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# **RESEARCH ARTICLE**

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# Microbiota of Bronchoalveolar Lavage Samples from Patients of Lower Respiratory Tract Infection – A Changing Trend

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# Abstract

Bronchoalveolar lavage (BAL) is the saline wash of the bronchial tree, which aids in diagnosing various pulmonary pathologies. The present study was contemplated with an aim to know the clinical, microbiological profile of BAL samples along with its sensitivity pattern and to assess its utility as a diagnostic tool. This was a prospective observational study, carried over 90 patients presenting with lower respiratory tract infections. The total microbial recovery rate from BAL was 39 (43.3%). The sensitivity, specificity and positive predictive value of BAL were found to be 76.4%, 89.7% and 90.6% respectively. Maximum isolates were bacteria (25.5%) followed by Mycobacterium tuberculosis (MTB) (16.6%) and fungi (1.1%). Predominant bacterial isolates were Gram-negative (81.5%) compared to Gram-positive (18.5%). Multidrug-resistance (MDR) in bacteria was seen in 59.2% of isolates. BAL is a valuable diagnostic tool to find not only bacterial but mycobacterial and fungal infections in patients with lower respiratory tract infection (LRTI). A trend towards LRTI with Gram negative infections is on the rise and they tend to be multi drug resistant. Hence checking susceptibility patterns is crucial to start evidence-based treatment.

Keywords: Bronchoalveolar lavage (BAL), LRTI, MTB, MDR, Bacteria, Fungi

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#### INTRODUCTION

Bronchoalveolar lavage (BAL) was introduced as a research-oriented procedure way back in 19701. It was originally used as a tool for obtaining secretions and cells from the lower respiratory tracts of patients with interstitial, occupational, or both types of pulmonary diseases<sup>2</sup>. It was used for first time in India in 1994 for its important role in diagnosis of infections and malignancies<sup>3</sup>. Lower Respiratory tract infections (LRTI), caused by various organisms like bacteria and fungi, can be isolated by culturing BAL samples<sup>4</sup>. Sputum samples are still considered effective in diagnosing infective lung conditions in our country, as they are easy to obtain. Although they have 24% diagnostic yield, contamination of sample from oral flora during expectoration makes it unsatisfactory for culture<sup>5</sup>. BAL has an advantage compared to repeat sputum sample collections as the contamination by oral flora can be avoided easily. Further, the quantitative culture with proper colony cut off provides a greater insight for therapy. Another distinct advantage of BAL is that being an outpatient procedure, it is very much useful in suspected cases of community acquired pneumonia (CAP). Several studies from western literatures show that BAL can be used to increase microbial isolation<sup>6</sup>. LRTI needs to be diagnosed with appropriate diagnostic tool for evidence-based management of patients. The clinical features of respiratory diseases vary with age, gender, season, associated risk factors and supplementary factors like environment and host<sup>7</sup>. As there are only few prospective studies reported from our country investigating the role of BAL in microbial isolation, the present study was undertaken to ascertain the clinical and microbiological profile of broncho alveolar lavage samples. We have also made an attempt to determine the susceptibility pattern, which can guide the clinicians to practice evidence-based management of LRTI.

The present study was carried out with an aim to find out the microbiota of BAL samples from patients of LRTI. The objective was to study the microbial profile of BAL samples, its utility as a diagnostic tool and to identify the divergent microbiota along with antimicrobial susceptibility pattern.

### MATERIALS AND METHODS

This observational prospective study was carried out in a study population of 90 patients in the Department of Microbiology, Mamata medical college, Khammam, Telangana, over a period of 01 year from April 2018 to April 2019 on both outpatients as well as admitted indoor patients in the Department of Pulmonology in Mamata General Hospital, affiliated to Mamata Medical College. The hospital mainly caters the rural population in the region. Institutional Ethical Committee (IEC) clearance was obtained before start of the study. The study was started initially as a short-term studentship (STS) project of Indian Council of Medical Research (ICMR - Ref No.2018-00184) and was further continued for total duration of 1 year. The inclusion criteria of the study were patients between 14 to 80 years of age presenting with symptoms of LRTI who were negative on sputum microscopy/culture and patients willing to participate in study. Exclusion criteria were patients below 14 years and above 80 years of age; patients with lung malignancy; unstable cardiac conditions and those who did not give consent to undergo the procedure.

The present study was planned in collaboration with Department of Pulmonology. Patients presenting with symptoms of LRTI like fever and cough, with or without expectoration, shortness of breath and chest pain ≤2 weeks were assessed with thorough history and physical examination. Relevant blood and radiological investigations in the form of chest X - ray was done in every case. As study protocol, sputum/ Induced sputum samples of all the patients were sent to microbiology lab. Gram stain was performed initially followed by Bartlett's grading to check sputum sample quality8. Zeihl-Neelson (ZN) stain was performed as per Revised National Tuberculosis Control Programme (RNTCP) guidelines. KOH mount was done in suspected cases of fungal etiology. Bacterial culture was done on blood agar, macconkey agar, chocolate agar and incubated at 37°C. Chocolate agar was incubated in candle jar for 24-48 hours to check the growth of fastidious bacteria like Haemophilus influenzae, Haemophilus parainfluenzae and Moraxella catarrhalis<sup>9</sup>. Further identification of isolates was done by routine biochemical tests. Sabourauds dextrose agar (SDA) was used for fungal culture. Patients, in whom sputum/induced sputum samples showed non satisfactory growth / no pathogenic organism/ mixed flora, were further followed up for a period of 03 days. If symptoms persisted during follow up, BAL was performed in all those cases. These were the patients with diagnostic dilemma as clinical, radiological (chest X-ray) and microbiology report were not analogous. High resolution computerised tomography (HRCT) was also done in those patients who had inconclusive evidence on chest X-ray<sup>10</sup>.

A standard BAL procedure was done in bronchoscopy room with 5 mm flexible fibre optic bronchoscope under sterile conditions. 10% lignocaine spray was used for local anaesthetic effect. Minimal sedation was used only in anxious patients. Access to BAL was achieved through nasal approach. BAL aliquots of 20 mL were used in 0.9% saline via syringe up to a total volume of 120-150 mL. Suction pressure of 100 mm of H<sub>2</sub>O was applied and about 70% of total instillation was retrieved. This fluid was then immediately sent to microbiological laboratory for examination.

Samples were subjected to Gram's stain for quality check. Samples were not processed further if more than 1% epithelial cells were seen on low power magnification, along with absence or presence of bacteria in less than than 1-2/ oil immersion field which was suggestive of oropharyngeal contamination<sup>11,12</sup>. On accepted samples Ziehl-Neelsen (ZN) staining was done for Mycobacterium tuberculosis (MTB). KOH mount was done for fungal elements. These were preliminary microscopy procedures on the sample. The sample was inoculated on Blood agar, MacConkey agar and chocolate agar using a sterile 4 mm nichrome loop (0.01ml) and incubated at 37°C for 18-24 hours for quantitative bacterial culture using standard laboratory techniques. Chocolate agar was placed in candle jar for 48-72 hrs to look for fastidious organisms. Sample was also inoculated on Sabaroud's dextrose agar (SDA) for fungal culture and incubated at 37°C for 04 weeks for growth. For MTB detection, Cartridge Based Nucleic Acid Amplification Test (CBNAAT) was performed as per RNTCP guidelines.

For bacterial growth on culture plates, the colony forming unit was calculated with cut

off > 105 CFU/ ml<sup>13</sup>. Colonisers were ruled out from true pathogens by strict diagnostic criteria if colony count was significant<sup>14</sup>. Antibiotic sensitivity testing was done on bacterial isolates as per Clinical and Laboratory Standards Institute (CLSI) 2018-19 guidelines by Kirby-Bauer's disc diffusion method along with local antibiogram guidelines as per institute protocol.

#### Definition

Pneumonia as in LRTI was diagnosed as patients with a chest radiograph finding suggestive of new/progressive/persistent (>24hours) infiltrate and with minimum of 02 criteria like purulent expectorant; body temperature of > 38.5°C or < 36.5°C; WBC count of >10,000 cells/μL or <4,000 cells/μL and breathlessness. HRCT evidence of a segment involvement, peribronchovascular or scattered ground glass or reticular opacity or consolidation congruent with lung involvement in acute phase was considered as evidence of pneumonia. Along with this Quantitative BAL cultures of >10<sup>5</sup> CFU/ml were considered as diagnostic of LRTI.

# Multidrug-resistant bacteria (MDR)<sup>7,15</sup>

Multidrug-resistant (MDR) bacteria were S. pneumoniae resistant to penicillin and other broad-spectrum agents like fluoroquinolones and macrolides, Methicillin resistant S. Aureus (MRSA) and MDR Gram negative bacilli. MDR Gram negative bacilli were Escherichia coli, Klebsiella spp., Enterobacter spp and Citrobacter spp, which were resistant to at least 03 of the following group of antibiotics, viz; piperacillin or ampicillin/sulbactam, aminoglycosides, third or fourth generation cephalosporins, carbapenems and fluoroquinolones. Pseudomonas aeruginosa resistant to at least 03 of the following antibiotic groups was also classified as an MDR, viz; piperacillin, aminoglycosides, cephalosporins, fluoroguinolones and carbapenems. *Acinetobacter* baumannii resistant to most of the antimicrobial agents with/without imipenem resistance was considered as MDR.

# Statistical analysis

Data was entered in Microsoft excel sheet. For analysis of continuous data, unpaired t test was used. For analysis of categorical data Pearson's  $\varkappa^2$  test was used. Differences were considered statistically significant, if p<0.05. IBM SPSS Statistics for Windows, version 24 (IBM Corp.,

Table 1. Clinical profile of patients (n=90)

Variables	Total (%)	
Male	61(67.7%)	
Female	29(32.2%)	
Mean Age(yrs)	43(14-80)	
Smoking history	32 (35.5%)	
Comorbidity		
Diabetes	23 (25.5%)	
Chronic Bronchitis	34 (37.7%)	
COPD*	29 (32.2%)	
Previous TB	11 (12.2%)	
Symptoms/Signs		
Fever	69 (76.6%)	
Sputum	68 (75.5%)	
Dyspnea	83 (92.2%)	
Leukocytosis/	41(45.5%)	
Leucopenia		
IPD**	22(24.4%)	

<sup>\*</sup>Chronic Obstructive Pulmonary disease, \*\*In patient department

Armonk, N.Y., USA) software program was used for statistical calculations.

## **RESULTS**

Study sample consisted of 90 patients who were clinically suspected but were sputum negative/unsatisfactory. Clinical profile of patients is shown in Table 1 with 61 (67.7%) males and 29 (32.2%) females with mean age of 43 years. Predisposing factors like smoking was seen in 32

**Table 2.** Contingency table – BAL Quantitative culture with LRTI {Total microbial isolation= 39(43.3%)}

	LR	ΓI <sup>£</sup>	Total
	Present	Absent	
BAL >10 <sup>5</sup> CFU/m <10 <sup>5</sup> CFU/m		4 35	43 47
TOTAL		39 (43.4%)	90

<sup>&</sup>lt;sup>£</sup>Lower Respiratory Tract Infection

(35.5%), diabetes in 23 (25.5%), chronic bronchitis in 34 (37.7%), past history of tuberculosis in 11 (12.2%) and chronic obstructive pulmonary disease (COPD) in 29 (32.2%) patients. Dyspnoea was the most common symptom (92.2%) followed by fever (76.6%) and cough with expectoration (75.5%). In patient treatment was needed for 22 (24.4%) patients.

Clinical diagnosis of LRTI could be achieved in 51 patients as per predefined criteria. 39 (76.4%) of those 51 patients had colony count of >105 CFU/mL, with true pathogens (Mycobacteria and fungi included) on BAL fluid culture. Of 90 patients 39 (43.4%) patients did not meet pneumonia criteria. 04 among those 39 patients, who had significant colony count of colonizers

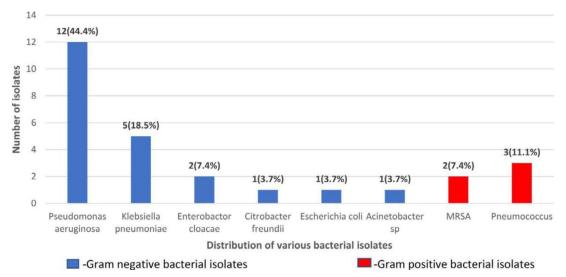


Fig. 1. Distribution of bacterial isolates from BAL fluid (> 105 CFU/ml) Total (n=27) (30%)

Sensitivity= 39/51=76.4%

Specificity=35/39=89.7%

Positive predictive value=39/43=90.6%

Negative predictive value=35/47=74.4%

**Table 3.** Distribution pattern of microorganisms on microscopy and culture from BAL fluid (n=90) Total recovery = 39(43.3%)

	Bacteria	Mycobacteria	Fungi	Total
Microscopy	13(14.4%) (Gram stain)	8(8.8%) (ZN Stain)	1(1.1%) (KOH mount)	22 (24.4%)
Culture/CBNAAT <sup>†</sup>	23 (25.5%) Culture	15 (16.6%) CBNAAT	1(1.1%) Culture	39 (43.3%)

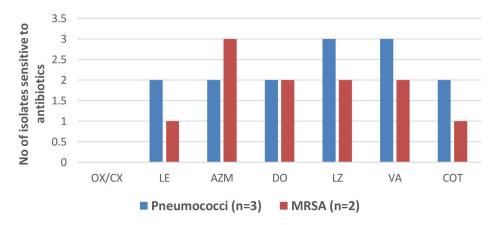
<sup>&</sup>lt;sup>†</sup> Cartridge based nucleic acid amplification test.

on BAL culture, were further retrospectively confirmed to actually harbour coloniser, even with BAL procedure as these patients did not respond to antibiotic treatment on follow up. Thus, contingency table was made showing sensitivity of 76.4%, specificity of 89.7%, positive predictive value 90.6% and negative predicative value 74.4% in BAL (Table 2).

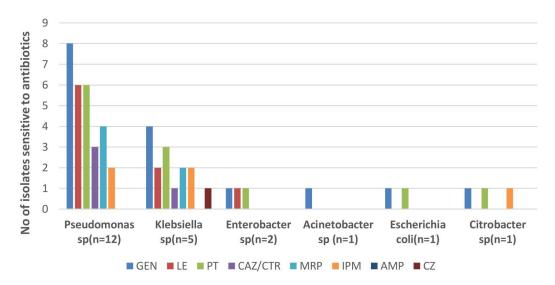
On microscopy, Gram stain could accurately identify bacteria in 13 (14.4%) samples; ZN stain for MTB in 08 (8.8%) and KOH mount for fungi in 01 (1.1%) sample. Bacterial growth was seen in 23 (25.5%) samples. CBNAAT was positive in 15 (16.6%) samples and Aspergillus flavus grew in a single sample (1.1%). Bacterial isolation was better in BAL when compared to mycobacteria and fungi as they usually contributed for majority of lower respiratory tract infections (LRTI). Out of 90 samples, 23 (25.5%) samples showed bacterial growth which was in significant colony count

(>10°CFU/ml) with monomicrobial growth in 19 samples (19 isolates) and polymicrobial growth in 04 samples (08 isolates) resulting in total 27 bacterial isolates. 49 (54.4%) samples showed no growth/ mixed growth/no pathogenic growth/ significant growth(>10°CFU/ml) with colonisers, whereas 18 (20%) samples had insignificant colony count. Insignificant growth or coloniser with significant colony count was not a useful indicator in diagnosing LRTI in the present study. Thus, BAL could accurately aid in diagnosing bacterial pneumonia in this study for up to 25.5%, MTB in 16.6% and fungal aetiology in 1.1% with total microbial recovery rate of 39 (43.3%) (Table 3).

Distribution of Gram positive and Gramnegative isolates is shown in Fig. 1. Of total 27 isolates, 22 (81.5%) were Gram negative isolates and 05 (18.5%) were Gram positive. Amongst Gram negative isolates, *Pseudomonas aeruginosa* 12 (44.40%) and *Klebsiella pneumoniae* 05 (18.5%)



**Fig. 2.** Antibiotic sensitivity pattern of Gram-positive isolates from BAL fluid(n=5). OX-Oxacillin; CX=Cefoxitin: LE=Levofloxacin: AZM=Azithromycin; DO= Doxycycline; LZ=Linezolid; VA=Vancomycin: COT=Cotrimoxazole



**Fig. 3.** Antibiotic sensitivity pattern of Gram-negative isolates from BAL fluid(n=22). PT= Piperacillin-Tazobactum: CAZ=Ceftazidime: CTR=Ceftriaxone: MRP=Meropenem: IPM= Imipenem: AMP= Ampicillin: CZ=Cefazolin.

were isolated predominantly in BAL samples, followed by *Enterobacter cloacae* complex 02 (7.40%), *Citrobacter freundii* 01 (3.7%), *Escherichia coli* 01 (3.7%) and *Acinetobacter spp* 01(3.7%). Among Gram positive isolates, Pneumococci 03 (11.10%) and MRSA 02 (7.40%) cases were found. Antibiotic susceptibility pattern of grampositive isolates is shown in Fig. 2. All 3 isolates of *pneumococci* were resistant to penicillin which was checked by Oxacillin disc and Penicillin-Minimum inhibitory concentration (MIC) E test. Only 01 (33.3%) among those was MDR. Both MRSA strains isolated were sensitive to vancomycin (by MIC-E test), linezolid, and ciprofloxacin. Thus, a total of 03 (11.1%) Gram positive MDR s were recovered.

Antibiotic sensitivity pattern in Gram negative isolates varied. 50% (6/12) isolates of *Pseudomonas aeruginosa* were MDR, whereas 66.7% (8/12) samples showed Carbapenem resistance. 60% (3/5) of *Klebsiella pneumoniae* were MDR. Single isolate of *Acinetobacter spp* was MDR with sensitivity to gentamicin and ampicillin sulbactam. *Enterobacter cloacae* complex (1 isolate) and single isolates of *Escherichia coli* and *Citrobacter freundii* were MDRs. Many Gram-negative isolates showed carbapenem and cephalosporin resistance except one *Klebsiella pneumoniae* which was sensitive to ceftriaxone (Fig. 3). MDR Gram negative bacteria isolated were 13 (48.1%). Thus, a total of 16 (59.2%) MDRs

were isolated from 27 bacterial isolates. Rifampicin resistance was detected in 26.6% (4/15) samples by CBNAAT for MTB. Antifungal susceptibility testing for Aspergillus could not be evaluated.

Outpatients without MDR bacteria in BAL were treated as per sensitivity reports. Patients who had MDR bacterial isolation were admitted in view of associated comorbid conditions and presumed morbidity. Renal function tests were done. Two patients with MRSA pneumonia recovered with linezolid therapy for 7-10 days. Single patient with MDR pneumococcal pneumonia recovered with levofloxacin for 7-10 days. Six (6) patients, positive for MDR Gram negative Enterobacterales, were treated with meropenem/ceftazidime-avibactum/ colistin/gentamicin, in whom 03 recovered and 03 were lost to follow up. 07 patients with MDR Pseudomonas aeruginosa and MDR Acinetobacter were treated with aminoglycoside/colistin, out of which, 05 cases recovered and 02 were lost to follow up. Patients diagnosed with MTB were treated as per RNTCP guidelines. Single patient who was diagnosed with pulmonary aspergillosis responded well to oral itraconazole therapy.

# **DISCUSSION**

LRTI affected predominantly middle-aged men in the present study (Mean age-43 yrs) which was comparable to study done in 2020 by Anisha et al<sup>16</sup>. This male predominance could be due to

more outdoor activity, smoking, cultural conditions existing in our country. Comorbidities like diabetes 23 (25.5%), chronic bronchitis 34 (37.7%), COPD 29 (32.2%) and previous history of MTB 11 (12.2%) led to discrepancy in diagnosing LRTI with sputum samples. Hence BAL had to be performed in these patients to aid the diagnosis.

Scientific literature is scarce with regard to the diagnostic utility of BAL in LRTI. Jimenez et al.<sup>17</sup> studied patients with suspected LRTI (CAP) without previous antibacterial treatment, BAL has adequate correlation equal to protected brush specimen (PSB) in diagnosing pneumonia. Sensitivity of BAL in the present study was less compared to the study done in 2019 by Travar et al.<sup>18</sup> for diagnosing ventilator associated pneumonia, where sensitivity was 91% but it had lesser specificity of 70% compared to the present study, where sensitivity and specificity were 76.4% and 89.7% respectively.

Overall microbial isolation, in the present study was 43.3%, which was found to be similar with other studies like Veleza et al. 19 and Kottmann et al.20, where the positive yield was 51.6% and 55.8% respectively. The comparative slight lesser positivity rate in the present study might be for the fact that the study was done in general population, with less sample size and immunocompetent subjects, whereas other studies quoted above were done in immunocompromised patients. 27 (30%) bacterial isolates grew in the present study which is less compared to a recent study done by Adhikari et al<sup>21</sup> in 2021, where 95.3% bacterial isolation rate was seen from BAL samples. Utility of BAL in diagnosing mycobacterial infections with Ziehl-Neelson stain was done by Vivek et al.<sup>22</sup> in 2016 with positivity rate equal to above study. CBNAAT being extremely sensitive yielded better diagnosis of TB in 15 (16.6%) cases in the present study. Our study had 1.1% fungal growth from BAL. In the study of Vivek et al<sup>22</sup>, more fungal isolates were found with a predominance of candida albicans (37%) followed by Aspergillus spp (14.8%). The reason for less fungal growth in the present study could be that all the patients were immunocompetent. BAL has shown to be good indicator of fungal infections with comparison equal to that of bronchial biopsy4.

In studies by Kottman et al.<sup>20</sup> and Bari et al.<sup>23</sup>, predominance of aerobic Gram-

negative bacteria over Gram positive bacteria was observed. Klebsiella was also the most common pathogen isolated from studies done by Sarmah et al.<sup>24</sup>, Lin et al.<sup>25</sup> and Singh et al.<sup>26</sup>. Pseudomonas aeruginosa was the commonest isolate in a study done by Magazine et al<sup>27</sup> in 2013. The results of present study were comparable to afore mentioned studies demonstrating gram negative organisms emerging predominantly as cause of LRTI from BAL samples. However, *Streptococcus pneumoniae* was the most common isolate from BAL, found in the study of Bansal et al.<sup>28</sup> and Capoor et al.<sup>29</sup> which contrasts with the present study.

The present study correlates with the study done by Magazine et al.27 where 87.9% isolates of P. aeruginosa had resistance to cephalosporins and 75% of the isolates of Acinetobacter spp were resistant to meropenem. The findings of the present study were discordant with those of a retrospective study done by Goel N et al.30, which revealed that only 25.6% of the isolates of Acinetobacter species, 22.8% the isolates of *P. aeruginosa* and 09% of the isolates of Klebsiella species were resistant to meropenem. This difference could be due to the different set up in which study was carried out. In the present study, patients who developed MDR infection of gram negative/gram positive organisms were outpatients, thus needing a more in-depth analysis to address this issue as treatment options for CAP with MDR Enterobacterales, Pseudomonas aeruginosa and Acinetobacter spp are limited due to lack of appropriate antibiotics<sup>31,32</sup>. Increasing carbapenem resistance in gram negative isolates from BAL samples has been seen in few studies similar to present study<sup>33,34</sup>.

Current medical literature suggests that Streptococcus pneumoniae is the most common Gram-positive bacterium causing pneumonia, whether community-acquired or hospital acquired. However, as it is evident from the present study, there was predominance of gram-negative organisms and emergence of MDR organisms from those proportion of patients where sputum sample did not yield any conclusive result. This changing trend of gram-negative predominance further needs to be investigated with large multicentre studies. This trend needs to

be kept in mind along with patient clinical profile as MDR organisms are emerging as a cause of LRTI due to improper antimicrobial stewardship even in untreated patients.

# Limitations of the study

Atypical bacterial pathogens and viruses could not be isolated due to resource constraints and non-affordability of patients in this region. Furthermore, studies need to be conducted to address this issue of emergence of multidrug resistant organisms in community acquired pneumonias.

# CONCLUSION

BAL, as a diagnostic tool can be used accurately to identify varied infections and their causative organisms with sensitivity pattern. BAL is useful not only in bacterial infections but also in fungal infections. A country like India, where tuberculosis is common, BAL may play a vital role in its diagnosis. Clinical profile of the patient needs to be kept in mind while evaluating LRTI. BAL provides that unique information which helps to switch from empirical to precise evidence-based therapy.

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## **CONFLICT OF INTEREST**

The authors declare that there is no conflict of interest.

# **AUTHORS' CONTRIBUTION**

DI contributed to formulating research hypothesis, collection of samples, interpretation and analysis of data along with manuscript writing. AB was involved data analysis, interpretation and contributed novel ideas in presentation of data. PI supervised and reviewed, edited the manuscript, and designed the figures and tables. PP contributed for sample collection, references gathering and patient communication.

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#### **DATA AVAILABILITY**

All datasets generated or analysed during this study are included in the manuscript.

#### **ETHICS STATEMENT**

This study was approved by the institutional ethical clearance committee. IEC/IRB No-05/2018.

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