

Supplementary Material

Extended Methods

Study Population

The study was conducted in six medical and surgical ICUs, overall comprising 45 beds, at an 800-bed university hospital. Data were prospectively collected from 2007 to 2017. Investigators made daily rounds of each ICU. Patients older than 18 years, mechanically-ventilated for 48 hours or more, with clinical suspicion of VAP were consecutively included in the study and only the first episode was analyzed. We excluded patients with severe immune-suppression (neutropenia after chemotherapy or hematopoietic transplant, drug-induced immune-suppression in solid-organ transplant or cytotoxic therapy, and human immunodeficiency virus infected patients) [1]. The institution's Internal Review Board approved the study (*Comitè Ètic d'Investigació Clínica*, registry number 2009/5427) and written informed consent was obtained from patients or their next-of-kin.

Definition of Pneumonia, Microbiologic Processing, and Antimicrobial Treatment

The clinical suspicion of pneumonia was based on clinical criteria (new or progressive radiological pulmonary infiltrate together with at least two of the following: (1) temperature $>38^{\circ}\text{C}$ or $<36^{\circ}\text{C}$; (2) leukocytosis $>12,000/\text{mm}^3$ or leucopenia $<4,000/\text{mm}^3$; or (3) purulent respiratory secretions) [2,3].

The microbiologic evaluation included the collection of at least one lower respiratory airways sample: tracheobronchial aspirates (TBAS) and/or bronchoscopic [4] or blind bronchoalveolar lavage (BAL) [5], if possible within the first 24 hours of inclusion [6]. Only samples of tracheal aspirates of high quality (i.e. <10 squamous cells and >25 leukocytes per optical microscopy field) were processed for culture. The same sampling method was performed on the third day if clinically indicated. Blood cultures and cultures from the pleural fluid were also collected if clinically justified. Urinary antigens of *Streptococcus pneumoniae* and *Legionella pneumophila* were not systematically collected.

Microbiologic confirmation of pneumonia was defined by the presence of at least one PPM, i.e. endogenous microorganisms associated with nosocomial respiratory infection in the host [7], in the respiratory samples with above pre-defined values (BAL $> 10^4$ and TBAS $> 10^5$ colony-forming units/ml, respectively, or any value if the patient was under antibiotic treatment), in pleural fluid or in blood cultures if an alternative cause of bacteremia was ruled out [8,9]. Microbiologic

identification and susceptibility testing were performed by standard methods [10]. Polymicrobial pneumonia was defined when more than one PPM were identified as causative agents.

The initial empiric antimicrobial treatment was administered to all patients according to local adaptation of American Thoracic Society (ATS)/Infectious Disease Society of America (IDSA) guidelines [11,12], based on the most frequently isolated PPM and their patterns of antimicrobial sensitivity in our institution, and subsequently revised according to the microbiologic results.

The empiric antimicrobial treatment was considered appropriate when the isolated pathogens were susceptible in vitro to at least one of the antimicrobials administered at adequate dose [13].

The initial response to treatment was evaluated after 72 hours of antimicrobial treatment. Non-response to treatment was defined by the presence of at least one of the following criteria: No improvement of the arterial PaO₂/FiO₂; need for intubation because of pneumonia (defined as need for intubation after 24 h from the beginning of antibiotics); persistence of fever (temperature ≥ 38 °C) or hypothermia (<35.5 °C) together with purulent respiratory secretions; increase in pulmonary infiltrates on chest radiograph ≥ 50 %; occurrence of septic shock or multiple organ dysfunction syndrome, defined as three or more organ system failures not present on day one. In patients with initial non-response to treatment [14,15], respiratory samples and blood cultures were obtained again, and the empiric antimicrobial treatment was revised.

Assessment of the Systemic Inflammatory Response

We evaluated the serum levels of interleukin (IL)-6, IL-8, IL-10, tumor necrosis factor-alpha (TNF-alpha), C-reactive protein (CRP) and Procalcitonin (PCT) within the first 24 hours and the third day after the diagnosis of pneumonia. We also determined mid-regional pro-adrenomedullin (MR-proADM) using a test based on the time-resolved amplified cryptate emission technology (KRYPTOR; BRAHMS AG; Hennigsdorf, Germany). MR-proADM has an analytical detection limit of 0.08 nmol/L (normal reference range 0.33 ± 0.7 nmol/L) and a functional assay sensitivity of 0.12 nmol/L. All methods for this analysis have been described in detail elsewhere [16,17].

Data Collection

All relevant data were collected at admission and at onset of pneumonia from the medical records and bedside flow charts, including laboratory, radiologic and microbiologic information. We calculated the Acute Physiology And Chronic Health Evaluation (APACHE-II) [18], the Simplified Acute Physiology Score (SAPS) [19], and the Sepsis-related Organ Failure Assessment (SOFA) [20] scores. Patients were followed until hospital discharge, death or up to 90 days after

the diagnosis of pneumonia. Septic shock [21] and acute respiratory distress syndrome (ARDS) [22] were defined according to the previously described criteria.

Outcomes Variables

The rate of microbiological confirmation of patients with $\text{PaO}_2/\text{FiO}_2 \leq 240$ was compared with that of patients with $\text{PaO}_2/\text{FiO}_2 > 240$ mmHg. The lowest value of $\text{PaO}_2/\text{FiO}_2$ on the day of VAP diagnosis was considered for patients' group allocation. Secondary outcomes were ICU and hospital length of stay and mortality, and survival at 90 days after VAP diagnosis.

Statistical analysis

Categorical and continuous data are presented as number (percentage) and as mean \pm SD (or median, inter-quartile range), respectively. Categorical variables were compared with the Chi-square or Fisher's exact tests. Quantitative continuous variables were compared using the unpaired Student's t-test or the Mann-Whitney test for normally and not-normally distributed variables, respectively. The Kaplan-Meier curves were used to compare survival in the two groups.

Univariate and multivariate logistic regression analyses were performed to assess the association of $\text{PaO}_2/\text{FiO}_2 \leq 240$ or > 240 with positive microbiology. Variables that showed a significant result univariately ($p < 0.10$) were included in the corresponding multivariate logistic regression backward stepwise model. Adjusted odds-ratio (OR) and 95% confidence intervals (CI) were calculated.

In addition, the association between $\text{PaO}_2/\text{FiO}_2$ at VAP onset as continuous variable and quantitative microbiological confirmation was assessed using receiver-operator-characteristics (ROC) curve analysis. The area under the ROC curve (AUC) with 95% confidence interval (CI), and optimal cut-off value of $\text{PaO}_2/\text{FiO}_2$ with sensitivity and specificity were calculated.

A two-sided p -value ≤ 0.05 was considered statistically significant. All statistical analyses were performed using SPSS 22.0.0.0 (Chicago, IL).

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