



# Clinical and Bacteriological Relevance of Procalcitonin: A Single Center, Retrospective Observational Study

Romya Singh <sup>a++</sup>, Chinmoy Sahu <sup>a##</sup>, Sangram Singh Patel <sup>a†</sup>,  
Nidhi Tejan <sup>a‡</sup> and Mohan Gurjar <sup>b^</sup>

<sup>a</sup> Department of Microbiology, Sanjay Gandhi Postgraduate Institute of Medical Sciences, Rae Bareilly Road, Lucknow, Uttar Pradesh 226014, India.

<sup>b</sup> Department of Critical Care Medicine, Sanjay Gandhi Postgraduate Institute of Medical Sciences, Rae Bareilly Road, Lucknow, Uttar Pradesh 226014, India.

## Authors' contributions

This work was carried out in collaboration among all authors. Author RS did the data acquisition, draft manuscript. Author CS did the conceptualization, manuscript editing and review. Author SSP did the validation and resources. Author NT did the formal analysis. Author MG did the manuscript review and Supervision. All authors read and approved the final manuscript.

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## ABSTRACT

**Introduction:** Clinical relevance of procalcitonin levels in cases of sepsis due to different pathogens and the relationship between procalcitonin levels and patient outcome has not been widely studied. The aim of this study is to highlight the clinical relevance of procalcitonin in sepsis due to various pathogens and in patient prognosis.

**Methods:** In this retrospective observational study 348 cases of sepsis were analysed and their procalcitonin levels were compared with the different pathogens isolated. The patient outcome as

<sup>++</sup> Senior Resident;

<sup>#</sup> Additional Professor;

<sup>†</sup> Associate Professor;

<sup>‡</sup> Assistant Professor;

<sup>^</sup> Professor;

\*Corresponding author: E-mail: sahu.chinmoy@gmail.com;

28 day mortality was also compared with different procalcitonin levels which was divided into four groups (group1: <0.5ng/ml, group 2: 0.5 - < 2ng/ml, group 3: 2 - < 10ng/ml, group 4: >= 10ng/ml). **Results:** The procalcitonin levels were significantly higher in cases of sepsis due to Gram negative bacilli (14.5ng/ml  $\pm$  2.8) compared to Gram positive cocci (8.59ng/ml  $\pm$  1.5) and yeast (2.96ng/ml  $\pm$  0.56). Multiple logistic regression showed significant difference between 28-day mortality and Multidrug resistant bacteria (MDR) pathogens ( $p=0.006$ ) and group 4 procalcitonin (PCT) levels ( $p=0.033$ ). **Conclusion:** The procalcitonin levels were significantly higher in sepsis due to Gram negative bacilli compared to Gram positive cocci, Gram positive bacilli and yeast. The patient clinical outcome observed as 28-day mortality was also higher in group 4 PCT levels ( $\geq 10$ ng/ml). Thus, we found PCT is a reliable marker for sepsis with Gram negative bacilli and for patient prognosis.

**Keywords:** Procalcitonin; sepsis; pathogens; marker; 28 day mortality.

## 1. INTRODUCTION

Procalcitonin is a prohormone of calcium homeostasis hormone calcitonin, it is produced in the neuroendocrine medullary C cells of thyroid gland at a very low concentration of <0.05ng/ml [1]. Bacterial infections selectively induce PCT production from multiple parenchymal tissues including liver, kidney, lung, intestine and fat tissues [2]. This results in accumulation of PCT because unlike neuroendocrine cells, parenchymal cells lack the ability to cleave PCT into mature form calcitonin [1]. In case of bacterial sepsis the bacterial endotoxin and other inflammatory cytokines induces the release of PCT [1]. There is production of different proinflammatory cytokines by different Toll like receptor signalling pathway by Gram negative bacteria, Gram positive bacteria or fungi which activates PCT [3]. Procalcitonin is a helpful marker in differentiating bacterial and viral infections [1] and also cases of true bacteremia and blood samples with contamination [4,5]. This may be a reason for different levels of PCT in different pathogens. PCT is more specific diagnostic biomarker for bacterial infections compared to CRP [6]. There are several studies which have reported the relevance of PCT in differentiating Gram positive, Gram negative bacteremia and fungemia [7,8]. The PCT levels in different pathogens and its correlation with severity and patient outcome is still not clear. This study evaluates clinical relevance of PCT levels in relation to sepsis due to various pathogens and its importance as a prognostic marker.

## 2. MATERIALS AND METHODS

### 2.1 Study Design and Population

It was a retrospective study conducted in a tertiary care hospital. This study analysed data

from positive blood samples showing growth of bacteria and/or fungus, during a period of one year from April 2021 to May 2022. This study was a part of a project and ethical clearance was taken from the institute ethics committee (PGI/DIR/RC/917/2021 dated 31/12/2021). Informed consent was waived off as no intervention was done and patient confidentiality was maintained.

### 2.2 Exclusion Criteria

Among the positive blood culture samples, samples showing normal skin commensals including *coagulase-negative Staphylococcus spp.* (CONS), *Propionibacterium acnes*, *Bacillus spp.*, *Corynebacterium spp.*, *Micrococcus spp.*, or "viridans"-group, streptococci in single blood culture was considered as contaminant and not included in the study. Also paediatric samples of age <18 years and samples with incomplete lab reports or clinical data were excluded from study analysis.

### 2.3 Data Collection

The data analysed in this study including the clinical details, laboratory tests and patient outcome were procured from the hospital information system and clinical records of the patient. For duplicate samples for each episode of bacteremia only the data for first positive sample was included in the study. A new episode of bacteremia in a same patient was defined if after 14 days of blood culture sample being negative, blood culture comes positive with the same or new organism. The data of new episode of bacteremia was included in the study. PCT levels were measured using the Roche Elecsys B.R.A.H.M.S. PCT test (Basel, Switzerland) (reference range, <0.05 ng/mL). Blood cultures were performed using the BACTEC blood culture bottles (Becton Dickinson, Sparks, MD). Blood

culture bottles when flagged positive were removed for Gram stain and subcultured on blood agar and McConkey agar. The isolated colonies were identified by automated system MALDI (matrix assisted laser desorption ionization time-of-flight [Biomerieux, USA]).

### 2.4 Statistical Analysis

Statistical analysis was performed using the software SPSS 20.0 (SPSS Inc., Chicago, IL, USA). Continuous variables were expressed as mean ± standard deviation. Kruskal–Wallis nonparametric analysis of variance was used for multi-group comparisons. Categorical variables were compared using the Chi-square test. All tests were performed as two-tailed tests. Multiple logistic regression analysis was used to determine the risk factors for mortality within 28 days. A p value of less than 0.05 was considered as statistically significant.

### 3. RESULTS

During the study period, 378 blood culture bottles were positive out of which 10 samples with

contamination, 5 with polymicrobial infection and 15 with incomplete lab results or clinical information were excluded from the study. After exclusion 348 positive blood samples with monomicrobial growth of bacteria or fungus were included in the study. In this study, 106 male and 242 females were included. Mean age of the patient included in the study 36.35±22 years.

### 3.1 Pathogens Isolated

Among the pathogen isolated Gram positive cocci were 151, Gram negative bacilli were 152, Gram positive bacilli were 7 and yeast isolates were 38. Most common Gram positive cocci was CONS (38.5%) and in Gram negative bacilli most common isolate was *Klebsiella species* (13.2%), some rare isolated pathogens like *Stenotrophomonas maltophilia*, *S.Typhi*, *Chryseobacterium indologenes*, *Serratia marcescens* were included in a category named other. The percentage of different pathogen isolated is shown in Fig.1 .

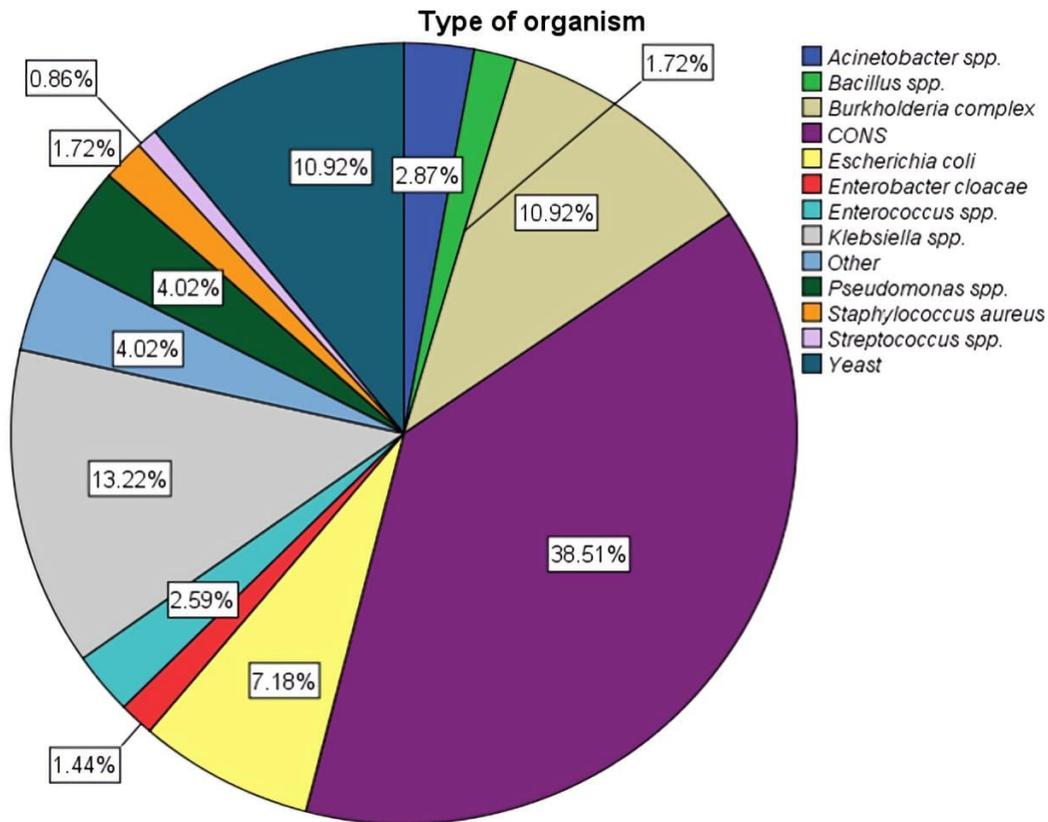


Fig . 1. The percentage of different pathogen isolated

### 3.1.1 PCT levels in different pathogens

On comparison of the PCT levels among different isolates significant difference was obtained between PCT values were significantly higher for Gram negative bacilli (14.5ng/ml  $\pm$  2.8) than Gram positive cocci(8.59ng/ml  $\pm$  1.5)and yeast(2.96ng/ml  $\pm$  0.56) shown in Fig. 2. On pairwise comparison of PCT values between different organism PCT values of CONS and *Klebsiella species* and also PCT values were significantly higher for *Klebsiella species*and *Escherichia coli* than yeast. Also the PCT values were significantly higher in *Burkholderia complex* and *Acinetobacter species* than CONS (p=0.036 and p=0.012 respectively). The PCT values were significantly different for yeast and *Acinetobacter species* (p= 0.032), shown in Fig. 3.No statistical difference was seen on intercomparison between the PCT values of other isolates included in the study.

Comparison was done between patient characteristics including age, gender, comorbidities and clinical course including length

of hospital stay and mortality in patients with low procalcitonin level (<0.5ng/ml) and high procalcitonin level (>0.5ng/ml). Statistically significant difference was found in procalcitonin levels of patient with malignancy. Patient mortality was also significantly different in low and high Procalcitonin levels shown in Table 1.

### 3.1.2 28 day mortality

On multivariate analysis, 28-day mortality was found not to be associated with age and gender while it was significantly associated with multidrug resistant (MDR) pathogen (p=0.006) and higher PCT level  $\geq$ 10 ng/ml (p=0.033), shown in Table 1.

### 3.1.3 Correlation between 28 day mortality and type of pathogen

In this study, there was no statistical difference of 28 day mortality rates between different types of pathogen isolated. Highest mortality was seen in *Klebsiella species* (23.46%).

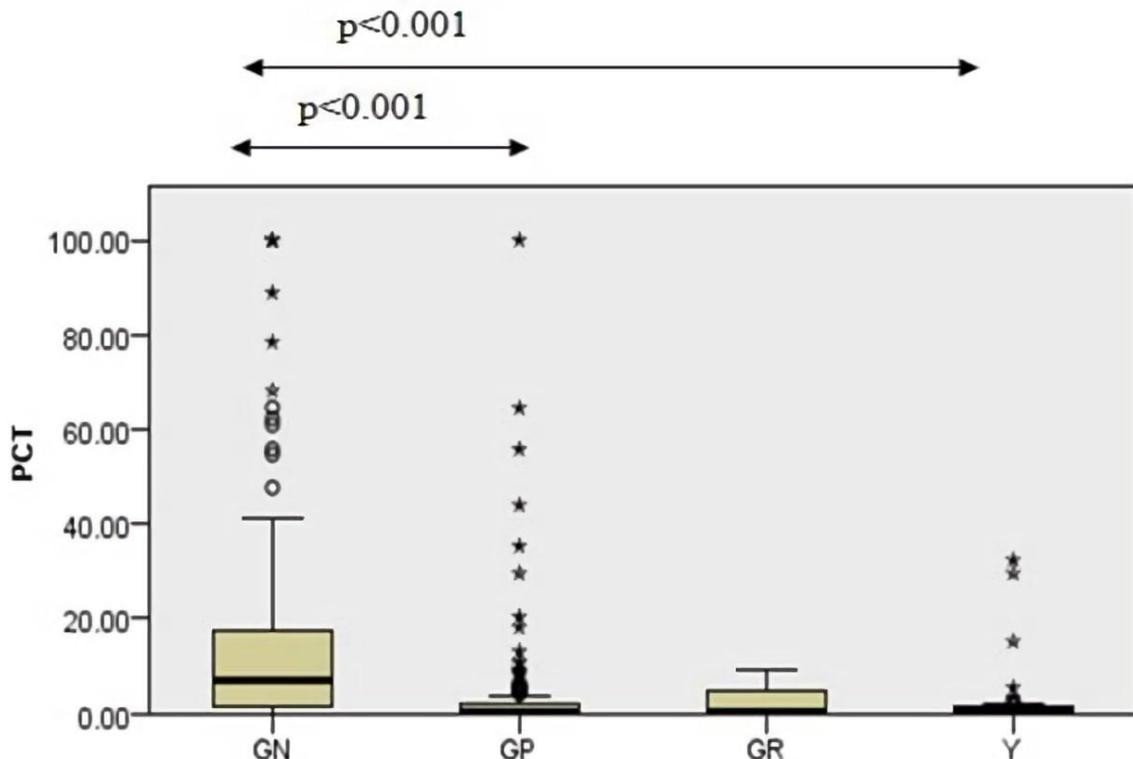


Fig. 2. Distribution of PCT in GN: Gram negative bacilli, GP: Gram positive cocci, GR: Gram positive bacilli and Y: Yeast

### 3.1.4 28 day mortality and PCT levels

The different PCT values in sepsis cases by different pathogens were classified into four groups i.e., group 1 (<0.5 ng/mL), group 2 (0.5 to <2.0 ng/mL), group 3 (2.0 to <10 ng/mL), and group 4 ( $\geq$  10 ng/mL) in accordance with one the study [9]. The 28-day mortality rates in groups 1, 2, 3, and 4 were 18.2%, 24.69%, 23.46%, and 33.33%, respectively. The 28-day mortality rate was significantly higher in group 4 than those of group 3 and 2.

## 4. DISCUSSION

The aim of present study was to investigate the level of PCT in cases of bacteremia and fungemia and also to find out PCT as a prognostic marker for patient outcome. The most predominant bacteria isolated in our study was

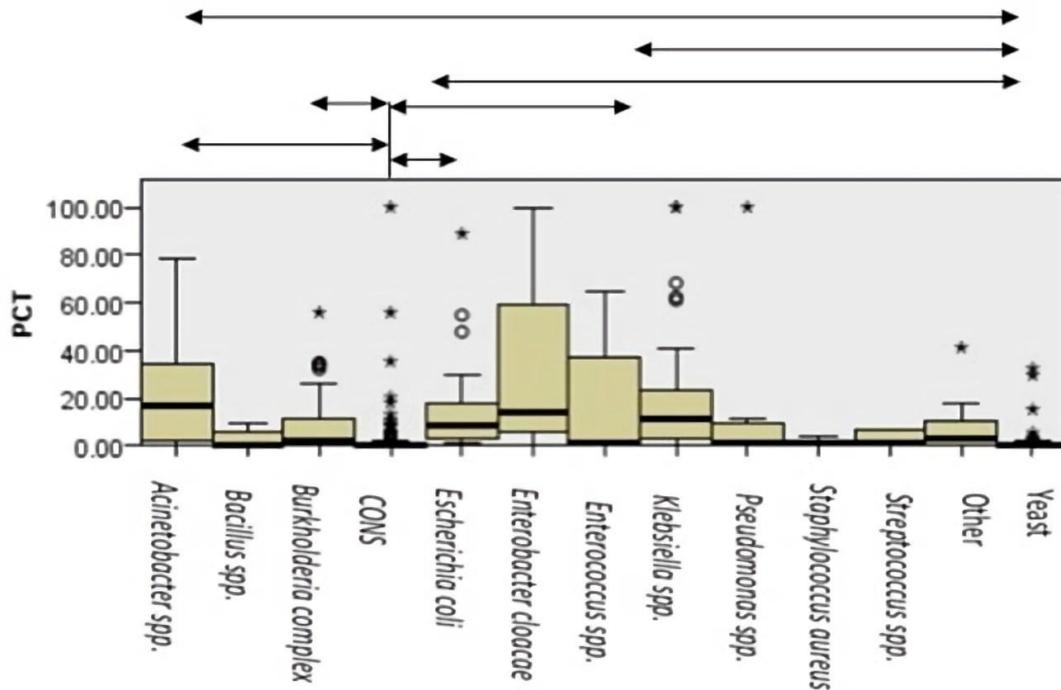
CONS (38.62%) which was higher than other studies [10,11]. In our study CONS positive cases from single blood culture positive cases was considered as contaminant and excluded from study. However complete exclusion of contaminants may not be possible, leading to higher isolation of CONS. In Gram negative bacteria sepsis with most common bacteria *Klebsiella species* (13.2%) followed by *Burkholderia complex* (10.9%), and *Escherichia coli* (7.2%) and yeast was isolated in 10.9% cases. In this study we found that the PCT values was significantly higher in cases of Gram negative bacteremia than the PCT values in Gram positive bacteremia and fungemia with yeast isolates. The PCT values in cases of *Klebsiella species* and *Escherichia coli* was found to be significantly higher than in CONS positive bacteremia, this finding is similar to other studies [12,13].

**Table 1. Multiple logistic regression analysis for 28-day-mortality in patients with bacteremia**

Independent variable	Odds ratio	Lower limit	Upper Limit	P value
Gender	0.56	0.272	1.156	0.117
Age	1.082	0.843	1.994	0.349
Malignancy	0.292	0.134	0.635	0.002
Gram positive cocci (GPC)	2.202	0.956	5.069	0.064
Gram positive bacilli (GPB)	1.006	0.166	6.092	0.995
Gram negative bacilli (GNB)	1.386	0.588	3.264	0.455
Yeast	0	0	0	0.999
Multidrug resistant bacteria(MDR)	3.683	1.449	9.360	0.006
PCT(0.5- <2.0 ng/ml )	0.800	0.380	1.686	0.558
PCT(2.0 - <10 ng/ml)	0.573	0.258	1.276	0.173
PCT( $\geq$ 10.0 ng/ml)	0.41	0.184	0.931	0.033

**Table 2. Comparison of patient characteristics between Low-Procalcitonin (<0.5 µg/L, and High-Procalcitonin ( $\geq$ 0.5 µg/L, Group B) patient groups**

Patient characteristics	Procalcitonin (<0.5ng/ml)n=95	Procalcitonin (>0.5ng/ml)n=253	P value
Age, Mean(S.D.)	37.6(12.4)	39.8(11.2)	0.78
Male	72(75.7)	188(74.3)	0.59
Comorbidity			
Diabetes mellitus	29(30.5)	82(32.4)	0.633
Chronic lung disease	15(15.8)	42(16.6)	0.550
Malignancy	9(9.4)	47(18.6)	0.022
Heart Failure	11(11.6)	31(12.3)	0.95
Chronic Kidney disease	9(9.4)	26(10.2)	0.64
Acute Pancreatitis	7(7.4)	19(7.5)	0.88
Length of hospital stay median in days (IQR)	7(4-10)	12(6-14)	0.61
Mortality	8(8.42)	67(26.4)	0.03



**Fig. 3. Distribution of PCT levels according to each pathogen. Arrow shows significant difference between PCT values of two pathogens (  $p < 0.05$  )**

There are several studies on contributing factors to mortality in sepsis. In one of the study the contributing factors included age, causative pathogen, primary source of infection and comorbidities in the patient [14]. In our study on multivariate analysis 28-day mortality was seen in high PCT levels ( $\geq 10$  ng/mL) and also in multidrug resistance (MDR) bacteremia including ESBL in *Klebsiella* species and *Escherichia coli*. The PCT levels were significantly higher in Gram negative sepsis and a better marker of sepsis and prognosis, compared to Gram positive bacterial and fungal sepsis which is similar to other study [15]. The different PCT levels in response to sepsis due to Gram-negative bacteria and Gram-positive bacteria is still not very clear. The most likely cause might be the difference in composition of cell membrane of Gram-negative and Gram-positive bacteria. The major difference is in the cell membrane composition is that Gram-negative bacteria have lipopolysaccharide (LPS) while it is absent in Gram-positive bacteria which have a thick peptidoglycan (PGN) layer [16]. One of the study by Oberhoffer et al [17], demonstrated that both LPS and sepsis-related cytokines increased PCT expression in human peripheral blood mononuclear cells (PBMCs) whereas in Gram positive bacterial sepsis due to

absence of LPS there is poor production of cytokine levels (TNF- $\alpha$  and IL-6) leading to poor PCT expression eventually [18,19]. Among the different causative pathogens isolated, highest mortality was seen in *Klebsiella* species (23.46%). It is studied that PCT levels are increased in normal neonates, in malignancy, trauma and in cases of major surgery, severe burns [10, 20]. Thus during interpretation of the PCT results the conditions where PCT values can be increased in absence of sepsis needs to be assessed carefully. In our study we have excluded the paediatric population to avoid false positive results.

## 5. CONCLUSION

The present study reported that the procalcitonin values were significantly higher in sepsis due to Gram negative bacteria as compared to Gram positive bacterial and fungal sepsis. In this study, we found procalcitonin to be a better prognostic marker, as 28-day mortality rates were significantly higher in patients with PCT levels  $\geq 10$  ng/ml. Thus, a vigilant monitoring of procalcitonin levels should be followed properly for better patient outcome. The antimicrobial therapy should be promptly started based on the hospital antibiogram and antibiotic susceptibility

results. Procalcitonin values should be evaluated at regular intervals for escalation or de-escalation of antibiotics. This will further reduce the unnecessary antibiotic usage burden and will also be helpful in control of emergence of multidrug resistant pathogens in hospital settings. However, in high-risk patient groups, with suspicion of sepsis, low procalcitonin levels should not be associated with delay in receipt of empirical antibiotics. Antibiotic stewardship programmes in hospital settings must correlate with the procalcitonin values, along with the blood culture and antibiotic sensitivity reports for avoidance of inappropriate administration of antibiotics to the admitted patients.

## CONSENT

It's not applicable.

## ETHICAL APPROVAL

This study was a part of a project and ethical clearance was taken from the institute ethics committee (PGI/DIR/RC/917/2021 dated 31/12/2021).

## COMPETING INTERESTS

Authors have declared that no competing interests exist.

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