



Performance of pneumococcal urinary antigen test in patients with community-onset pneumonia: a propensity score-matching study

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Background/Aims: Although pneumococcal urinary antigen tests (PUATs) have universally been used for the diagnosis of pneumococcal pneumonia, data on the efficacy of these exams are limited. The objective of our study was to investigate the clinical impact of the PUAT in patients with community-onset pneumonia (CO-pneumonia).

Methods: We conducted a retrospective cohort study of patients diagnosed with CO-pneumonia. Patients were classified according to their PUAT results and were matched using the propensity score-matching method. The primary outcome was 30-day mortality.

Results: A total of 1,257 patients were identified and 163 (13.0%) demonstrated positive PUAT results. The sensitivity and specificity values of PUAT for overall pneumococcal pneumonia were 56.5% and 91.4%, respectively. In the full cohort, there were no significant differences in 30-day mortality between the two groups (6.1% in the positive PUAT group vs. 8.2% in the negative PUAT group, $p = 0.357$). However, in the propensity-matched cohort, the 30-day mortality rates were lower in the positive PUAT group (5.6% vs. 17.4%, $p = 0.001$). With respect to secondary outcomes, the proportion of patients with potentially drug-resistant pathogens, changes in antibiotics, and failure rates of initial antibiotic therapy were significantly lower in the positive PUAT group than in the negative PUAT group of the propensity-matched cohort.

Conclusions: We found that the sensitivity of the index test was low and specificity was high in this clinical setting. And our findings suggest that positive PUAT results may be associated with favorable clinical outcomes in patients with CO-pneumonia.

Keywords: Pneumonia; Streptococcus pneumoniae; Urinalysis; Antigens; Mortality

INTRODUCTION

Pneumonia is the leading cause of infectious disease-related deaths in adults worldwide [1]. Selecting appropriate pathogen-directed antibiotics in patients with community-onset pneumonia (CO-pneumonia) can de-

crease treatment cost, drug-related adverse events, and antibiotic resistance and allow for the narrowing of empiric antibiotic therapy [2-4]. Despite the development of diagnostic techniques for pneumonia, causative organisms have not been detected in more than half of the patients [5]. Determining the etiology of pneumonia has

remained challenging because of several reasons, including difficulties in obtaining good-quality sputum and the unreliability of subsequent culture results, low sensitivity of blood cultures, and the administration of antibiotics before sample collection [5,6]. *Streptococcus pneumoniae* is the most frequently isolated pathogen; therefore, initial empirical antibiotic treatment regimens for most patients with community-acquired pneumonia (CAP) aim at covering *S. pneumoniae* [3]. The early identification of *S. pneumoniae* may lead to the selection of more reliable pathogen-targeted antibiotics [3].

The pneumococcal urinary antigen test (PUAT) is an assay widely used to identify the C-polysaccharide antigen common to all serotypes excreted into the urine using a membrane immunochromatographic test [7]. In addition to the advantages of rapidity and simplicity, it has a high sensitivity and specificity over Gram stain and sputum cultures, as well as availability in patients who cannot demonstrate expectorated sputum [3]. In addition, even after antibiotic therapy has been started, the PUAT has the ability to detect pneumococcal pneumonia [3,8]. Current guidelines recommend that the PUAT can only be performed in patients with severe CAP or moderate- or high-severity pneumonia [3,9].

To date, the clinical benefits derived from the results of PUAT have not been fully established [10]. Thus, the aim of our study was to investigate the diagnostic performances of the PUAT, and the association between the results of PUAT and clinical outcomes in patients with CO-pneumonia.

METHODS

Study design and population

We conducted a retrospective cohort study of patients with CO-pneumonia at a university affiliated medical center. Adult patients (aged ≥ 18 years) who were hospitalized with pneumonia between January 2012 and December 2015 were investigated. Patients were identified by the use of the international diagnostic codes version 10 to screen for possible cases as follows: J18.0 to 18.9 as representative codes of pneumonia in the primary discharge diagnosis [11]. After reviewing the medical records and radiological findings of relevant patients, we confirmed the diagnosis of pneumonia as the presence

of a new infiltrate on a chest radiograph and by clinical signs [2]. We excluded the following types of patients: (1) those who were readmitted due to pneumonia within 10 days of leaving the hospital; (2) those who were transferred from other hospitals after hospitalization for > 48 hours; (3) those with obstructive pneumonia; (4) those who had immunocompromised status, such as neutropenia (absolute neutrophil count $< 1,500$ cells/ μL) after chemotherapy or human immunodeficiency virus infection; (5) those who did not receive guideline-concordant antibiotic therapy; and (6) those who had hospital-acquired pneumonia (HAP). Although the concept of healthcare-associated pneumonia (HCAP) was eliminated in the revised 2016 American Thoracic Society (ATS)/Infectious Disease Society of America (IDSA) guidelines for the management of HAP and ventilator-acquired pneumonia [12], we included HCAP as a category of CO-pneumonia in the present study.

Urine samples were collected in the emergency department or ward in the first 24 hours to detect urinary antigens for *S. pneumoniae* using BinaxNOW (Binax Inc., Scarborough, ME, USA). According to the results of PUAT, we classified the study patients into positive and negative PUAT groups. Empirical guideline-concordant antibiotics were maintained without regard to the results of tests. Demographics, radiological findings, laboratory findings, microbiological results, and clinical outcomes were compared between the two groups.

Ethical consideration

Ethical committees approved to review clinical data of relevant patients obtained from medical records such as clinical parameters, laboratory, radiological, and microbiological findings. Information obtained during the study was kept confidential and only intended for research purpose.

The study protocol was approved by the Ethical Review Committee of Jeju National University Hospital (Institutional Review Board no. 2018-04-001). Informed consent was waived because of the retrospective nature of the study.

Microbiology and antibiotics

Sampling to determine the microbial etiology of pneumonia included sputum, tracheobronchial aspirates, bronchoalveolar lavage fluid, pleural fluid, or blood

through a semi-quantitative manner. The antibiotic sensitivity of all isolates was determined using a disc diffusion method. Serologic tests were performed to detect antibodies against *Mycoplasma pneumoniae* or *Chlamydia pneumoniae*. A patient was considered to have a potentially drug-resistant (PDR) pathogen if one of the following pathogens was isolated: methicillin-resistant *Staphylococcus aureus* (MRSA), *Pseudomonas aeruginosa*, extended-spectrum beta-lactamase (ESBL)-producing or carbapenem-resistant *Klebsiella pneumoniae* and *Escherichia coli*, *Acinetobacter baumannii*, or *Stenotrophomonas maltophilia* [13].

Changes in antibiotic regimens were defined as either escalation or de-escalation after culture sensitivities or clinical stabilities were identified. Inappropriate antibiotic therapy was noted if the empirical antibiotic treatment was not effective against the identified pathogen based on *in vitro* susceptibility testing [14]. Initial treatment failure was defined as death during initial treatment or a change of initial therapeutic agent after 48 hours due to clinical instability [15]. Guideline-concordant antibiotic therapy for CAP was defined as antibiotic regimens recommended by the 2007 ATS/IDSA guidelines for CAP: specifically, beta-lactam plus macrolide or fluoroquinolone [3]. Since the data of the present study were collected before the 2016 revised ATS/IDSA guidelines were released, guideline-concordant antibiotic therapy for HCAP was defined as antibiotic regimens recommended by the 2005 ATS/IDSA guidelines, i.e., anti-pseudomonal beta-lactam plus either a fluoroquinolone or an aminoglycoside, along with anti-MRSA coverage if risk factors were present [16].

Statistical analyses

Data are presented as median (interquartile range) for continuous variables and as a number (%) for categorical variables. Continuous variables were compared using the Student's *t* test for normally distributed variables and the Mann-Whitney *U* test for non-normally distributed variables. Categorical variables were compared using the Pearson chi-square test, and Fisher's exact test was used when any cell contained less than five items. To evaluate the reliability of the PUAT, we calculated its sensitivity and positive and negative predictive values, using the following three different reference groups of patients as the gold standard: (1) definitive pneumococcal pneu-

monias (CO-pneumonia with *S. pneumoniae* isolated in blood or pleural fluid culture); (2) probable pneumococcal pneumonias (CO-pneumonia with *S. pneumoniae* as the predominant morphotype on Gram stain or culture of sputum, transtracheal aspirates, and bronchoalveolar lavage); and (3) all pneumococcal pneumonias (definite plus probable) [17]. When determining specificity, we considered the control group as all patients without a diagnosis of pneumococcal pneumonia, including those with unknown etiology. We also calculated positive and negative likelihood ratios (LRs) as a measure of the extent to which the pretest odds were altered by the test results; low negative LR (< 0.1) and high positive LR (> 10) findings were considered to be useful for ruling out and ruling in decisions, respectively [18].

To improve the balance of baseline characteristics and reduce the effects of selection bias and potential confounding in this retrospective cohort study, estimated propensity scores were used to match the positive PUAT group to the negative PUAT group. The propensity score was calculated by logistic regression analysis using the covariates of baseline characteristics (Table 1). Standardized differences were estimated for all baseline covariates before and after matching to assess prematch imbalance and postmatch balance. All statistical analyses were performed using SPSS version 18.0 (SPSS Inc., Chicago, IL, USA). All tests were two-sided, and *p* values of < 0.05 were considered to be statistically significant.

RESULTS

Study population

Fig. 1 shows patient enrolment. Overall, 1,257 patients were analyzed, of whom 163 (13.0%) had positive PUAT results and 1,094 (87.0%) had negative PUAT results. The proportion of patients with positive PUAT results was similar in both CAP and HCAP groups (13.1% vs. 12.8%, *p* = 0.875). The microbiological methods by which *S. pneumoniae* was identified in 115 patients as a cause of CO-pneumonia are as follows: blood cultures in seven patients (6.1%), pleural fluid in two patients (1.7%), transtracheal aspiration in 15 patients (13.0%), bronchoalveolar lavage in six patients (5.2%), and sputum in 87 patients (75.7%). In two patients, *S. pneumoniae* was simultaneously identified in blood and sputum cultures,

Table 1. Baseline characteristics of the patients

Characteristic	Entire cohort (n = 1,257)			Propensity score-matched cohort (n = 322)		
	Positive PUAT (n = 163)	Negative PUAT (n = 1,094)	p value	Positive PUAT (n = 161)	Negative PUAT (n = 161)	p value
Age, yr	73 (66-81)	72 (61-80)	0.124	73 (66-81)	74 (65-81)	0.683
Male sex	105 (64.4)	673 (61.5)	0.477	103 (64.0)	111 (68.9)	0.345
Tube feeding	6 (3.7)	45 (4.1)	0.794	6 (3.7)	6 (3.7)	1.000
Hospitalization for 2 or more days in the past 90 days	25 (15.3)	207 (18.9)	0.271	25 (15.5)	29 (18.0)	0.551
Residence in a nursing home or long-term care facility	30 (18.4)	139 (12.7)	0.047	30 (18.6)	29 (18.0)	0.885
Recent outpatient intravenous therapy ≤ 30 days	6 (3.7)	39 (3.6)	0.941	6 (3.7)	4 (2.5)	0.551
Attendance at a dialysis center in the previous 30 days	1 (0.6)	24 (2.2)	0.238	1 (0.6)	2 (1.2)	1.000
Comorbidity						
Malignancy ^a	29 (17.8)	197 (18.0)	0.947	28 (17.4)	27 (16.8)	0.882
Chronic liver disease ^b	10 (6.1)	54 (4.9)	0.516	10 (6.2)	11 (6.8)	0.821
Chronic heart disease ^c	23 (14.1)	169 (15.4)	0.658	23 (14.3)	19 (11.8)	0.508
Chronic kidney disease ^d	11 (6.7)	126 (11.5)	0.068	11 (6.8)	16 (9.9)	0.315
Diabetes mellitus ^e	41 (25.2)	251 (22.9)	0.533	40 (24.8)	48 (29.8)	0.317
Chronic respiratory disease ^f	38 (23.3)	295 (27.0)	0.324	38 (23.6)	38 (23.6)	1.000
Central nervous system disorders ^g	49 (30.1)	264 (24.1)	0.102	48 (29.8)	54 (33.5)	0.472
Two or more comorbidities	60 (36.8)	436 (39.9)	0.458	59 (36.6)	66 (41.0)	0.423
Clinical parameters						
Altered mental state ^h	15 (9.2)	100 (9.1)	0.980	15 (9.3)	14 (8.7)	0.846
Respiratory failure ⁱ	58 (35.6)	359 (32.8)	0.484	57 (35.4)	62 (38.5)	0.564
Sepsis or septic shock at onset ^j	22 (13.5)	138 (12.6)	0.752	22 (13.7)	23 (14.3)	0.872
Severe pneumonia	26 (16.0)	165 (15.1)	0.773	26 (16.1)	27 (16.8)	0.881
Intensive care unit admission	21 (12.9)	132 (12.1)	0.745	20 (12.4)	21 (13.0)	0.867
Need for ventilation	8 (4.9)	69 (6.3)	0.487	8 (5.0)	10 (6.2)	0.628
Radiological findings						
Multi-lobar involvement	86 (52.8)	504 (46.1)	0.127	84 (52.2)	83 (51.6)	0.921
Pleural effusion	34 (20.9)	189 (17.3)	0.264	32 (19.9)	31 (19.3)	0.888
Laboratory findings						
White blood cells, /mm ³	11,600 (8,600–16,100)	10,600 (7,675–14,100)	0.011	11,600 (8,500–16,000)	11,800 (8,650–15,850)	0.877
C-reactive protein, mg/dL	12.0 (5.9–18.9)	9.7 (4.2–17.2)	0.020	12.0 (5.8–18.7)	13.3 (6.3–21.7)	0.432
Indices for disease severity						
CURB-65 score	1 (1–2)	1 (1–2)	0.054	1 (1–2)	1 (1–2)	0.737
PSI score	98 (76–123)	92 (70–121)	0.080	98 (76–123)	99 (76–131)	0.610

Values are presented as median (interquartile range) or number (%). Septic shock was defined as sepsis with persisting hypotension requiring vasopressors to maintain a mean arterial pressure of ≥ 65 mmHg and having a serum lactate level of > 2 mmol/L (18 mg/dL) despite adequate volume resuscitation.

PUAT, pneumococcal urinary antigen test; CURB-65, confusion, urea, respiratory rate, blood pressure, age ≥ 65 years; MRSA, methicillin-resistant *Staphylococcus aureus*; PSI, pneumonia severity index.

Table 1. Continued

- ^aMalignancy includes active at the time of presentation or requiring anticancer treatment within the previous 5 years.
- ^bChronic liver disease included preexisting viral or toxic hepatopathy at the time of pneumonia diagnosis.
- ^cChronic heart disease was identified based on past history or administration of diuretics for the treatment of heart disease.
- ^dChronic kidney disease included preexisting renal disease with documented abnormal serum creatinine levels.
- ^eDiabetes mellitus included a history of diagnosis of intolerance to glucose, hemoglobin A1c $\geq 6.5\%$, or treatment with oral hypoglycemic agents or insulin.
- ^fChronic respiratory disease included simple chronic bronchitis, chronic obstructive pulmonary disease, and structural lung diseases such as bronchiectasis and interstitial lung diseases.
- ^gCentral nervous system disorders included acute or chronic vascular or nonvascular encephalopathy with or without dementia.
- ^hAltered mental state was defined as Glasgow Coma Score ≤ 13 documented by the physician.
- ⁱRespiratory failure was defined when partial pressure of oxygen in arterial blood (PaO₂) was 60 mmHg or less or when the PaO₂/fraction of inspired oxygen (FiO₂) ratio was 300 mmHg or less.
- ^jSepsis was defined as organ dysfunction identified as an acute change in the Sequential Organ Failure Assessment score of ≥ 2 consequent to pneumonia.

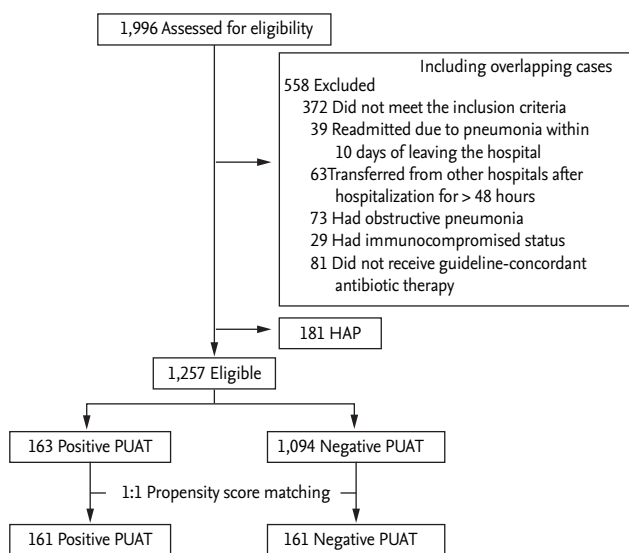


Figure 1. Flow diagram of patient enrollment. HAP, hospital-acquired pneumonia; PUAT, pneumococcal urinary antigen test.

and PUAT was positive in seven of nine (77.8%) patients with definitive pneumococcal pneumonia. Also, in 108 patients with probable pneumococcal pneumonia, 58 (53.7%) tested positive. Taking into account both of the groups together (i.e., all pneumococcal pneumonia), the results of PUAT was positive in 65 patients (56.5%).

Baseline characteristics are presented in Table 1. Prior to matching, patients in the positive PUAT group were more likely to reside in a nursing home or long-term care facility. In addition, the median white blood cell

and C-reactive protein levels were higher in the positive PUAT group than in the negative PUAT group. The propensity score-matching process provided 161 pairs of patients with positive and negative PUAT findings and achieved a good balance for all baseline comorbidities, clinical parameters, and severity indexes.

Diagnostic accuracy of the PUAT in patients with CO-pneumonia

Table 2 shows the calculated values of sensitivity, specificity, positive and negative predictive values, and positive and negative LRs by the different methods. In patients with all pneumococcal pneumonia, the sensitivity of PUAT was 56.5% (95% confidence interval [CI], 47.0 to 65.7) and the specificity was 90.9% (95% CI, 89.0 to 92.6). Additionally, the positive predictive value was 39.9% (95% CI, 34.1 to 46.0) and the negative predictive value was 95.1% (95% CI, 94.1 to 96.0). Positive and negative LRs were 6.2 (95% CI, 4.9 to 8.0) and 0.5 (95% CI, 0.4 to 0.6).

Microorganisms

Table 3 shows the distributions of microorganisms on the basis of the culture results. Overall etiology was established in 97 (59.5%) and 518 (47.3%) patients in the positive PUAT and the negative PUAT groups, respectively. Among patients with positive PUAT results, other microorganisms were identified in 36 (22.1%) patients using conventional methods. In both the full and propensity-matched cohorts, the rate of *S. pneumoniae* identified was significantly higher in the positive PUAT group (39.9% vs. 4.6%, $p < 0.001$ and 39.5% vs. 4.3%, $p <$

Table 2. Diagnostic accuracy of the PUAT in patients with community-onset pneumonia

Diagnosis of pneumococcal pneumonia	Sensitivity	Specificity	Positive PV	Negative PV	Positive LR	Negative LR
Definitive	77.8 (40.0–97.2)	87.3 (85.4–89.1)	4.2 (3.0–6.1)	99.8 (99.4–100)	6.1 (4.2–9.0)	0.3 (0.1–0.9)
Probable	53.7 (43.9–63.3)	90.9 (89.1–92.5)	35.6 (30.0–41.6)	95.4 (94.5–96.2)	5.9 (4.6–7.6)	0.5 (0.4–0.6)
Overall	56.5 (47.0–65.7)	90.9 (89.0–92.6)	39.9 (34.1–46.0)	95.1 (94.1–96.0)	6.2 (4.9–8.0)	0.5 (0.4–0.6)

Values are presented as 95% confidence interval.

PUAT, pneumococcal urinary antigen test; PV, predictive value; LR, likelihood ratio.

Table 3. Microorganisms isolated from the patients on the basis of culture results

Variable	Entire cohort (n = 1,257)			Propensity score-matched cohort (n = 322)		
	Positive PUAT (n = 163)	Negative PUAT (n = 1,094)	p value	Positive PUAT (n = 161)	Negative PUAT (n = 161)	p value
Microorganisms^a						
Identified pathogens	97 (59.5)	518 (47.3)	0.004	95 (59.0)	85 (52.8)	0.262
Gram-positive bacteria						
<i>Streptococcus pneumoniae</i>	65 (39.9)	50 (4.6)	< 0.001	64 (39.8)	7 (4.3)	< 0.001
<i>Staphylococcus aureus</i>	2 (1.2)	43 (3.9)	0.083	2 (1.2)	9 (5.6)	0.032
Methicillin-sensitive <i>S. aureus</i>	0	20 (1.8)	0.096	0	6 (3.7)	0.030
Methicillin-resistant <i>S. aureus</i>	2 (1.2)	23 (2.1)	0.762	2 (1.2)	3 (1.9)	1.000
Other Gram-positive bacteria	0 (0)	17 (1.6)	0.151	0	2 (1.2)	0.498
Gram-negative bacteria						
<i>Pseudomonas aeruginosa</i>	8 (4.9)	46 (4.2)	0.680	8 (5.0)	6 (3.7)	0.585
<i>Haemophilus influenza</i>	1 (0.6)	13 (1.2)	1.000	1 (0.6)	1 (0.6)	0.000
<i>Klebsiella pneumoniae</i>	9 (5.5)	69 (6.3)	0.698	8 (5.0)	11 (6.8)	0.478
ESBL (+)	1 (0.6)	8 (0.7)	1.000	1 (0.6)	1 (0.6)	1.000
ESBL (–)	8 (4.9)	61 (5.6)	0.727	7 (11.5)	10 (6.2)	0.455
<i>Acinetobacter</i> species	0	11 (1.0)	0.377	0	4 (2.5)	0.123
Other Gram-negative species ^b	2 (1.2)	14 (1.3)	1.000	2 (1.2)	1 (0.6)	1.000
<i>Mycoplasma pneumonia</i>	14 (8.6)	61 (5.6)	0.130	14 (8.7)	9 (5.6)	0.279

Values are presented as number (%). Percentages refer to the division by the total number of patients.

PUAT, pneumococcal urinary antigen test; ESBL, extended-spectrum β -lactamase.

^aAllowed for overlapping.

^bOther Gram-negative species included *Escherichia coli*, *Enterobacter* species, *Serratia marcescens*, and *Legionella pneumophila*.

0.001). Following propensity score matching, the rate of *Staphylococcus aureus* was statistically higher in the negative PUAT group (1.2% vs. 5.6%, $p = 0.032$).

Associations between the PUAT and clinical outcomes

The primary and secondary study outcomes are summarized in Table 4. In the full cohort of 1,257 patients,

there were no significant differences in 30-day mortality between groups (6.1% in the positive PUAT group vs. 8.2% in the negative PUAT group; odds ratio [OR], 0.73; 95% CI, 0.37 to 1.43; $p = 0.357$) (Fig. 2A). However, in the propensity-matched cohort, the 30-day mortality rates were lower in the positive PUAT group when compared with the negative PUAT group (5.6% vs. 17.4%; OR, 0.28; 95% CI, 0.13 to 0.62; $p = 0.001$) (Fig. 2B). In subgroup

Table 4. The associations between the PUAT and clinical outcomes before and after propensity score matching

Outcomes	Positive PUAT	Negative PUAT	p value	Effect size
Primary outcome				
30-day mortality, %				
Crude (full cohort)	6.1	8.2	0.357	0.73 (0.37 to 1.43) ^a
Propensity score matching ^b	5.6	17.4	0.001	0.28 (0.13 to 0.62) ^a
Secondary outcomes				
PDR pathogens ^c , %				
Crude (full cohort)	6.7	9.8	0.216	0.67 (0.35 to 1.27) ^a
Propensity score matching ^b	4.3	10.6	0.034	0.39 (0.16 to 0.96) ^a
Duration of antibiotic therapy, day				
Crude (full cohort)	11.1 ± 6.0	11.6 ± 8.2	0.590	-0.50 (-1.54 to 0.54) ^d
Propensity score matching ^b	11.1 ± 6.0	11.9 ± 6.7	0.294	-0.80 (-2.19 to 0.59) ^d
Change of antibiotics, %				
Crude (full cohort)	17.8	23.0	0.134	0.72 (0.47 to 1.11) ^a
Propensity score matching ^b	17.4	26.7	0.044	0.58 (0.34 to 0.99) ^a
Use of inappropriate antibiotics, %				
Crude (full cohort)	6.1	6.7	0.796	0.91 (0.46 to 1.81) ^a
Propensity score matching ^b	5.6	6.2	0.813	0.89 (0.35 to 2.26) ^a
Failure of initial antibiotic therapy, %				
Crude (full cohort)	15.3	21.5	0.071	0.66 (0.42 to 1.04) ^a
Propensity score matching ^b	14.9	30.0	0.004	0.45 (0.26 to 0.79) ^a
Length of hospital stay, day				
Crude (full cohort)	10.8 ± 12.0	9.6 ± 8.8	0.737	1.20 (-0.71 to 3.11) ^d
Propensity score matching ^b	10.9 ± 12.1	12.3 ± 10.5	0.153	-1.40 (-3.87 to 1.07) ^d

Values are presented as percentage or mean ± SD.

PUAT, pneumococcal urinary antigen test; PDR, potentially drug-resistant; CURB-65, confusion, urea, respiratory rate, blood pressure, age ≥ 65 years; ESBL, extended-spectrum β-lactamase.

^aOdds ratio (95% confidence interval).

^bThe variables included as covariates in the propensity score matching were age; sex; tube feeding; comorbidities (e.g., malignancy, chronic liver disease, chronic heart disease, chronic kidney disease, diabetes mellitus, chronic respiratory disease, central nervous system disorders, and two or more comorbidities); altered mental state; respiratory failure; sepsis or septic shock; intensive care unit admission; need for ventilation; antibiotic use before admission; multi-lobar involvement; pleural effusion; white blood cells; C-reactive protein; CURB-65 score; and pneumonia severity index.

^cPDR pathogens included methicillin-resistant *Staphylococcus aureus*, *Pseudomonas* species, *Acinetobacter* species, *Stenotrophomonas maltophilia*, and ESBL-producing Enterobacteriaceae.

^dMean difference (95% confidence interval).

analysis for overall patients with severe pneumonia, the 30-day mortality was not significantly different between the positive and negative PUAT groups (23.1% vs. 33.1%, $p = 0.297$). On the other hand, in the propensity-matched cohort of severe pneumonia, the 30-day mortality rates were lower in the positive PUAT group (23.1% vs. 59.3%, $p = 0.008$).

In the full cohort, there were no significant differences in all secondary study outcomes between groups. In the propensity-matched cohort, the proportion of patients with PDR pathogens was significantly lower in the positive PUAT group (4.3% vs. 10.6%; OR, 0.39; 95% CI, 0.16 to 0.96; $p = 0.034$). Likewise, the proportion of patients with a change of antibiotics and failure of initial

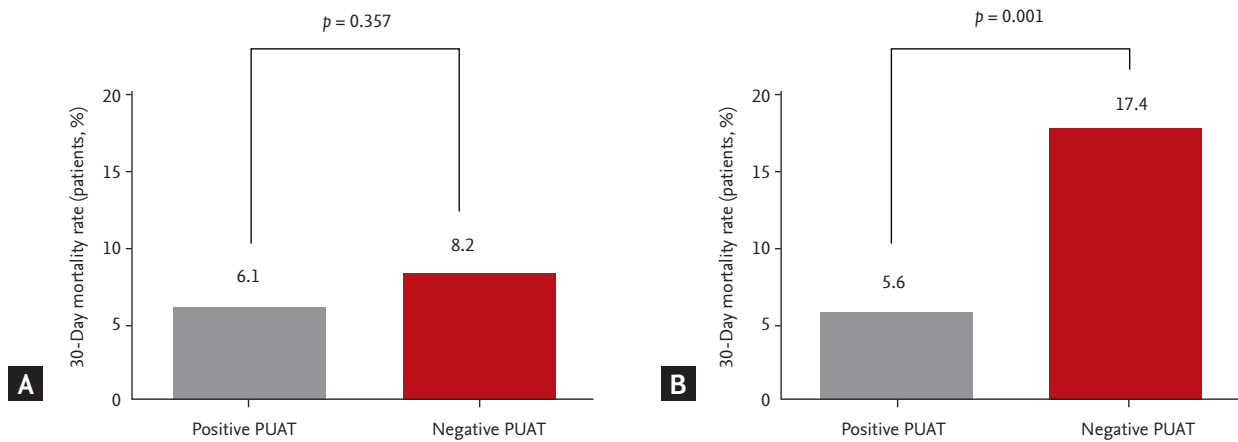


Figure 2. Associations between the pneumococcal urinary antigen test (PUAT) and 30-day mortality rate. (A) The full cohort and (B) the propensity score-matched cohort.

antibiotic therapy was also statistically lower in the positive PUAT group compared to the negative PUAT group ([17.4% vs. 26.7%; OR, 0.58; 95% CI, 0.34 to 0.99; $p = 0.044$] and [14.9% vs. 30.0%; OR, 0.45; 95% CI, 0.26 to 0.79; $p = 0.004$], respectively).

DISCUSSION

In our study, we identified three major findings. First, approximately 13% of patients with CO-pneumonia had positive PUAT results, and the sensitivity and specificity values for the diagnosis of definitive, probable, and overall pneumococcal pneumonia were 77.8% and 87.3%, 53.7%, and 90.9%, and 56.5% and 90.9%, respectively. Second, the clinical parameters of the positive PUAT group were not worse than in the negative PUAT group. Third, in the propensity-matched cohort, the 30-day mortality rates were lower in the positive PUAT group.

In patients with CO-pneumonia, the selection of empirical antibiotic regimen is based on the prediction of the most common pathogens [3,9]. Since the widespread use of pneumococcal conjugate vaccines was initiated, recent studies have reported a decline in *S. pneumoniae* as a cause of CAP [2,19,20]. However, *S. pneumoniae* is still the most common cause at 5% to 21% among microorganisms leading to hospitalization in patients with CAP [7,21-23]. The PUAT is simple, rapid, and useful for detecting pneumococcal pneumonia when samples for culture cannot be obtained in a timely fashion or when

antibiotic therapy has already commenced [3,8]. Previously, a meta-analysis demonstrated that the pooled sensitivity and specificity of the PUAT for definitive pneumococcal pneumonia in the same clinical setting used in the present study were 75% and 80%, respectively [24]. Our results indicated that the diagnostic performances of the PUAT for definitive pneumococcal pneumonia were comparable with the results of a meta-analysis [24].

The PUAT has been reported to have a higher diagnostic yield in patients with severe pneumonia [25]. Although a previous study revealed that the PUAT was more sensitive in patients with high-risk pneumonia as compared to those without (94% vs. 63%, $p < 0.001$) [25], the rate of severe pneumonia in our study was similar between both groups (16.0% vs. 15.1%, $p = 0.773$). And, while a recent multicenter USA study reported that patients with HCAP were less likely to have positive PUAT results than those with CAP [10], there were no differences in positive PUAT results between the CAP and HCAP groups in our study.

The association between PUAT result and clinical outcomes in patients with CO-pneumonia has been investigated in a few studies [26,27]. A large prospective study further demonstrated that positive PUAT results were significantly associated with 30-day mortality, intensive care unit admission, use of mechanical ventilation, treatment failure, and adverse outcomes in patients with bacteremic pneumococcal pneumonia [26]. However, this study contained the possibility of selection bias, and patients with positive PUAT results were

significantly worse in clinical parameters such as respiratory rates, arterial oxygen pressure, and pH levels, and multi-lobar involvement [26]. In the present study, we found that 30-day mortality rates were significantly lower in the positive PUAT group after adjustment for initial presentations using the propensity score-matching process. Similarly, in a recent registry-based retrospective study using a modified CRB65 (confusion, respiratory rate, blood pressure, and age ≥ 65 years) score by subtracting 1 from the scoring system if the PUAT was positive, positive PUAT results were associated with lower risk of 30-day mortality [27]. We additionally investigated the association between the PUAT results and secondary outcomes. In the propensity-matched cohort, the proportion of patients with PDR pathogens, changes in antibiotics, and failure rates of initial antibiotic therapy were significantly higher in the negative PUAT group than in the positive PUAT group.

Reasons for why PUAT is negative in some patients with pneumococcal pneumonia have been proposed to include the following: low levels of C-polysaccharide antigen and sequestration of the antigen by binding to serum antibodies in immune complexes, reduced urinary excretion of the antigen, delayed times from the onset of symptoms to diagnosis, and previous antibiotic treatment [26]. We tried to find clinical parameters of false negative PUAT results. Compared to patients with true positive PUAT results, those with false negative PUAT results had chronic heart disease less frequently (18.5% vs. 8.0%, $p = 0.049$). However, because of the small sample size ($n = 165$) and low statistical power ($p = 0.049$), we could not allow for this as a crucial parameter.

There remains controversy with regard to the change of antibiotics on the basis of the results of PUAT [28,29]. In a prospective study including 219 patients with non-severe CAP, the targeted use of amoxicillin based on PUAT results was able to decrease antibiotic resistance and reduce the use of unnecessary antibiotics [28]. In contrast, another randomized controlled trial reported that *S. pneumoniae*-targeted therapy with positive PUAT results increased the rate of clinical relapse compared with empirical therapy [29]. Possible explanations for treatment failure may include the following: first, the possibility of polymicrobial infections cannot be excluded [29]. Especially in the case of polymicrobial infections with PDR pathogens, narrowing the anti-

biotic treatment can lead to treatment failure. Second, a high proportion of pathogens remain unidentified in patients with CO-pneumonia. This means that it is difficult to predict the rate of polymicrobial infections. Third, pneumococcal capsular polysaccharides could react with other microorganisms such as *Staphylococcus*, *Streptococcus* species, and Gram-negative strains [30]. Other microorganisms identified in patients with positive PUAT results through conventional methods, might be regarded as having cross-reactivity [30]. Possible cross-reacting microorganisms could elicit treatment failure [30]. Finally, the presence of drug-resistant *S. pneumoniae* may also be a plausible explanation [28]. The emergence and spread of drug-resistant *S. pneumoniae* have been reported in recent years, and antibiotic resistance threatens the successful management of pneumococcal pneumonia [31]. In a meta-analysis encompassing 10 studies that involved 3,430 patients, the combined relative risks of all-cause mortality for the penicillin-resistant *S. pneumoniae* groups was 1.29 (95% CI, 1.01 to 1.66) when compared with the penicillin-susceptible *S. pneumoniae* group [32].

In principle, patients with positive PUAT results would have a good response to empirical guideline-concordant antibiotic therapy, which has been associated with a significant decrease in in-hospital mortality, length of hospital stay, and the duration of parenteral therapy [33]. We maintained guideline-concordant empirical antibiotics without regard to the results of tests and investigated the association between the PUAT results and clinical outcomes in the present study.

The strength of our study is in the propensity score-matching process, although our subjects were a retrospective cohort. Meanwhile, there are some study limitations. First, because our study was conducted retrospectively involving patients admitted to a single institution, our data should be interpreted with caution. In particular, even though pneumococcal vaccination might produce a false positive result in the PUAT in clinical practice [34], we could not identify the rate of pneumococcal vaccination due to the retrospective nature of the study. Second, we did not perform viral testing in most of the included patients. Mixed infections consisted of approximately 5% to 10% of the etiology for CAP in previous studies [20,27]. Most polymicrobial agents in these studies were viral pathogens, which do not usually require specific anti-

microbial therapy [20,35]. Because we did not perform routine viral testing, the role of viral agents in CO-pneumonia might be underestimated in clinical outcomes. Third, previous studies have reported the relationship between serotype of *S. pneumoniae* and disease outcome [36]. A pooled estimates for invasive pneumococcal disease demonstrated that serotypes 1, 7F, and 8 were associated with decreased risk ratios, while serotypes 3, 6A, 6B, 9N, and 19F were associated with increased risk ratios [36]. However, the serotype of *S. pneumoniae* was not investigated in this clinical setting.

In conclusion, our study revealed that the PUAT had low sensitivity and high specificity for overall pneumococcal pneumonia. After baseline characteristics were adjusted through the propensity score-matching process, empirical guideline-concordant antibiotic therapy in CO-pneumonia was associated with a lower rate of 30-day mortality in patients with positive PUAT results.

KEY MESSAGE

1. The pneumococcal urinary antigen test (PUAT) had low sensitivity and high specificity for overall pneumococcal pneumonia.
2. Baseline clinical parameters were similar between the positive PUAT and negative PUAT groups.
3. In the propensity-matched cohort, the 30-day mortality rates were lower in the positive PUAT group.

Conflict of interest

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

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