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# A Pilot Study on the Diagnostic Accuracy of Proadrenomedullin and Proatrial Natriuretic Peptide in Lower Respiratory Tract Infections



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**Abstract:** *Background*: Pneumonia is the leading cause of death among infectious diseases in developed countries. However, the severity of pneumonia requiring hospitalization often makes the initial diagnosis difficult because of an equivocal clinical picture or interpretation of the chest film. The objective of the present study was to assess the usefulness of the plasma levels of mid-regional proadrenomedullin (MR-proADM) and mid-regional proatrial natriuretic peptide (MR-proANP) in differentiating pneumonia from other lower respiratory tract infections (LRTIs).

*Methods*: A retrospective study was conducted. The plasma levels of MR-proADM and MR-proANP were measured in 85 patients hospitalized for LRTIs, 56 of whom with diagnosis of pneumonia and 29 with other LRTIs.

*Results*: The patients with pneumonia had increased MR-proADM levels (median 1.46 nmol/L [IQR 25-75, 0.82-2.02 nmol/L]) compared with the patients with other LRTIs (median 0.88 nmol/mL [0.71-1.39 nmol/L]) (p= 0.04). However, the MR-proANP levels did not show differences between the groups. The optimal threshold of MR-proADM to predict pneumonia was 1.5 nmol/L, which yielded a sensitivity of 51.7% (95% CI, 38.0-65.3), a 79.3% specificity (95% CI, 60.3-92.0), and an odds ratio of 6.64 (95% CI, 1.32-32.85). The combination of this parameter with C-reactive protein in an "and" rule increased the specificity for detecting pneumonia significantly.

Conclusion: MR-proADM levels (but not MR-proANP levels) are increased in patients with pneumonia although its discriminatory power is moderate.

Keywords: Adrenomedullin, atrial natriuretic peptide, biomarkers, diagnosis, pneumonia.

## **INTRODUCTION**

Pneumonia is the sixth leading cause of death in general and the first among all infectious diseases in developed countries [1-3]. However, discriminating pneumonia from other chest infections where antibiotics are not required (i.e. acute bronchitis), particularly in its early stages, is sometimes difficult. In older patients, the clinical presentation is unclear, and interpreting chest radiographs can be difficult in patients with severe disease or previous pulmonary disease [4, 5].

The search for biomarkers to serve as diagnostic tools for clinicians has not yet produced a reliable tool. One biomarker of interest is mid-regional proadrenomedullin (MR-proADM), a more stable mid-region fragment of the parent precursor adrenomedullin [6]. Adrenomedullin is a peptide that is produced by multiple tissues types; it has pluripotent functions, including vasodilatory, antimicrobial, and anti-inflammatory activities [7]. In animal models of sepsis, exogenous adrenomedullin reduced acute lung injury, vascular permeability, and death [8-10]. On the other hand, mid-regional proatrial natriuretic peptide (MR-proANP) is a more stable mid-region fragment precursor of atrial natriuretic peptide, and the stimuli for its production most likely include proinflammatory factors and hypoxia. Its expression has been found in many tissues, but it is mainly produced in the atrium of the heart [11-12]. In previous studies, MR-proADM and MR-proANP levels have correlated with severity of pneumonia and death [13-17]. However, data on the diagnostic usefulness of these biomarkers is lacking in the literature.

Therefore, we examined the MR-proADM and MRproANP levels in a cohort of patients with lower respiratory tract infections (LRTIs) requiring hospitalization. We sought to describe the pattern of the initial biomarkers in pneumonia and other LRTIs, and to assess the potential diagnostic role in pneumonia.

#### MATERIALS AND METHODS

#### **Study Sample**

Since 2010, consecutive patients over age 18 with LRTIs requiring hospital admission were asked to provide

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additional blood samples for investigation. The study protocol was approved by the local ethical committee, and all of the participants provided written informed consent for the collection and storage of blood upon admission. Plasma was obtained by centrifugation and stored frozen at -80°C. For the present study, 85 plasma samples were randomly selected.

#### **Diagnostic Criteria**

LRTI was defined by the presence of at least one respiratory symptom (e.g., cough, sputum production, dyspnea, tachypnea, pleuritic pain) plus at least one finding during auscultation (i.e., rales) or one sign of infection (temperature >38.0°C, shivering, leukocyte count >10 or <4  $x10^9$  cells) regardless of antibiotic use. For CAP, a new infiltrate on the chest radiograph was also required. COPD was defined by post-bronchodilator spirometric criteria according to the Global initiative for Chronic Obstructive Lung Disease (GOLD)-guidelines as a FEV1/FVC ratio < 70%. Acute bronchitis was defined as a LRTI in the absence of an underlying lung disease or focal chest signs and infiltrates on chest X-rays [18, 19]. A panel of three specialists had been previously reviewed chest X-rays from the patient population. Panel consensus in classifying LRTI was required for study inclusion.

The exclusion criteria were: 1) severe immunosuppression (HIV infection or severe haematological diseases); 2) immunosuppressive therapy (prednisone or an equivalent dose of 20 mg daily for 2 weeks or any immunosuppressive regimen (azathioprine, cyclosporine, cyclophosphamide and/or other immunosuppressant drugs); 3) leukopenia (<1,000 leukocytes per mm<sup>3</sup>) or neutropenia (< 500 neutrophils per mm<sup>3</sup>) and/or chemotherapy in the previous year; 4) pulmonary abscess (radiological cavitation), aspiration pneumonia and obstructive pneumonia; 5) possible nosocomial origin (30 days from hospital discharge); and 6) known active neoplasm.

#### **Data and Severity Assessment**

The patients' clinical parameters were included in an electronic database. Comorbidities were documented and defined as the presence of one or more of the following: congestive heart failure, chronic obstructive pulmonary disease, chronic renal disease, chronic liver disease, cerebrovascular disease, malignancy, or diabetes mellitus. The microbiological work-up in the patients with pneumonia included sputum and/or bronchoalveolar lavage fluid samples for the Gram stain and culture, a urine sample for detecting Legionella pneumophila and Streptococcus pneumoniae antigens, and serological testing for Mycoplasma pneumoniae and Chlamydophila pneumoniae. The detection of respiratory pathogens was performed according to standard methods. In patients with severe exacerbations of COPD, a sputum sample was collected for the Gram stain and sputum culture. No microbiological tests were performed in the patients with acute bronchitis.

The severity assessment was made with the CURB-65 score and performed upon admission to the emergency department. The CURB-65 score consists of five variables: confusion, urea, respiratory rate  $\geq 30$  breaths/min, blood pressure (systolic) < 90 mm Hg or diastolic  $\leq 60$  mmHg, and age  $\geq 65$  years [20]. One point is given for each parameter

present, which results in CURB-65 scores of 0-5. The criteria for ICU admission were based on the American Thoracic Society (ATS) guidelines [21]. In brief, ICU admission was considered in patients with severe CAP, which was defined as the presence of one of two major criteria (the need for mechanical ventilation or septic shock), the presence of two of three minor criteria (systolic blood pressure <90 mmHg, multilobar disease, or PaO2/FIO2ratio <250) or more than two CURB points. For COPD patients, the ICU criteria included severe acidosis or respiratory failure (pH <7.25, pO2 < 40 mmHg), no response to initial treatment in the emergency department or worsening mental status (confusion, coma) despite adequate therapy [19].

#### Laboratory Measurements

Both MR-proADM and MR-proANP were measured in duplicated in stored EDTA plasma samples by the TRACE® technology, as previously described [22, 23]. The procedure was fully automated in a KRYPTOR Compact analyzer (B.R.A.H.M.S. AG, Hennigsdorf, Germany). The other biochemical determinations were routinely performed as part of the admission study using conventional laboratory methods.

#### **Statistical Analysis**

The categorical variables were expressed as counts (percentages), and the continuous variables were expressed as medians and interquartile ranges (IQR), as appropriate. The  $\chi 2$  and Fisher's exact tests were used, as appropriate, to compare the qualitative variables. In the case of quantitative variables, the Mann-Whitney U-test was used.

To assess the accuracy of the biomarkers for predicting the presence of pneumonia, we generated receiver operating characteristic curves and determined the area under the curve (AUC) and its 95% confidence interval (95% CI). For each ROC curve, a cut-off point was determined as the value of the biomarker that maximised the sum of the sensitivity and specificity for diagnosing pneumonia. Pairwise comparisons of the AUC were conducted, following the Hanley and McNeil procedure. To adjust for confounders, a logistic regression model estimated the simultaneous impact of CRP and MR-proADM, along with significant variables in the univariate analysis predicting pneumonia. Associations were considered to be statistically significant if the p-value was <0.05. The statistical software packages used were SPSS 15.0 (SPSS Inc., Chicago, IL, USA) and MedCalc version 12.3.0 (MedCalc, Mariakerke, Belgium).

### RESULTS

#### **Baseline Characteristics**

A total of 85 patients who were hospitalized for LRTIs were recruited; 56 were diagnosed with pneumonia and 29 with other LRTIs (13 with acute exacerbation of chronic obstructive pulmonary disease and 16 with acute bronchitis). The mean patient age was 71.0 years (range 20-96); 69.4% were males, and 62.4% had comorbidities. Of the participants, 8.2% arrived from nursing home facilities. The patients with pneumonia had a median CURB65 score of 2.5. During the follow-up, 3 (3.5%) patients were transferred to the ICU, and 7 (8.2%) died (6 with pneumonia and 1 with other LRTIs). The mean in-patient stay was 8.4 days (range 2-41).

Microbiological work-up were performed in patients with pneumonia or with severe exacerbations of COPD. A final etiological diagnosis was established in 29 (42.0%) patients. The microorganisms identified were: *Streptococcus pneumoniae* (19), *Chlamydophila pneumoniae* (6), *Legionella pneumophila, Staphylococcus aureus, Escherichia coli*, and *Aspergillus fumigatus* (1 each).

#### **Data Analysis**

A comparison of the clinical, analytical and follow-up data between the groups is shown in Table 1. In a univariate analysis, 5 variables were different between the patients with pneumonia and those with other LRTIs: acute onset of symptoms (48% vs 14%), heart rate (100 vs 89 beats/min), the presence of rales (64% vs 14%), CRP (222 vs 102 mg/L) and MR-proADM (1.46 vs 0.88 nmol/L). However, in the multivariate analysis, the presence of rales, CRP and MR-proADM maintained significant differences only.

Fig. (1) shows MR-proANP levels in the study population. The patients with pneumonia had similar MR-proANP levels (median 233 pmol/L [IQR 25-75, 115-398 pmol/L]) compared with those with other LRTIs (median

The AUC for MR-proADM to discriminate between the patients with pneumonia and other LRTIs was 0.65 (95% CI, 0.53-0.76), which was not inferior to the result obtained for CRP (0.78 [95% CI, 0.67-0,86], p=0.11). The optimal threshold (minimal false negative and false positive results) of MR-proADM to predict pneumonia was 1.5 nmol/L, which yielded a sensitivity of 51.7% (95% CI, 38.0-65.3), a 79.3% specificity (95% CI, 60.3-92.0). As Table **2** shows, combining MR-proADM with CRP in an "and" rule increased the specificity to 96% (95% CI, 82-99) and the likelihood ratio positive to 9.2 (95% CI, 1.3-65.1) for detecting pneumonia.

#### DISCUSSION

In the present study, MR-proADM levels but not MRpro-ANP levels were found to be increased in patients with pneumonia compared with other LRTIs.

Table 1.	Comparison between	the patients with p	neumonia and those	with other lower	respiratory trac	t infections (LRTIs).

	Pneumonia N=56	Other LRTIs N=29	p Value	p Value*	
Clinical findings					
Days of illness	3 (2-5)	5 (3-7)	0.14		
Previous antibiotic treatment	14 (25)	11 (38)	0.21		
Acute onset	27 (48)	4 (14)	0.00	0.39	
Chills	24 (43)	7 (24)	0.08		
Cough	46 (82)	24 (82)	0.94		
Sputum	26 (46)	14 (48)	0.87		
Pleuritic chest pain	20(36)	6 (21)	0.15		
Physical findings					
Confusion	10 (18)	1(3)	0.05		
Body temperature (°C)	37.3 (36.4-38)	37 (36.6-37.7)	0.27		
Systolic blood pressure (mmHg)	118 (91-139)	127 (108-142)	0.20		
Heart rate (beats/min)	100 (80-111)	89 (77-101)	0.03	0.08	
Respiratory rate (breaths/min)	28 (20-36)	24 (24-29)	0.97		
Rales	36 (64)	4 (14)	0.00	0.01	
Laboratory					
WCC (white cell count) (x10 <sup>9</sup> )	12.3 (7.7-17.3)	11.5 (7.7-14.4)	0.37		
Haematocrit %	aematocrit % 41 (36-45)		0.89		
Urea mg/dL	56 (40-81)	37 (28-76)	0.05		
Sodium mmol/L	138 (133-140)	138 (136-140)	0.77		
pO <sub>2</sub> (mm Hg)	62 (54-79)	64 (53-75)	0.86		
C-reactive protein (mg/L)	222 (149-278)	102 (59-152)	0.00	0.01	
MR-proADM (nmol/L)	1.46 (0.82-2.02)	0.88 (0.71-1.39)	0.04	0.03	
MR-proANP (pmol/L) 233 (115-398)		259 (110-399) 0.88			

The data are expressed as the median (IQR) or N (%). \* Multivariate analysis.



**Fig. (1).** Box plots of the MR-proANP plasma levels in the study population. The patients with pneumonia had similar MR-proANP levels (median 233 pmol/L [IQR 25-75, 115-398 pmol/L]) compared with those with other LRTIs (median 259 pmol/L [110-399 pmol/L]) (p=0.88). The Box plots represent the  $25^{\text{th}}$  and  $75^{\text{th}}$  percentiles, with the internal horizontal lines showing the median. T-shaped bars represent the  $10^{\text{th}}$  and  $90^{\text{th}}$  percentiles. IQR, interquartile range; MR-proANP, mid-regional proatrial natriuretic peptide.



**Fig. (2).** Box plots of the MR-proADM plasma levels in the study population. The patients with pneumonia had increased MR-proADM levels (median 1.46 nmol/L [IQR 25-75, 0.82-2.02 nmol/mL]) compared with those with other LRTIs (median 0.88 nmol/L [0.71-1.39 nmol/L]) (p=0.04). The MR-proADM units are expressed as natural logarithm. The box plots represent the 25<sup>th</sup> and 75<sup>th</sup> percentiles, with the internal horizontal lines showing the median. T-shaped bars represent the 10<sup>th</sup> and 90<sup>th</sup> percentiles. IQR, interquartile range; MR-proADM, mid-regional proadrenomedullin.

The diagnosis of pneumonia is sometimes difficult in clinical practice. In older patients, the clinical presentation is unclear, and the accuracy of chest radiography, which remains the reference technique for diagnosing pneumonia, is 65% compared with a CT scan [24]. In addition, the search of biomarkers that could help in managing infections has produced conflicting results [25].

 Table 2.
 MR-proADM
 characteristics
 alone
 or
 combined

 with C-reactive protein to predict pneumonia.

Serum Markers	Sensitivity	Specificity	LR+	LR-
MR-proADM	51.7	79.3	2.5	0.6
>1.5 nmol/L	(38.0 - 65.3)	(60.3 - 92.0)	(1.2 - 5.3)	(0.4 - 0.8)
CRP	75.4	77.2	3.3	0.3
>150 mg/L	(61.7 - 86.2)	(54.6 - 92.2)	(1.5 - 7.3)	(0.2 - 0.5)
MR-proADM >1.5 nmol/L and CRP >150 mg/L	34.0 (22.7 – 47.4)	96.3 (81.7 – 99.3)	9.2 (1.3 - 65.1)	0.6 (0.5 - 0.8)

The data are presented as % (95% CI). LR+: likelihood ratio positive; LR-: likelihood ratio negative.

In a study of 151 patients with systemic inflammatory response syndrome, the AUC for detecting a bacterial cause was 0.72 for procalcitonin (PCT), 0.74 for neutrophil count, and 0.81 for CRP [26]. In a study of 545 patients with LRTIs (373 with CAP), the AUCs of a clinical model, PCT and CRP were 0.79, 0.76 and 0.88, respectively, to detect patients with pneumonia [27].

Both, MR-proANP and MR-proADM have been recently incorporated into the race of biomarkers primarily by adding prognostic information. Thus, in a study of 300 patients with pneumonia, the MR-proANP levels were significantly higher in the patients with high PSI risk class (IV-V) than in those with low PSI risk class (I-III) [13]. In a multicentre prospective study of 1,653 patients with pneumonia, MRproADM had a higher AUC for 30-day mortality than PCT (0.76 vs 0.65, respectively) [14]. In the multicentre ProHOSP study [15], the authors compared the prognostic accuracy of several biomarkers with the PSI and CURB65 clinical scores in 1.359 patients with LRTIs. The discriminatory power of biomarkers for predicting death ranged from an AUC of 0.60 for PCT to 0.76 for MR-proADM and 0.79 for MR-proANP. The CURB65 and PSI score had an AUC of 0.74 and 0.84, respectively. In the multicentre CAPNETZ study [16], the authors compared the prognostic accuracy of several biomarkers with the PSI and CRB65 clinical scores in 728 patients with CAP. The AUC of MR-proADM for 28-day mortality (0.85) was superior to those of MR-proANP (0.81)and CRB-65 (0.72) for predicting mortality. Recently, MRproADM has shown a better discriminatory power than PSI score in identifying risk of death among CAP patients with PSI class IV and V (0.81 vs 0.66, respectively) [28]. In addition, the utility of MR-proANP or MR-proADM have also been analysed to detecting the aetiology, the presence of bacteraemia, the risk of developing complications, or the need for ICU admission in patients with LRTIs [17, 29-31].

To the best of our knowledge, this is the first study analysing the diagnostic performance of MR-proANP and MR-proADM in an LRTI population. Our results supported the notion that high MR-proADM levels are found in patients with pneumonia compared with other LRTIs. At cutoff levels of 1.5 nmol/L, the discriminatory power of MRproADM was moderate (sensitivity and specificity of approximately 52% and 80%, respectively). However, the combination of MR-proADM with CRP (at cut-off levels of 150 mg/L) in an "or/and" rule, significantly improved the operating characteristics. Indeed, the presence of any of both biomarkers showed a sensitivity of 86%, whereas the presence of both biomarkers increased the specificity to 96% and the likelihood ratio positive to 9.2. By this rule, approximately 15% of patients with pneumonia were missed. However, the levels of MR-proANP did not differ between the groups. The fact that MR-proADM seems to be superior to MR-proANP might be explained by the multiple functions of adrenomedullin. biological Indeed. adrenomedullin is one of the most potent vasodilation agents with immune modulating, metabolic and bactericidal properties. In contrast, natriuretic peptides such as ANP correlate with the presence of heart of renal failure, which may be present only in the subgroup of patients with LRTIs and poor prognoses [6-12].

The present study has some limitations. First, the design was retrospective, and the effects of sample storage on the measurements have not been evaluated. Second, the population evaluated included adult patients with LRTIs who required hospital care; therefore, it may not be possible to extrapolate our conclusions to populations with mild LRTIs. Finally, immunosuppressed patients were excluded from the analysis, and this patient group has difficult diagnoses that could benefit from the use of biomarkers.

In our opinion, further studies with larger samples are needed to analyze the usefulness of MR-proADM testing alone or in combination with other biomarkers in the management of respiratory infections.

#### CONCLUSION

Among hospitalized patients with LRTIs, MR-proADM levels are increased in patients with pneumonia. The optimal threshold of MR-proADM to predict pneumonia was 1.5 nmol/L, which yielded a specificity of 79.3%. MR-proANP levels did not show differences between groups.

#### **ABBREVIATIONS**

CAP	=	Community-acquired pneumonia
LRTI	=	Lower respiratory tract infection
MR-proADM	=	Mid-regional proadrenomedullin
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MR-proANP = Mid-regional proatrial natriuretic peptide

#### **CONFLICT OF INTEREST**

The authors state that have no conflict of interest including any financial, personal or other relationships with other people or organizations with the submitted work that could inappropriately influence, or be perceived to influence, their work.

#### ACKNOWLEDGEMENTS

Declared none.

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Revised: June 12, 2014

Accepted: June 25, 2014

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Received: October 11, 2013