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Both early-onset and late-onset ventilator-associated pneumonia are caused mainly by potentially multiresistant bacteria

Abstract Objective: To compare the causative pathogens of early-onset and late-onset ventilator-associated pneumonia (VAP) diagnosed by bronchoalveolar lavage quantitative cultures. Most previous reports have been based on endotracheal aspirate cultures and gave uncertain findings. Design: Prospective evaluation of consecutive patients with clinical suspicion for VAP. Setting: Multidisciplinary intensive care unit of a university hospital. Patients and participants: During a 3-year period 473 patients with clinical suspicion of VAP entered the study. Diagnosis of VAP was confirmed by cultures of bronchoalveolar lavage (> 10^4 cfu/ml) specimens in 408 patients. Interventions: Protected bronchoalveolar lavage samples were taken. Initial antibiotic therapy was modified upon bronchoalveolar lavage culture results. Measurements and results: Among 408 patients 191 had earlyonset (<7 days mechanical ventilation) and 217 late-onset (>7 days) VAP. Potentially multiresistant bacteria, mainly Pseudomonas aeruginosa and methicillin-resistant Staphylococcus aureus (MRSA), were

the most commonly isolated pathogens in both types of VAP. No difference was noted in the contribution of potentially multiresistant pathogens (79% vs. 85%), P. aeruginosa (42% vs. 47%), or MRSA (33% vs. 30%) between early-onset and lateonset VAP. Initial antibiotic therapy was modified in 58% of early-onset VAP episodes and in 36% of lateonset VAP episodes. No difference in mortality was found between the two types of VAP. Conclusions: Both early-onset and late-onset VAP were mainly caused by potentially multiresistant bacteria, most commonly P. aeruginosa and MRSA. Antimicrobial agents against these pathogens should be prescribed empirically, at least in our institution.

Keywords Ventilator-associated pneumonia · Resistant bacteria · Mechanical ventilation · Bronchoalveolar lavage

Introduction

The initial, empirical antibiotic therapy of ventilator-associated pneumonia (VAP) is often based on timing of its occurrence in relation to the onset of mechanical ventilation [1]. This is due to reported differences between causal pathogens associated with early-onset (<5-7 days mechanical ventilation) compared to late-onset ($\geq 5-7$ days mechanical ventilation) VAP [2, 3]. Early-onset VAP is most often caused by core micro-organisms (e.g.,

Streptoccoccus pneumoniae, enteric Gram-negative bacilli, methicillin-susceptible *Staphylococcus aureus*), whereas late-onset VAP is most commonly due to potentially multiresistant bacteria (e.g., *Pseudomonas aeruginosa*, *Acinetobacter baumanni*, methicilline resistant *S. aureus*) [3, 4, 5, 6]. However, Ibrahim et al. [7] reported that both early-onset and late-onset VAP may be associated with similar, usually multiresistant, pathogens. Nevertheless, most of the above reports were based on studying patients with VAP diagnosed on the basis of clinical criteria and endotracheal aspirate cultures, which make their conclusions uncertain [8].

Therefore the aim of our study was to compare the causative pathogens of early-onset and late-onset VAP diagnosed by bronchoalveolar lavage (BAL) quantitative cultures, because BAL samples are considered to give more accurate results than endotracheal aspirates [9, 10, 11, 12].

Methods

Patients

The study was conducted at the University of Thrace teaching hospital (500 beds) during a 3-year period (September 2000-September 2003) and was approved by the Human Studies Ethics Committee. All patients admitted to the multidisciplinary intensive care unit (15 beds) were potentially eligible for this investigation. Inclusion criteria were: age over 18 years, at least 48 h of mechanical ventilation, and clinical suspicion of VAP [3, 7, 12], defined by new and persistent (present for >72 h) infiltrate on chest radiography, plus two of the following items: (a) purulent tracheal secretions (>25 neutrophils per high power field using Gram's stain), (b) body temperature above 38.3°C or below 36°C, and (c) leukocytosis (>10,000 cells/µl) or leukopenia (<5,000 cells/µl). Patients were excluded if (a) they were temporarily transferred to our intensive care unit due to lack of available bed in another hospital, (b) they had received immunosuppressants or long-term corticosteroid therapy (>0.5 mg/kg prednisolone/day for >1 month) during the previous year, (c) they had neutropenia (leukocyte count <1,000/µl or neutrophil count <500/µl), or (d) they had concomitant acquired immune-deficiency syndrome.

Of the 1,467 patients admitted to the intensive care unit during the study period 1,059 (72%) required mechanical ventilation longer than 48 h, and 535 of these (51%) had clinical suspicion of VAP. Sixty-two (12%) were excluded (28 had received immunosuppressants or long-term corticosteroid therapy, 8 were temporarily transferred to our intensive care unit, 3 had acquired immune deficiency syndrome, and 23 had neutropenia). Therefore 473 patients with clinical suspicion of VAP were evaluated with BAL. Diagnosis of VAP was finally confirmed by quantitative cultures of BAL in 408 (86%), and these patients represented the study population.

Study design, data collection, and end-points

In patients with clinical suspicion for VAP in daily rounds, protected BAL samples were taken by fiberoptic bronchoscopy, as previously described [13]. BAL specimens were cytocentrifuged and divided into two preparations, one for Gram stain and the other for quantitative culture. The culture results were available within 2 days. The diagnosis of VAP was confirmed if at least one bacterial species grew at a concentration above the predetermined threshold (>10⁺ cfu/ml) [6]. BAL specimens were always obtained before introduction of any antibiotics. Initial antibiotic therapy was modified in the light of quantitative BAL culture results when at least one cultured isolate proved resistant in vitro to the administered regimen or combination therapy was necessary because of isolation of P. aeruginosa [8]. Deescalation of the initial regimen for a narrower spectrum alternative after BAL results was not used. The treatment regimen that replaced the initial one consisted of at least one antibiotic to which all isolates were susceptible in vitro. In the presence of P. aeruginosa at least two active agents (combination therapy) were used [8]. Quantitative BAL cultures yielding less than significant growth of organisms were not taken into account in order to modify the initial antibiotic therapy; nevertheless these patients were not included as only patients with bacterial species growth greater than 10^4 cfu/ml in BAL cultures were studied. Only the first VAP episode of each patient was recorded. No patient received oropharyngeal or selective digestive decontamination. Histamine type 2 receptors antagonists were used in all patients to prevent upper digestive tract hemorrhage.

At the time of study entry and before bronchoscopy we recorded each patient's age, sex, admission diagnostic category (medical vs. surgical, trauma vs. nontrauma surgical), concomitant diseases, indication for mechanical ventilation, severity of illness based on Acute Physiology and Chronic Health Evaluation (APACHE) II [14], and severity of organ dysfunction based on Sequential Organ Failure Assessment (SOFA) [15]. The presence of potential specific risk factors for development of VAP were also recorded and included the administration of histamine type 2 receptor antagonists, antacids, sucralfate, corticosteroids, or vasopressors, tracheostomy, dialysis, reintubation, duration of mechanical ventilation, and previous 2 weeks use of antibiotics. The initial, empirical antibiotic regimen was always chosen by the attending physicians, usually based on American Thoracic Society recommendations [3].

We compared the findings in patients with early-onset VAP (<7 days mechanical ventilation) with those in patients with lateonset VAP (\geq 7 days mechanical ventilation). We used a cutoff of 7 days because it corresponded to the median time of occurrence of VAP in the 408 patients studied, and because mechanical ventilation lasting 7 days or more is associated with increased likelihood of infection with a potentially multiresistant mico-organism [2, 8]. Among 408 patients 191 (47%) had early-onset and 217 (53%) lateonset VAP.

The primary end-point was comparison of causative pathogens between early-onset and late-onset VAP. Secondary end-points were the comparison of frequency of initial antibiotic treatment modification, and 15-day, 28-day, intensive care unit, and hospital mortality between early-onset and late-onset VAP.

Statistics

Statistical analysis was performed using Norusis MJ SPSS version 11.0 (SPSS, USA). The χ^2 test with Yates' correction was used to compare categorical variables. Continuous variables, normally or abnormally distributed, were compared using Student's *t* test or Wilcoxon's signed rank test, respectively. Differences with a *p* value less than 0.05 were considered statistically significant.

Results

Characteristics of patients with early-onset and late-onset VAP at study entry are presented in Table 1. With the exception of vasopressors use and tracheostomy, which

Table 1 Characteristics of 408 patients with ventilator associated pneumonia (<i>VAP</i>) con-		Early-onset VAP (<i>n</i> =191)	Late-onset VAP (<i>n</i> =217)	р
firmed by quantitative cultures	Age, median (years; range)	61 (21-75)	59 (25-87)	0.18
of bronchoalveolar lavage at	Sex: male/female	105/86	140/77	0.06
study entry (early-onset devel-	SOFA score, mean ±SD	4.8 ± 4.1	4.3±4.3	0.20
oping in less than 7 days of	APACHE II score, mean ±SD	15.5 ± 4.5	16.2 ± 4.8	0.14
mechanical ventilation, late-	Diagnostic category			
onset developing in 7 days of	Medical	114 (60%)	119 (55%)	0.37
mechanical ventilation or more,	Surgical, nontrauma	47 (25%)	68 (31%)	0.16
SOFA Sequential Organ Failure	Surgical, trauma	30 (16%)	30 (14%)	0.69
Assessment, APACHE II Acute	Underlying malignancy	17 (9%)	32 (15%)	0.09
Physiology and Chronic Health	Indication for MV			
Evaluation II, ARDS acute	ARDS	12 (6%)	20 (9%)	0.36
respiratory distress syndrome,	Status asthmaticus	11 (6%)	12 (6%)	0.90
COPD chronic obstructive	Drug overdose	13 (7%)	16 (7%)	0.97
pulmonary disease,	Abdominal surgery	5 (3%)	13 (6%)	0.15
MV mechanical ventilation)	Other than abdominal surgery	21 (11%)	19 (9%)	0.55
	Community-acquired pneumonia	22 (12%)	16 (7%)	0.20
	Exacerbation of COPD	25 (13%)	19 (9%)	0.21
	Congestive heart failure	17 (9%)	23 (11%)	0.68
	Neurological emergency	42 (22%)	38 (18%)	0.31
	Miscellaneous	23 (12%)	41 (19%)	0.07
	Risk factors for VAP			
	Reintubation	30 (16%)	32 (15%)	0.89
	Tracheostomy	28 (15%)	80 (37%)	< 0.001
	Vasopressors	55 (29%)	97 (45%)	0.001
	Dialysis	35 (18%)	42 (19%)	0.88
	Corticosteroids	95 (50%)	119 (55%)	0.35
	Previous use of antibiotics	161 (84%)	187 (86%)	0.69
	Duration of MV, mean ±SD	2.3±1.2	8.9±6.5	0.03
	Mortality, patients died			
	15-day	8 (4%)	10 (5%)	0.97
	28-day	17 (9%)	21 (10%)	0.92
	Crude intensive care unit	24 (13%)	30 (14%)	0.81
	Crude hospital	28 (15%)	38 (18%)	0.51

were more frequent in late-onset VAP, other characteristics did not differ significantly between the two groups.

Pathogens identified and initial treatment modifications

Micro-organisms responsible for VAP are shown in Table 2. Potentially multiresistant bacteria, mainly P. aeruginosa and MRSA, were the most commonly isolated pathogens in both early-onset and late-onset VAP. No significant difference was noted in the contribution of potentially multiresistant pathogens between early-onset and late-onset VAP (79% vs. 85%, p=0.06). This was also the case for P. aeruginosa, the most common Gramnegative pathogen (42% vs. 47%, p=0.26) and for MRSA, the most common Gram-positive pathogen (33% vs. 30%, p=0.39). Characteristics of patients with early-onset VAP caused either by P. aeruginosa/MRSA or by other pathogens at study entry are presented in Table 3. Risk factors of early-onset VAP due to P. aeruginosa or MRSA were previous use of antibiotics, corticosteroid therapy, vasopressors use, trauma, and neurological emergency.

Table 2 Micro-organisms responsible for 408 episodes of ventilator associated pneumonia (*VAP*) confirmed by quantitative cultures of bronchoalveolar lavage (*early-onset* developing in less than 7 days of mechanical ventilation, *late-onset* developing in 7 days of mechanical ventilation or more, *MRSA* methicillin-resistant *Staphylococcus aureus*, *MSSA* methicillin-sensitive *Staphylococcus aureus*)

	Early-onset VAP (<i>n</i> =191)	Late-onset VAP (<i>n</i> =217)	р
Potentially multiresistant	219 (79%)	257 (85%)	0.06
bacteria			
Pseudomonas aeruginosa	116 (42%)	141 (47%)	0.26
MRSA	93 (33%)	90 (30%)	0.39
Acinetobacter baumannii	6 (2%)	12 (4%)	0.30
Stenotrophomonas	4 (1%)	14 (5%)	0.04
maltophilia			
Other bacteria	59 (21%)	45 (15%)	0.06
MSSA	15 (5%)	9 (3%)	0.21
Haemophilus influenzae	7 (3%)	5 (2%)	0.66
Escherichia coli	5 (2%)	9 (3%)	0.51
Morganella morganii	5 (2%)	7 (2%)	0.88
Enterococcus species	6 (2%)	4 (1%)	0.65
Streptococccus pneumoniae	6 (2%)	4 (1%)	0.65
Other cocci	15 (5%)	7 (2%)	0.08
Total number of bacteria	278 (100%)	302 (100%)	-

Table 3 Characteristics of 191 patients with early-onset ventilator associated pneumonia (*VAP*) developing in less than 7 days of mechanical ventilation caused by *P. aeruginosa* (*PA*) or methicillin-resistant *Staphylococcus aureus* (*MRSA*) vs. by other pathogens at study entry (*SOFA* Sequential Organ Failure Assessment, *APACHE II* Acute Physiology and Chronic Health Evaluation II, *ARDS* acute respiratory distress syndrome, *COPD* chronic obstructive pulmonary disease, *MV* mechanical ventilation)

Characteristics	PA/MRSA (<i>n</i> =142)	Other pathogens (<i>n</i> =49)	р
Age, median (years, range)	56 (21-75)	59 (28-69)	0.23
Sex: male/female	78/64	27/22	0.98
SOFA score, mean ±SD	4.9 ± 3.6	4.6 ± 2.2	0.55
APACHE II score,	16.5 ± 3.2	15.1 ± 2.8	0.47
mean ±SD			
Diagnostic category			
Medical	75 (53%)	39 (80%)	0.002
Surgical, nontrauma	39 (27%)	8 (16%)	0.17
Surgical, trauma	28 (20%)	2 (4%)	0.01
Underlying malignancy	10 (7%)	7 (14%)	0.21
Indication for MV			
ARDS	7 (5%)	5 (10%)	0.33
Status asthmaticus	8 (6%)	3 (6%)	0.81
Drug overdose	8 (6%)	5 (10%)	0.44
Abdominal surgery	4 (3%)	1 (2%)	0.82
Other than abdominal	18 (13%)	3 (6%)	0.31
surgery			
Community acquired	12 (8%)	10 (20%)	0.04
pneumonia			
Exacerbation of COPD	17 (12%)	8 (16%)	0.59
Congestive heart failure	16 (11%)	1 (2%)	0.09
Neurological emergency	40 (28%)	2 (4%)	< 0.001
Miscellaneous	17 (12%)	6 (12%)	0.83
Risk factors for VAP			
Reintubation	19 (13%)	11 (22%)	0.20
Tracheostomy	17 (12%)	11 (22%)	0.12
Vasopressors	47 (33%)	8 (16%)	0.04
Dialysis	25 (18%)	10 (20%)	0.82
Corticosteroids	85 (60%)	10 (20%)	< 0.001
Previous use of antibiotics	141 (99%)	20 (41%)	< 0.001
Duration of MV, mean ±SD	2.9±1.6	2.1±1.8	0.75
Mortality, deaths			
15-day	7 (5%)	1 (2%)	0.64
28-day	15 (11%)	2 (4%)	0.27
Crude intensive care unit	22 (15%)	2 (4%)	0.06
Crude hospital	26 (18%)	2 (4%)	0.02

Among 408 VAP episodes 236 (58%) were monomicrobial (one bacterial species in BAL) and 172 (42%) polymicrobial (two or more bacterial species in BAL) in origin. Monomicrobial were 124 of 191 episodes of early-onset VAP (65%) and 112 of 217 episodes of late-onset VAP (52%); polymicrobial were the remaining 67 episodes of early-onset VAP (35%) and 105 episodes of late-onset VAP (48%; p=0.008).

After BAL culture results, initial antibiotic therapy was modified in 189 of 408 VAP episodes (46%): in 111 of 191 episodes of early-onset VAP (58%) and in 78 of 217 episodes of late-onset VAP (36%; *p*<0.001). Strains responsible for this modification in early-onset and lateonset VAP are presented in Table 4. In early-onset VAP

Table 4 Strains responsible for inadequate initial antibiotic therapy in episodes of ventilator associated pneumonia (*VAP*) developing in less than 7 days of mechanical ventilation (*early-onset*) and developing in 7 days of mechanical ventilation or more (*late-onset*) (*MRSA* methicillin-resistant *Staphylococcus aureus*)

Organism (n)	Organism
	sensitivity
P. aeruginosa (14)	Sensitive ^a
P. aeruginosa (16)	Sensitive ^a
P. aeruginosa (14)	Sensitive ^a
P. aeruginosa (4)	Sensitive ^a
P. aeruginosa (12)	Resistant
P. aeruginosa (9)	Resistant
P. aeruginosa (15)	Resistant
MRSA (16)	Resistant
MRSA (15)	Resistant
MRSA (4)	Resistant
P. aeruginosa (13)	Resistant ^b
P. aeruginosa (11)	Resistant ^b
P. aeruginosa (9)	Resistant
P. aeruginosa (5)	Resistant
P. aeruginosa (4)	Resistant
P. aeruginosa (2)	Resistant
P. aeruginosa (4)	Resistant ^b
MRSA (19)	Resistant
MRSA (21)	Resistant
	P. aeruginosa (14) P. aeruginosa (16) P. aeruginosa (14) P. aeruginosa (14) P. aeruginosa (12) P. aeruginosa (12) P. aeruginosa (15) MRSA (16) MRSA (15) MRSA (15) MRSA (4) P. aeruginosa (13) P. aeruginosa (11) P. aeruginosa (11) P. aeruginosa (2) P. aeruginosa (2) P. aeruginosa (4) P. aeruginosa (4) MRSA (19)

^a But a second drug was required

^b In at least one antibiotic, thus addition of at least one susceptible drug was required

initial antibiotic regimen modification was due to isolation of *P. aeruginosa* in 76 VAP episodes (69%), MRSA in 27 VAP episodes (24%), and both *P. aeruginosa* and MRSA in the remaining 8 of 111 VAP episodes (7%). Among 84 *P. aeruginosa* strains 36 (43%) were resistant to the initial regimen, and 48 (57%) were sensitive but modification of initial therapy was required due to addition of a second effective agent. In late-onset VAP initial antibiotic treatment modification was due to isolation of *P. aeruginosa* in 38 VAP episodes (49%), MRSA in 30 (38%), and both *P. aeruginosa* and MRSA in the remaining 10 of 78 VAP episodes (13%). In contrast to early-onset, in late-onset VAP all strains of *P. aeruginosa* responsible for treatment modification were resistant to the initial regimen.

Antibiotic therapy used at pre-BAL and post-BAL periods, and susceptibility to antibiotics of cultured pathogen strains in early-onset and late-onset VAP are shown in Tables 5 and 6, respectively.

Table 5Antimicrobials usedbefore bronchoalveolar lavage(*pre-BAL*) and after (*post-BAL*)in 408 episodes of ventilatorassociated pneumonia (*VAP*)

Antimicrobial therapy	Early-onset VAP (n=191)		Late-onset VAP (n=217)	
	Pre-BAL	Post-BAL	Pre-BAL	Post-BAL
Monotherapy	108	23	0	0
Combinationtherapy	83	168	217	217
Aminoglycosides	34	97	144	142
Ticarcillin/clavulanicacid	26	42	65	63
Piperacillin/tazobactam	36	57	73	62
Aztreonam	10	4	23	15
Ceftazidime	25	37	38	30
Cefamandole, cefotaxime	55	21	16	5
Clindamycin	16	2	2	2
Imipenem	30	67	70	68
Quinolones	31	66	67	35
Vancomycin	14	113	50	151
Antifungalagents	1	4	10	11
Total	278	510	558	584

Table 6 Percentages of susceptibility to ten antimicrobial agents of the 580 strains responsible for 408 episodes of ventilator associated pneumonia (*VAP*) (*early-onset* developing in less than 7 days of mechanical ventilation, *late-onset* developing in 7 days of mechanical ventilation or more)

Antimicrobial agent	Early-onset VAP (191/278 ^a)	Late-onset VAP (217/302 ^a)	р
Ticarcillin/ clavulanic acid	96	85	< 0.001
Piperacillin/ tazobactam	88	85	0.34
Cefamandole	71	13	< 0.001
Cefotaxime	75	13	< 0.001
Ceftazidime	85	74	0.001
Aztreonam	50	34	< 0.001
Imipenem	87	67	< 0.001
Gentamycin	62	60	0.69
Amikacin	62	50	0.005
Ciprofloxacin	75	65	0.009

^a Number of episodes/number of pathogens

Mortality rate

Crude intensive care unit and hospital mortality of patients with VAP were 13% and 16%, respectively. No significant difference was found in 15-day, 28-day, intensive care unit, or hospital mortality between earlyonset and late-onset VAP (Table 1). Hospital mortality was higher in early-onset VAP caused by *P. aeruginosa* and/or MRSA than in early-onset VAP caused by other pathogens (Table 3).

Discussion

The main finding of the present study, which confirmed the responsible pathogens of VAP by BAL quantitative cultures, was that both early-onset and late-onset VAP cases are mainly caused by potentially multiresistant micro-organisms, most commonly *P. aeruginosa* and MRSA. As a result, initial antibiotic therapy was modified more frequently in early-onset than in late-onset VAP because attending physicians did not expect such high prevalence of potentially multiresistant organisms in early-onset VAP.

Several studies have shown that early-onset and lateonset VAP are caused by different pathogens [2, 3, 4, 5, 6]. In contrast, Ibrahim et al. [7] reported that pathogens associated with early-onset and late-onset VAP may be similar and frequently multiresistant; however, because microbiological diagnosis of VAP in this study was usually made on the basis of endotracheal aspirate cultures, which were not always quantitative, its findings were considered uncertain [8]. The present study confirms by BAL quantitative cultures for the first time, to our knowledge, that early-onset and late-onset VAP episodes are caused by similar pathogens, which are usually multiresistant.

The importance of our findings is that they may influence antimicrobial prescribing practices in the intensive care unit. Indeed, these findings suggest that antimicrobial agents against P. aeruginosa and MRSA should be prescribed empirically, at least in our institution, to patients suspected of having either early-onset or lateonset VAP. This may help to reduce the occurrence of inadequate or ineffective antimicrobial therapy, which has been associated with poorer patient outcomes [16, 17, 18]. According to the American Thoracic Society consensus statement [3], early-onset VAP is usually due to core pathogens, and monotherapy is recommended. Highly resistant P. aeruginosa and MRSA are not included among these core bacteria. Therefore initial antibiotic therapy prescribed by the attending physicians, usually based on the American Thoracic Society recommendations, resulted in undertreating of patients with early-onset VAP of the present study, and therapy modification was required in 58% of cases. In patients who develop late-onset VAP the most commonly encountered pathogens are potentially multiresistant Gram-negative bacteria, including P. aeruginosa and Acinetobacter species as well as MRSA, and the American Thoracic Society recommendations for the empirical treatment of these patients include the use of combination antimicrobial therapy with drugs that are active against *P. aeruginosa* and vancomycin for severely ill patients with suspected MRSA infection [3]. Despite prescription of rather adequate combination antibiotic therapy in most patients with late-onset VAP of our study, initial therapy modification was required in 36% of VAP episodes because of frequent isolation of resistant *P. aeruginosa* strains or not inclusion in the initial regimen of drugs active against MRSA (Table 4).

Several studies [19, 20, 21, 22] have demonstrated that the success rate of empirical monotherapy in early-onset VAP is similar to that of combination therapy, and this is probably the case in patients who have not received antibiotics previously [2, 8]. However, the findings of the present study as well as those of others [7] suggest that combination antipseudomonal therapy and vancomycin are initially required in early-onset VAP.

The high rate of early-onset VAP due to potentially multiresistant bacteria in this study may be due in part to the prior use of antibiotics; over 99% of patients with early-onset VAP caused by *P. aeruginosa* and/or MRSA received antimicrobial therapy prior to the development of this infection (Table 3). Previous investigations have shown a strong association between prior antibiotic use and the subsequent development of VAP, particularly VAP caused by potentially antibiotic-resistant pathogens [2, 7, 23, 24, 25, 26, 27]. Another factor potentially contributing to the explanation of our findings in patients with early-onset VAP is frequent short-term corticosteroid use (Table 3). In fact, corticosteroids predispose to VAP with *P. aeruginosa* [3].

Our findings of similar mortality rates in early-onset and late-onset VAP (Table 1) are consistent with those reported by other investigators in different countries [7, 28, 29]. One potential explanation suggested by the results of the present and other [7] studies is that patients with either early-onset or late-onset VAP have similar rates of infection with high-risk multiresistant pathogens, mainly *P. aeruginosa* and MRSA, which are associated with higher rates of attributable hospital mortality [24]. Hospital mortality in the present study was higher in early-onset VAP caused by *P. aeruginosa* and/or MRSA than in early-onset VAP caused by other pathogens (Table 3).

The present study has the advantage of using protected BAL quantitative cultures to identify the causative pathogens in a large sample size of VAP. Although we are aware of the potential limitations of this method [11, 30, 31, 32], it is currently considered one of the best invasive techniques for the microbiological diagnosis of VAP [9, 10, 11, 12]. However, the findings of this study may not be applicable to other intensive care units with lower rates of VAP caused by P. aeruginosa and MRSA. That is because our patient population may not be similar to those of other intensive care units. The high incidence of prior exposure to antibiotics among our patients may not be representative of practices at other institutions. Significant variations in the bacterial pathogens associated with VAP have been found between four European intensive care units [33]; multiresistant bacteria associated with early-onset VAP were isolated in two of these, thus supporting the existence of variability between intensive care units of different hospitals in terms of the causal agents of VAP.

In conclusion, the present study demonstrates that in our intensive care unit both early-onset and late-onset VAP were caused mainly by potentially multiresistant bacterial pathogens, most commonly *P. aeruginosa* and MRSA. Consequently, the empirical antibiotic therapy prescribed was inadequate in a significant proportion of patients, and therapy modification was required. These findings, rather than providing information generally applicable, emphasize the need of intensive care unit-specific knowledge of causal agents associated with VAP [34]; this information can influence local antibiotic prescribing practices and reduce the rate of administration of inadequate antimicrobial therapy.

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