

Clinical manifestations of pneumonia according to the causative organism in patients in the intensive care unit

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Background/Aims: Whether the causative organism influences the clinical course of pneumonia in the intensive care unit (ICU) is controversial. We assessed the clinical manifestations and prognosis of pneumonia according to the causative pathogens in patients in a medical ICU.

Methods: A retrospective observational study was performed in a medical ICU. Among 242 patients who were admitted to the ICU, 103 who were treated for pneumonia were analyzed.

Results: The causative pathogen was identified in 50 patients (49.0%); 22 patients (21.6%) had multidrug-resistant (MDR) pathogens. The distribution of causative micro-organisms was *Staphylococcus aureus* (20%), *Pseudomonas* species (16%), *Klebsiella pneumoniae* (14%), and *Acinetobacter baumannii* (12%). No significant difference in ICU mortality rate, duration of ICU stay, duration of mechanical ventilation, or frequencies of re-intubation and tracheostomy were detected based on the identification of any pathogen. In sub-analyses according to the pneumonia classification, the number of pathogens identified did not differ between pneumonia types, and a higher incidence of identified MDR pathogens was detected in the hospital-acquired pneumonia group than in the community-acquired or health-care-acquired pneumonia groups. However, the clinical outcomes of pneumonia according to identification status and type of pathogen did not differ significantly between the groups.

Conclusions: Neither the causative micro-organism nor the existence of MDR pathogens in critically ill patients with pneumonia was associated with the clinical outcome of pneumonia, including ICU mortality. This result was consistent regardless of the pneumonia classification.

Keywords: Pneumonia; Pathogenicity; Drug resistance, multiple; Intensive care units

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INTRODUCTION

Pneumonia is the second most common cause of in-hospital infection. Pneumonia is very common in the intensive care unit (ICU) setting and can be fatal. The inci-

dence of pneumonia is about 17% in the medical ICU [1] but can be 6 to 20 times higher in mechanically ventilated patients [2]. The mortality rate of hospital-acquired pneumonia (HAP) depends on the clinical situation, but can range from 20% to 50% [3,4]. Determining the prog-

nosis of pneumonia is important for anticipating the disease course and establishing a proper management plan.

The causative micro-organism can influence the clinical presentation, outcome of antibiotic therapy, and prognosis in patients with pneumonia. However, whether the causative organism influences the clinical course of pneumonia in the ICU is controversial. Identifying a pathogen by culture can be a poor prognostic factor [5], and infection with multidrug-resistant (MDR) pathogens such as *Acinetobacter baumannii* or methicillin-resistant *Staphylococcus aureus* (MRSA) is a risk factor for hospital mortality [6-8]. However, one study reported that the pathogen classification or the existence of MDR pathogens does not affect the mortality rate after adjusting for the effect of antibiotics [9]. Although one study has reported the epidemiology and causative micro-organisms of pneumonia in Korea [10], the prognosis of pneumonia according to the causative micro-organism in the ICU is not well known. Thus, in this study, we elucidated the clinical manifestations and prognosis of patients with pneumonia according to the causative pathogen in the medical ICU.

METHODS

Study design and participants

A retrospective observational study was performed in the medical ICU of the Seoul National University Hospital between January 2011 and August 2011. We included patients with pneumonia treated in the medical ICU. Patients were enrolled if they had pneumonia on admission or developed pneumonia during their ICU stay. A total of 242 patients were admitted and treated in the medical ICU over the study period; in addition, data from 102 patients with pneumonia were analyzed retrospectively.

Pneumonia was clinically classified based on the American Thoracic Society/Infectious Disease Society of America guidelines [11,12]. According to these guidelines, healthcare-associated pneumonia (HCAP) was defined as pneumonia in any patient admitted to an acute care hospital for ≥ 2 days within 90 days of the infection; who resided in a nursing home or long-term care facility; who received recent intravenous antibiotic therapy,

chemotherapy, or wound care within 30 days of onset of the current infection; or who attended a hospital or hemodialysis clinic within 30 days. HAP was defined as pneumonia that developed ≥ 48 hours after admission. Community-acquired pneumonia (CAP) was defined as pneumonia that did not meet any of the HCAP and HAP criteria.

The causative organisms were considered to be micro-organisms that had been isolated from specimens, including blood, bronchoalveolar lavage fluid, bronchial wash, and pleural effusion, and which grew to greater than a threshold concentration in quantitative cultures. Specimens obtained by endotracheal aspiration or in sputum expectorant were evaluated as appropriate using counts of white blood cells and epithelial cells, and the micro-organisms identified were considered the causative pathogen. Growth below the threshold was considered to be caused by colonization or contamination. *Legionella* and *Streptococcus* urinary antigen tests, viral polymerase chain reaction, and antigen tests were also used to identify the pathogens. Pathogens may have been confirmed in additional samples after a patient developed pneumonia, but a secondary infection due to a hospital- or ICU-acquired pathogen was ruled out. Specimens sampled within 3 days after a patient developed pneumonia were considered significant.

Empirical antibiotic regimens to treat pneumonia were reviewed, and their response and relevance to subsequent changes in the antibiotic regimen were analyzed. The initial empirical antibiotic regimen was administered according to the American Thoracic Society/Infectious Disease Society of America guidelines [11,12]. Initial non-responders to empirical antibiotics were defined as cases in which the initial antibiotic was changed due to expansion or a switch in the antibiotic spectrum due to persistence or worsening of the clinical course within 1 week after developing pneumonia. This did not include cases in which empirical antibiotics were changed to the targeted regimen based on drug susceptibility in clinically improving patients.

We analyzed ICU mortality, duration of ICU stay, duration of mechanical ventilation, frequency of re-intubation, and frequency of tracheostomy as clinical outcomes of pneumonia. These outcomes were compared between patients with CAP, HCAP, and HAP.

This retrospective data collection was approved by the

Institutional Review Board of Seoul National University (IRB no: H-1208-150-424) and was performed in accordance with the Declaration of Helsinki.

Statistical analysis

Data are presented as means and standard deviations for continuous variables and as numbers (percentages) for categorical variables. We analyzed baseline characteristics, clinical outcomes, and prognoses according to specific pathogens using the Kruskal-Wallis test. We used the Mann-Whitney test to compare data for prognosis according to whether a pathogen or MDR pathogen was identified. Associations between prognostic factors and survival rate were investigated using the Cox proportional-hazard analysis after adjusting for age, sex, Acute Physiology and Chronic Health Evaluation (APACHE) II score [13], Sequential Organ Failure Assessment (SOFA) score [14], Charlson comorbidity index [15], pneumonia

severity index [16], the identification of the pathogen or MDR pathogen, and the pneumonia classification. Likelihood ratio tests were used to examine the goodness of fit of the model, and no lack of fit was found. Adjusted odd ratios (aOR) are presented with 95% confidence intervals (CIs). A $p < 0.05$ was considered significant. Data were analyzed using SPSS version 18 (SPSS Inc., Chicago, IL, USA).

RESULTS

Baseline characteristics and clinical outcomes of the patients

During the study period, 242 patients were admitted to the ICU; 102 had pneumonia, and all were eligible for this study. The study population was comprised of 20 patients (19.6%) with CAP, 26 (25.5%) with HCAP, and 56

Table 1. Baseline characteristics of patients with pneumonia treated in the intensive care unit

Characteristic	Overall (n = 102)	Pathogen not identified (n = 52)	Pathogen identified (n = 50)	p value
Age, yr	64.1 ± 14.3	66.0 ± 11.5	62.1 ± 16.5	0.410
Male sex	77 (75.5)	43 (82.7)	34 (68)	0.086
APACHE II score	26.1 ± 9.4	26.0 ± 9.4	26.2 ± 9.5	0.815
SOFA score	8.9 ± 4.4	8.5 ± 3.7	9.4 ± 5.1	0.541
Pneumonia severity index	142.7 ± 39.7	142.3 ± 41.6	143.1 ± 38.1	0.723
Charlson comorbidity index	3.7 ± 2.5	3.7 ± 2.7	3.6 ± 2.4	0.837
Comorbidity				
Chronic lung disease	29 (28.4)	17 (32.7)	12 (24)	0.333
Chronic heart disease	8 (7.8)	3 (5.8)	5 (10)	0.429
Chronic liver disease	17 (16.7)	6 (11.5)	11 (22)	0.158
Chronic renal disease	15 (14.7)	5 (9.6)	10 (20)	0.141
Solid tumor	42 (41.2)	28 (53.8)	14 (28)	0.008
Diabetes mellitus	32 (31.4)	12 (23.1)	20 (40)	0.067
Hematologic malignancy	18 (17.6)	6 (11.5)	12 (24)	0.101
Pneumonia type				
Community-acquired pneumonia	20 (19.6)	11 (21.2)	9 (18)	0.690
Healthcare-associated pneumonia	26 (25.5)	15 (28.8)	11 (22)	0.430
Hospital-acquired pneumonia	56 (54.9)	26 (50)	30 (60)	0.313
MDR pathogen	22 (21.6)	0	22 (44)	< 0.001

Values are presented as mean ± SD or number (%).

APACHE, Acute Physiology and Chronic Health Evaluation; SOFA, Sequential Organ Failure Assessment; MDR, multidrug-resistant.

(54.9%) with HAP.

The baseline characteristics and clinical outcomes of the patients treated for pneumonia in the ICU are presented in Table 1. Mean age of the entire population was 64.1 ± 14.3 years, and 75.5% were men. No significant differences were found between the CAP, HCAP, and HAP groups for age, sex distribution, or severity as evaluated by the APACHE II score and the SOFA score. However, the Charlson comorbidity index and pneumonia severity index scores were significantly higher in the HCAP and HAP groups than in the CAP group; the scores did not differ between the HCAP and HAP groups.

The causative pneumonia pathogens were identified in 50 patients (49.0%). No significant differences in baseline characteristics were observed according to the pathogens identified except for a history of solid tumor

within 5 years and the presence of a MDR pathogen.

Distribution of causative pathogens and clinical outcomes

Among 50 cases in which a pathogen was identified, sputum (24, 48%), blood (17, 34%), endotracheal aspirate (12, 24%), bronchoalveolar lavage (4, 8%), serology (4, 8%), and bronchial wash (2, 4%) were the specimens in which pathogens were confirmed. The pathogens causing pneumonia were identified in nine patients (45%) with CAP, in 11 (42.3%) with HCAP, and in 30 (53.6%) with HAP. No significant difference in the incidence of pathogens was observed among the pneumonia types after adjusting for culture frequency. The incidence of pneumonia with a MDR pathogen was 22 patients (21.6%) in the entire population. The incidence was sig-

Table 2. Microetiological diagnoses and prognoses of pneumonia when the causative pathogen was identified (n = 50)

Variable	Frequency	MDR proportion	ICU mortality, %	ICU LOS, day	MV duration, day
Gram-positive bacteria					
<i>Staphylococcus aureus</i>	10 (20)	6 (60)	4 (40)	13.0 ± 8.8	7.7 ± 7.4
<i>Streptococcus pneumoniae</i>	3 (6)	0	1 (33.3)	40 ± 17.6	22.7 ± 26.7
<i>Enterococcus faecium</i>	1 (2)	1 (100)	1	5	5
Gram-negative bacteria					
Enterobacteriaceae					
<i>Klebsiella pneumoniae</i>	7 (14)	2 (28.6)	4 (57.1)	7.4 ± 5.5	2.6 ± 3.0
<i>Acinetobacter baumannii</i>	6 (12)	5 (83.3)	3 (50)	12.7 ± 15.5	12.8 ± 15.5
<i>Escherichia coli</i>	3 (6)	2 (66.7)	1 (33.3)	11.0 ± 10.1	3.3 ± 4.9
<i>Enterobacter aerogenes</i>	2 (4)	1 (50)	0	13.0 ± 17.0	12.5 ± 17.7
Nonfermentative gram-negative bacilli					
<i>Pseudomonas</i> species	8 (16)	5 (62.5)	3 (37.5)	13.1 ± 8.0	5.9 ± 8.0
<i>Stenotrophomonas maltophilia</i>	2 (4)	2 (100)	2 (100)	4.0 ± 1.4	3.5 ± 2.1
Mycobacterium					
<i>Mycobacterium tuberculosis</i>	1 (2)	0	0	52	9
Non-tuberculosis <i>Mycobacterium</i>	1 (2)	0	1 (100)	1	0
Fungi					
<i>Aspergillus</i> species	5 (10)	0	3 (60)	23.2 ± 29.8	19.0 ± 29.1
Virus					
<i>Cytomegalovirus</i>	4 (8)	0	2 (50)	20.5 ± 14.6	12.5 ± 8.2
Atypical pathogen					
<i>Pneumocystis jiroveci</i>	2 (2.0)	0	1 (50)	14.5 ± 6.4	13.5 ± 7.8
<i>Mycoplasma</i> species	1 (1.0)	0	1 (100)	15	15

Values are presented as number (%) or mean ± SD.

MDR, multidrug-resistant; ICU, intensive care unit; LOS, length of stay; MV, mechanical ventilation.

nificantly higher in the HAP group than in the HCAP and CAP groups (HAP, 32.1%; HCAP, 11.5%; CAP, 5.0%; $p = 0.015$), but was not different between the CAP and HCAP groups. The MDR micro-organisms were: carbapenem-resistant *A. baumannii*, MRSA, coagulase-negative *Staphylococcus*, vancomycin-resistant enterococci, extended-spectrum β -lactamase-producing *Escherichia coli*, *Klebsiella*, and MDR *Pseudomonas* species.

Table 2 shows the distribution of specific pathogens and the respective prognoses. *S. aureus* was the most frequently isolated pathogen among all patients in all groups (10, 20%). *Pseudomonas* species were detected in eight patients (16%), *Klebsiella pneumoniae* in seven (14%), and *A. baumannii* in six (12%). No significant difference was observed in the clinical outcomes between specific pathogens.

Regimen and response to empirical antibiotic treatment

The empirical antibiotics and their responses are shown in Table 3. The most commonly used antibiotics as an empirical regimen, including combination therapy, were β -lactam/ β -lactamase inhibitors (62, 60.8%), fol-

Table 3. Empirical antibiotic regimen and response (n = 102)

Variable	No. (%)
Empirical antibiotic treatment	
β -Lactam/ β -lactamase inhibitor	62 (60.8)
Quinolone	57 (55.9)
Carbapenem	20 (19.6)
Vancomycin	19 (18.6)
Cephalosporin	15 (14.7)
Trimethoprim/sulfamethoxazole	14 (13.7)
Antifungal agents	10 (9.8)
Antiviral agents	5 (4.9)
Macrolides	5 (4.9)
Colistin	3 (2.9)
Initial responder to empirical antibiotic therapy	59 (57.8)
Antibiotics response in cases with identified pathogens (n = 50)	
Initially appropriate antibiotics	30 (60)
Subsequent change to targeted antibiotics	28 (56)
Appropriate change according to drug susceptibility	25 (89.3)

lowed by quinolones (57, 55.9%). About 58% of the patients responded initially to the empirical antibiotic regimen. Appropriate antibiotics were initially used in 60% of cases when a retrospective review was performed considering pathogen drug susceptibility and the antibiotic spectrum in cases in which the pathogen was identified. In these cases, a subsequent change to the targeted antibiotic occurred in 56% of cases, and most (89.3%) were appropriate considering the drug susceptibility of the pathogen.

Clinical outcomes according to identification of the causative organism and pneumonia classification

The prognoses of pneumonia based on identifying the pathogens are shown in Table 4. The ICU mortality rate was 47.1% in the entire study population, and did not differ according to whether the pathogen was identified (unidentified pathogen 46.2% vs. identified pathogen 48.0%, $p = 0.853$). Other clinical outcomes such as duration of ICU stay, need for and duration of mechanical ventilation, and frequencies of re-intubation and tracheostomy were different based on whether the pathogen was identified. These results were consistent with cases in which MDR pathogens were identified.

A subgroup analysis was performed according to pneumonia type, and the clinical outcomes did not differ significantly between the groups.

Prognostic factors for mortality

A Cox proportional-hazard analysis was performed to evaluate the relative risk of ICU mortality caused by pneumonia, and the data are presented in Table 5. The ICU mortality rate was significantly higher in the group with higher APACHE II and SOFA scores in both univariate and multivariate analyses ([APACHE II score: aOR, 1.07; 95% CI, 1.03 to 1.11]; [SOFA score: aOR, 1.26; 95% CI, 1.14 to 1.38]). Age was also a significant prognostic factor for ICU mortality in the multivariate analysis (aOR, 1.04; 95% CI, 1.01 to 1.07). The pneumonia severity index was significantly associated with ICU mortality in the univariate analysis, but significance was lost after adjustment. Age, sex, the Charlson comorbidity index, identification of the pathogen or MDR pathogen, initial responder to empirical antibiotic therapy, and pneumonia classification were not significant prognostic indicators in the ICU mortality evaluation.

Table 4. Clinical outcomes and prognoses of pneumonia according to whether the pathogen was identified

Variable	Pathogens			MDR pathogens		
	Absence	Presence	p value	Absence	Presence	p value
Overall (n = 102)	52	50		80	22	
ICU mortality	24 (46.2)	24 (48)	0.853	38 (47.5)	10 (45.5)	0.865
ICU length of stay, day	19.0 ± 26.8	15.4 ± 15.4	0.176	18.7 ± 24.3	11.8 ± 7.6	0.305
MV duration, day	15.5 ± 27.3	10.1 ± 13.7	0.060	14.4 ± 24.1	7.0 ± 7.1	0.062
Re-intubation	9 (17.3)	5 (10)	0.286	13 (16.3)	1 (4.5)	0.189
Tracheostomy	13 (25)	11 (22)	0.722	18 (22.5)	6 (27.3)	0.777
CAP group (n = 20)	11	9		19	1	
ICU mortality	2 (18.2)	5 (55.6)	0.160	6 (31.6)	1 (100)	0.350
ICU length of stay, day	14.9 ± 10.1	15.9 ± 16.0	0.909	15.4 ± 13.1	15	0.794
MV duration, day	11.5 ± 10.4	7.3 ± 7.7	0.340	9.3 ± 9.4	15	0.257
Re-intubation	3 (27.3)	2 (22.2)	1.000	5 (26.3)	0	1.000
Tracheostomy	4 (36.4)	0	0.094	4 (21.1)	0	1.000
HCAP group (n = 26)	15	11		23	3	
ICU mortality	7 (46.7)	2 (18.2)	0.217	9 (39.1)	0	0.294
ICU length of stay, day	15.5 ± 8.3	18.0 ± 18.6	0.612	17.4 ± 13.9	9.7 ± 6.7	0.377
MV duration, day	11.9 ± 7.8	11.4 ± 14.9	0.275	12.6 ± 11.4	5.0 ± 5.6	0.137
Re-intubation	3 (20)	1 (9.1)	0.614	4 (17.4)	0	1.000
Tracheostomy	1 (6.7)	2 (18.2)	0.556	3 (13)	0	1.000
HAP group (n = 56)	26	30		38	18	
ICU mortality	15 (57.7)	17 (56.7)	0.939	23 (60.5)	9 (50)	0.461
ICU length of stay, day	22.7 ± 36.8	14.2 ± 14.5	0.277	21.1 ± 32.4	12.0 ± 8.1	0.493
MV duration, day	19.2 ± 37.5	10.5 ± 14.8	0.226	18.1 ± 33.0	6.9 ± 7.3	0.106
Re-intubation	3 (11.5)	2 (6.7)	0.655	4 (10.5)	1 (5.6)	0.662
Tracheostomy	8 (30.8)	9 (30)	0.951	11 (28.9)	6 (33.3)	0.741

Values are presented as number (%) or mean ± SD.

MDR, multidrug-resistant; ICU, intensive care unit; MV, mechanical ventilation; CAP, community-acquired pneumonia; HCAP, healthcare-associated pneumonia; HAP, hospital-acquired pneumonia.

DISCUSSION

In our study, patients with HCAP and HAP had significantly more severe comorbidity and pneumonia than those of patients with CAP, but the number of pathogens identified did not differ between the pneumonia types. More MDR pathogens were identified in the HAP than those in the CAP and HCAP groups. However, the influence of the pathogen or MDR pathogen identified on the clinical outcomes, such as ICU mortality rate, ICU duration, duration of mechanical ventilation, and frequencies of re-intubation and tracheostomy, was not significant in the entire population or in subgroups ac-

ording to the pneumonia classification.

It is controversial whether the pneumonia prognosis differs according to whether the pathogen is identified. In previous studies on the pneumonia prognosis, the severities of underlying disease and comorbidities were evaluated as important prognostic factors. Some HCAP and HAP studies have shown that age, multi-organ dysfunction, septic shock, the Charlson comorbidity index, and SOFA and APACHE II scores are significant prognostic factors for patients with pneumonia [6,9,17]. Other studies of patients with severe CAP who required mechanical ventilation or ICU admission reported acute renal failure, septic shock, and the Simpli-

Table 5. Risk of intensive care unit mortality according to prognostic factors in the entire population

Prognostic factor	Univariate		Adjusted	
	OR (95% CI)	p value	OR (95% CI)	p value
Age, yr	1.01 (0.99–1.02)	0.595	1.04 (1.01–1.07)	0.003
Male sex	0.63 (0.32–1.22)	0.168	0.59 (0.28–1.25)	0.169
APACHE II score	1.06 (1.03–1.09)	< 0.001	1.08 (1.04–1.12)	< 0.001
SOFA score	1.20 (1.12–1.28)	< 0.001	1.31 (1.18–1.45)	< 0.001
Charlson comorbidity index	1.03 (0.92–1.14)	0.658	1.14 (0.99–1.32)	0.066
Pneumonia severity index	1.01 (1.00–1.02)	0.027	0.998 (0.99–1.01)	0.680
Identified pathogen	1.21 (0.68–2.14)	0.523	1.86 (0.83–4.17)	0.131
Multidrug-resistant pathogen	1.39 (0.68–2.84)	0.365	0.52 (0.21–1.29)	0.159
Initial responder to empirical antibiotic therapy	1.03 (0.58–1.83)	0.924	0.62 (0.33–1.15)	0.131
Pneumonia classification				
Community-acquired pneumonia	1 (reference)		1 (reference)	
Healthcare-associated pneumonia	0.92 (0.34–2.48)	0.872	0.35 (0.10–1.22)	0.100
Hospital-acquired pneumonia	1.56 (0.69–3.55)	0.289	1.62 (0.53–4.91)	0.395

OR, odds ratio; CI, confidence interval; APACHE, Acute Physiology and Chronic Health Evaluation; SOFA, Sequential Organ Failure Assessment.

fied Acute Physiology Score II as significant prognostic factors [18,19].

In this study, the evaluation of ICU mortality according to the prognostic factors of pneumonia showed that higher APACHE II and SOFA scores were significantly associated with a lower survival rate, but the identification status and the microbe causative of pneumonia were not. This result suggests that the severity of pneumonia or underlying disease may be a more important prognostic factor than microbial etiology, and these data coincide with those of a recent prospective cohort study in Spain, which showed that comorbidities are a stronger determinant of mortality than microbial etiology in a comparison between CAP and HCAP [20].

Several limitations in this study should be mentioned. This study was performed retrospectively in a single tertiary referral hospital; thus, there may have been selection bias because several of the patients enrolled had severe or advanced disease and some patients may have been immunocompromised. The small study population may have limited the statistical power and subgroup analysis. In the pathogen evaluation, MRSA and gram-negative rods may be more easily detected because they are well seen in gram-stained specimens and cultures than anaerobes and viruses. Finally, the initial an-

tibiotic therapy and responses were not analyzed in this study.

In conclusion, we found that neither identifying the causative micro-organism nor the existence of a MDR pathogen was associated with the clinical outcomes of pneumonia, including ICU mortality. The severity of the underlying disease or pneumonia itself may be a significant prognostic indicator, regardless of whether the causative micro-organism is identified or if the pneumonia is classified.

KEY MESSAGE

1. The causative pathogen was identified in 49.0% of patients with pneumonia treated in the medical intensive care unit (ICU); 21.6% of patients had a multidrug-resistant pathogen.
2. In the clinical outcomes of pneumonia including ICU mortality, the severity of the underlying disease or pneumonia itself may be a significant prognostic indicator, regardless of the causative micro-organism or the pneumonia classification.

Conflict of interest

No potential conflict of interest relevant to this article was reported.

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