquisition source and increased PBS and additionally for *E. faecium* cases, presentation with neutropaenia. Previous studies have highlighted exposure to penicillins and carbapenems as predictive factors for ampicillin-resistant enterococcal bacteraemia [13] and notable prognostic factors were clinical and co-morbid severity, nosocomial acquisition, isolation of *E. faecium* and several other underlying conditions [14–18]. However, in almost all of these studies, all enterococcus species were generally pooled into a single group and evaluated together [13–15, 17, 18]. Owing to differences in their epidemiology and ampicillin susceptibility, we investigated predictive factors of ampicillin-resistant strains in *E. faecium* alone and prognostic factors for both *E. faecalis* and *E. faecium* bacteraemia separately. The high proportion of *E. faecium* (48.1%) in our case series is noteworthy in contrast to some previous studies, which reported bacteraemia frequencies for this species ranging from 14.1% to 37.0% [13–18]. Moreover, our observed rate of ampicillin-resistant strains (47.2%) in

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Table 1. Comparison of the baseline demographics and clinical characteristics of patients with ampicillin-susceptible and ampicillin-resistant *Enterococcus faecium* bacteraemia

	Ampicillin susceptibility				
Variables	Susceptible (*n* = 10)		Resistant (*n* = 103)		*P*-value
Demographics					
Age, median (IQR)	74	(70–78)	61	(50–71)	0.015
Sex (males)	6	(60)	57	(55)	1
Nosocomial infection	5	(50)	96	(93)	0.001
ICU-acquired	0	(0)	31	(30)	0.059
Underlying disease					
Heart disease	4	(40)	25	(24)	0.276
Stroke/hemiplegia	1	(10)	8	(7.8)	0.580
Chronic pulmonary disease	2	(20)	8	(7.8)	0.216
Systemic autoimmune diseases	0	(0)	12	(12)	0.596
Chronic kidney disease	2	(20)	13	(13)	0.620
Diabetes mellitus	2	(20)	25	(24)	1
Liver disease	1	(10)	50	(49)	0.022
Hematological malignancy	1	(10)	25	(24)	0.449
Solid tumour	5	(50)	41	(40)	0.738
Prior operation (6 m)	1	(10)	51	(50)	0.020
Indwelling devices					
Central venous catheter	2	(20)	52	(51)	0.097
Arterial line	0	(0)	30	(29)	0.060
Urinary devices	1	(10)	33	(32)	0.277
Mechanical ventilation	0	(0)	23	(22)	0.209
Bile duct devices	1	(10)	24	(23)	0.454
Surgical drain	1	(10)	41	(40)	0.088
Immunosuppression					
Solid organ transplantation	0	(0)	33	(32)	0.033
BMT/HSCT	1	(10)	10	(9.7)	1
Neutropenia	0	(0)	22	(21)	0.205
Chemotherapy (30 days)	3	(30)	26	(25)	0.715
Immunosuppressant	2	(20)	60	(58)	0.041
Prior antibiotic exposure (30 days)	9	(90)	100	(97)	0.313
Penicillins	2	(20)	60	(58)	0.041
Cephalosporins	9	(90)	91	(88)	1
Carbapenems	1	(10)	54	(52)	0.016
Quinolones	5	(50)	47	(46)	1
Glycopeptides	1	(10)	43	(42)	0.086
Sulfamethoxazole/Trimethoprim	0	(0)	53	(52)	0.002
Days from admission to onset, median (IQR)	3	(1–43)	51	(18–91)	0.010
Prior hospitalisation (1 year)	10	(100)	69	(67)	0.031
Prior ICU admission (1 year)	0	(0)	49	(48)	0.005
Prior enterococcal isolation (1 year)	2	(20)	48	(47)	0.181
Source of bacteraemia					

(*Continued*)

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Table 1. (*Continued.*)

Variables	Ampicillin susceptibility				*P*-value
	Susceptible ($n = 10$)		Resistant ($n = 103$)		
Intra-abdominal	5	(50)	34	(33)	0.310
Catheter-related bloodstream infection	1	(10)	11	(11)	1
Febrile neutropenia	0	(0)	5	(4.9)	1
Urinary tract	1	(10)	6	(5.8)	0.487
Unknown	3	(30)	44	(43)	0.518
Other	0	(0)	3	(2.9)	1
Pitt bacteraemia score, median (IQR)	1	(0–2)	2	(1–6)	0.056
Charlson comorbidity index, median (IQR)	3	(2–5)	4	(3–5)	0.386

Data represent the number (%) of patients unless otherwise indicated.
IQR, interquartile range; ICU, intensive care unit; BMT, bone marrow transplantation; HSCT, hematopoietic stem cell transplantation.

Table 2. Comparison of the baseline demographics and clinical characteristics of 30-day survivors and non-survivors with *Enterococcus faecalis* bacteraemia

Variables	30-day mortality				*P*-value
	Survivors ($n = 84$)		Non-survivors ($n = 14$)		
Demographics					
Age, median (IQR)	71	(60–78)	67	(63–75)	0.784
Sex (males)	52	(62)	10	(71)	0.486
Polymicrobial bacteraemia	32	(38)	7	(50)	0.389
Persistent bacteraemia	7	(8.3)	1	(7.1)	0.865
Nosocomial infection	61	(73)	12	(86)	0.319
ICU-acquired	3	(3.6)	2	(14)	0.086
Underlying disease					
Heart disease	26	(31)	4	(29)	0.843
Stroke/hemiplegia	17	(20)	3	(21)	0.895
Chronic pulmonary disease	4	(4.8)	1	(7.1)	0.694
Systemic autoimmune diseases	4	(4.8)	0	(0)	0.422
Chronic kidney disease	7	(8.3)	1	(7.1)	0.865
Diabetes mellitus	16	(19)	2	(14)	0.678
Liver disease	17	(20)	4	(29)	0.442
Hematological malignancy	5	(6.0)	2	(14)	0.220
Solid tumor	39	(46)	7	(50)	0.803
Prior operation (6 m)	37	(44)	10	(71)	0.064
Indwelling devices					
Central venous catheter	31	(37)	6	(43)	0.665
Arterial line	7	(8.3)	2	(14)	0.489
Urinary devices	25	(30)	3	(21)	0.515
Mechanical ventilation	9	(11)	2	(14)	0.713
Bile duct devices	11	(13)	0	(0)	0.165
Surgical drain	15	(18)	5	(36)	0.128

(*Continued*)

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Table 2. (*Continued.*)

Variables	30-day mortality				*P*-value
	Survivors (*n* = 84)		Non-survivors (*n* = 14)		
Immunosuppression					
Solid organ transplantation	7	(8.3)	2	(14)	0.472
BMT/HSCT	2	(2.4)	1	(7.1)	0.318
Neutropenia	6	(7.1)	3	(21)	0.053
Chemotherapy (30 days)	14	(17)	5	(36)	0.071
Immunosuppressant	23	(27)	3	(21)	0.643
Prior antibiotic exposure	66	(79)	13	(93)	0.227
Penicillins	23	(27)	5	(36)	0.506
Cephalosporins	58	(69)	12	(86)	0.204
Carbapenems	17	(20)	2	(14)	0.616
Quinolones	20	(24)	2	(14)	0.418
Glycopeptides	19	(23)	3	(21)	0.932
Sulfamethoxazole/Trimethoprim	13	(16)	3	(21)	0.577
Days from admission to onset, median (IQR)	17	(1–47)	27	(7–62)	0.646
Prior hospitalisation (1 year)	47	(56)	8	(57)	0.979
Prior ICU admission (1 year)	20	(24)	5	(36)	0.340
Prior enterococcal isolation (1 year)	29	(35)	3	(21)	0.350
Source of bacteraemia					
Intra-abdominal	20	(24)	1	(7.1)	0.162
Catheter-related bloodstream infection	9	(11)	2	(14)	0.751
Febrile neutropenia	3	(3.6)	0	(0)	0.490
Urinary tract	19	(23)	0	(0)	0.055
Unknown	24	(29)	10	(71)	0.001
Other	9	(11)	1	(7.1)	0.693
Pitt bacteraemia score, median (IQR)	2	(1–4)	5	(1–9)	0.002
Charlson comorbidity index, median (IQR)	3	(2–6)	3	(2–8)	0.449

Data represent the number (%) of patients unless otherwise indicated.
IQR, interquartile range; ICU, intensive care unit; BMT, bone marrow transplantation; HSCT, hematopoietic stem cell transplantation.

bacteraemia was markedly higher than earlier reports of 9.8% to 40.6% [13, 15–17]. A possible explanation for this higher frequency may be due to a relatively high rate of ampicillin-resistant strains in Japan with frequencies ranging from 32.7% to 40.6% [13, 26]. Another reason for our high rate may be the specialist tertiary nature of a university hospital which provides advanced medical care for patients referred from community hospitals where almost all patients would have received earlier treatment, such as antibiotic therapy.

A notable finding of the study was the increase in the rate of ampicillin-resistant *E. faecium* in healthcare-associated community-onset bacteraemia. According to previous studies, prior hospitalisation and the use of β-lactams and fluoroquinolones were associated with colonisation of patients by ampicillin-resistant enterococci on admission [27, 28]. We did not find prior hospitalisation to be significantly associated with ampicillin-resistant bacteraemia and neither was there a correlation between the amounts of prescribed β-lactams and fluoroquinolones and rates of ampicillin-resistant strains, which might suggest an absence of antimicrobial selection pressure.

The multivariable analysis suggested that nosocomial acquisition was the only predictive factor for ampicillin-resistant *E. faecium* bacteraemia. This contrasts with a report from Spain that the previous administration of β-lactams and urinary catheterisation were predictive factors of such cases [18]. This difference might have been due to the proportions of nosocomial infection and ampicillin-resistant *E. faecium* which were markedly higher here (nosocomial infection: 67.3% *vs.* 89.3%, ampicillin-resistant *E. faecium*: 59.2% *vs.*91.2%) compared with the Spanish study. Predictive factors for ampicillin-resistant strains in enterococcal bacteraemias caused by all enterococcal species has been investigated previously in Japan [13] which reported that exposure to penicillins and carbapenems and bacteraemia related to mucositis with febrile neutropenia were risk factors for ampicillin-resistant

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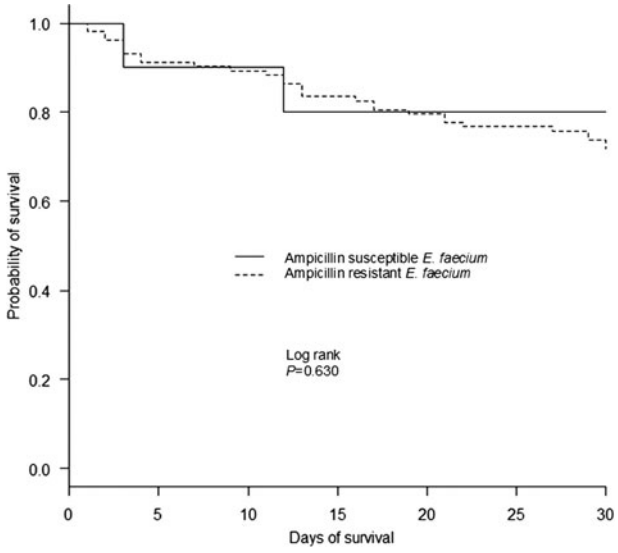


Fig. 2. Kaplan–Meier curve for all-cause 30-day mortality, according to the isolated *E. faecium* susceptibility to ampicillin.

strains [13]. Several studies have reported an association between the incidence of ampicillin-resistant enterococci and the prior administration of penicillins, cephalosporins, β-lactams, imipenem and fluoroquinolones [13, 18, 29–32].

The prognostic factors for 30-day mortality in patients with both *E. faecalis* and *E. faecium* bacteraemia were increased PBS and bacteraemia from an unknown source. The severity of both bacteraemia and comorbidities were previously reported as prognostic factors for enterococcal bacteraemia [15, 16, 18]. A population-based cohort study in Denmark which sought risk factors of 30-day mortality for *E. faecalis* and *E. faecium* bacteraemia separately identified an unknown focus of infection as one of the prognostic factors associated with *E. faecalis* [16]. Although the reasons for such an association are unclear, inadequate clinical investigation or an unrecognised primary focus of infection such as endocarditis may lead to antimicrobial treatment and surgical management for an insufficient duration [10, 11].

We were unable to estimate the attributable mortality for enterococcal bacteraemia due to the absence of records of direct cause of death on medical charts and autopsies were rarely performed. We also did not include a matched control group without enterococcal bacteraemia. However, the survival time analysis indicated that the isolation of ampicillin-resistant strains from the patients' blood was not associated with a poor prognosis and may imply that ampicillin susceptibility of the infecting strain is not critical to patient outcomes; nevertheless, more complex patient conditions such as focus of bacteraemia, comorbidity, neutropenia and severity of infection, most notably affected the prognosis.

The primary limitations of the study were associated with the retrospective design. Some data were missing or inaccurate due to incomplete documentation in the medical records and we were unable to determine the source of certain bacteraemias due to insufficient investigations; this, therefore, increased the number of cases with an unknown source. In addition, even with electronic medical charts, it was difficult to identify the exact time when the index cultures were collected and antibiotics were given. We obtained patient referral medical information if the patients had previously received medical care in different hospitals in order to collect as much information as possible on variables. However, some data from other hospitals at which the patients may have been treated in the previous year were sometimes incomplete. We did not record the duration of the antibiotic therapy and glycopeptide use and may have underestimated the incidence of enterococcal bacteraemia due to false-negative blood cultures; additionally, blood cultures might not have been collected in some cases in which bacteraemia was suspected.

In conclusion, the prognostic factors for *E. faecalis* bacteraemia were the severity of BSI and infection from an unknown source; these factors also applied to *E. faecium* bacteraemia. The Nosocomial acquisition was the sole predictive factor for ampicillin-resistant *E. faecium* bacteraemia. We suggest that when severe *E. faecium* bacteraemia is suspected in a setting with a high-degree of ampicillin resistance then glycopeptides should be administered as empirical treatment.

Supplementary material. The supplementary material for this article can be found at https://doi.org/10.1017/S0950268818002479

Conflict of interest. None.

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