

Original Research Article

<https://doi.org/10.20546/ijcmas.2019.801.072>

Study of Bacterial Isolates in Community Acquired Pneumonia

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ABSTRACT

Keywords

Pneumonia,
Infection, Sputum
culture, Klebsiella

Article Info

Accepted:
07 December 2018
Available Online:
10 January 2019

Community Acquired Pneumonia (CAP) is an infection of pulmonary parenchyma. Despite availability of potent antibiotics, CAP remains a common and serious illness with significant morbidity and mortality. Objective of the study is to identify the bacteria causing community acquired pneumonia and risk factors associated with it. 100 clinically diagnosed CAP patients attending medical out-patient and admitted in Upgraded Osmania General Hospital selected. Study was conducted during Sept 2016 to Oct 2017. Sputum samples were cultured and organism identified by standard biochemical tests. Out of 100 included, 52 had identifiable etiology. Most frequent organism was *Klebsiella pneumoniae* (n=27) followed by *Staphylococcus aureus* (n=14). People in the age group of 45-65 years were more susceptible. Major risk factor was smoking.

Introduction

Community Acquired Pneumonia (CAP) is a commonly encountered lower respiratory tract infection by clinicians. It is defined as, “an infection of the pulmonary parenchyma. Infectious Diseases Society of America defines Community Acquired pneumonia (CAP) as “an acute infection of the pulmonary parenchyma that is associated with at least some symptoms of acute infection (cough, dyspnoea, fever) accompanied by the presence of an acute infiltrate on a chest radiograph or auscultatory findings (ronchi, crepitations) consistent with pneumonia in a patient not hospitalized or residing in a long-term care facility for more than 14 days before onset of

symptoms”. CAP is usually acquired via inhalation or aspiration of pulmonary pathogenic organisms into a lung segment or lobe. Less commonly, from secondary bacteraemia from a distant source or by contiguous extension from infected pleural or mediastinal space. Pneumonia may present as acute (community acquired or nosocomial), sub-acute or chronic. CAP commonly affects people of all ages, with higher incidence occurring in very young to very old age groups.

In the United State, pneumonia is the sixth leading cause of death with annual incidence of CAP ranging from 4 to 5 million cases. But the problem is much greater in developing

countries, though definite statistics are lacking, pneumonia remains a leading cause of death in India according to study by Bansal S (2004). Pneumonia is increasingly common in patients with co-morbidity like chronic obstructive pulmonary disease (COPD), Diabetes mellitus (DM), renal failure, Congestive heart failure (CHD) and Bronchiectasis. The cause of CAP is often difficult to establish. Despite the progress made in the clinical diagnosis of pneumonia, it takes a few days to identify the causative microorganism and the aetiology of half of all patients with CAP remains uncertain as per study conducted by Ishida T (1998). The bacteriological profile of CAP is not the same across various countries. It also varies within the same country with time, due to differences in the frequency of use of antibiotics, environmental pollution, awareness of the disease and life expectancy. Clinicians need reliable data on the prevalence of different etiological agent in their area of residence.

The present study has been conducted in Upgraded Department of Microbiology, Osmania General Hospital, Hyderabad, Telangana, with the objective to know the prevalence of etiological microorganism of CAP and risk factors associated with it.

Materials and Methods

This study was undertaken in a 750 bedded multi-specialty referral hospital in Hyderabad catering to both urban and semi-urban populations. This prospective study was carried out after taking clearance from ethical committee, in the Department Of Microbiology, Osmania general hospital, Hyderabad, Telangana.

Source of data

Patients attending Osmania General Hospital above 15 years of age clinically diagnosed as

CAP were selected from Medicine Department. The study conducted during a time period of 1 year from September 2016 to October 2017.

Sample size

100 patients of CAP attending medical out-patient department and admitted in Upgraded Osmania General Hospital, Hyderabad were included in the study after taking informed consent

Inclusion criteria

All patients over 15yrs attending medical out-patient department or admitted with at least two of the following symptoms.

- Fever
- Cough
- Production of purulent sputum
- Breathing difficulty
- Chest pain
- Leucocytosis (WBC > 10,000/cumm)
- New infiltrate in chest radiograph
- Patients not on antibiotic therapy.

Exclusion criteria

- Patients already on antibiotic therapy
- Patients not willing to give informed consent
- Patients with Pulmonary infarction, pulmonary edema, interstitial lung disease.
- Patients receiving immunosuppressive therapy.
- HIV patients

Sample collection

Sputum (deeply coughed) from the patients is collected in sterile wide mouthed leak proof container. In patients who could not expectorate sputum spontaneously, sputum induction was done using 3% hyper-tonic saline nebulization. Label the sample

appropriately and transport it to laboratory immediately.

The following data were recorded on enrolling: age, gender, comorbidities, antimicrobial treatment prior to enrolment, duration of symptoms before the diagnosis of pneumonia, clinical symptoms (body temperature, pleuritic chest pain, purulent sputum), haematology (total WBC with differential counts, platelet count, hemoglobin), chest radiographic pattern, and smoking and alcohol consumption.

Sputum processing:

Macroscopic appearance

Nature of the sputum was observed-purulent, muco-purulent, mucoid, or blood stained.

Microscopic examination

Gram's stain

Bartlett's grading system was used for assessing the quality of sputum samples.

Culture

Sputum was inoculated onto 5% sheep Blood agar, Chocolate agar and Mac Conkey agar.

Plates were incubated for 18-24 hours at 37⁰c in candle jar.

The organisms isolated were identified by standard biochemical reactions.

Results and Discussion

112 patients with age >15 years of age, attending medical out-patient or admitted in Osmania General Hospital, Hyderabad, between September 2016 and October 2017 were included in the study. After sputum microscopy, 12 were excluded from the study

because, 7 sputum samples did not satisfy Barlett scoring criteria and 5 were positive for *Candida* species. From the 100 which were included in the study, 71 were males and 29 were females (Fig. 1 and 2).

This study was conducted to find out the bacterial etiology in patients with Community acquired pneumonia and sensitivity profile, as it is one of the leading causes of the morbidity and mortality in the world as per study conducted by Bansal (2004). Aetiological agents vary from area to area, so do their antibiotic susceptibility profile.

In the present study, 52% of bacterial isolates were recovered from 100 sputum samples which were included in the study. A similar percentage of was reported by Madhulata *et al.*, (2013) whereas 71.6% positivity of culture was shown by Ramana *et al.*, (2013) from Andhra Pradesh

Males were found to be more commonly affected with a M: F ratio of 2.4:1 which correlated to a study by Madhulata *et al.*, (2013) who also found males were commonly affected, with the M: F ratio being 2.7:1. A study by Wattanathum *et al.*, (2003) showed Male to female ratio 1.6:1, Basheer shah *et al.*, (2010) and Rohinikumar *et al.*, (2015) found male to female ratio of 1.3 and 1.7:1 respectively.

In our study, age of patients ranged from 15 – 93 yrs. The most affected age group was 45-65 yrs, which correlated with study by Reechaipichitkel Wipa *et al.*, (2002) who found the mean age was 56.9 years.

Smoking is well known and important risk factor for community acquired pneumonia through alteration in mechanisms of host defense system. It causes changes in mucociliary clearance, bacterial adherence and respiratory epithelium. Tobacco smoking is most important risk factor for development of

COPD and it is recognized as risk factor for other respiratory infections. In the present study, most common identified risk factor was smoking 55% followed by Alcohol consumption in 30%, Diabetes Mellitus in 20% and COPD in 11%. Study conducted by Bansal *et al.*, (2004) showed 71%, Shah Bashir Ahmed *et al.*, (2010) found smoking as a predisposing factor in 65% followed by COPD in 57% and Madhulata (2013) reported smoking as risk factor in 45% followed by COPD in 26% and Diabetes in 8%. In contrast

Oberoi (2006) found 26.6% and Rohinikumar (2015) found smoking as risk factor in 37% cases (Fig. 4).

Maximum number of patients presented with cough, fever, sputum production, pleuritic chest pain, and dyspnea, this correlated with previous studies (Fig. 3). Sputum culture was positive in 52%. Similar observations were reported by Madhulata *et al.*, (2013) and Chawla *et al.*, (2008) (Table 1–9).

Table.1 Age and Sex wise distribution of cases (n=100)

Age	No. of cases	Males	Females
15-25	8	5	3
26-35	13	8	5
36-45	12	8	4
46-55	13	8	5
56-66	26	19	7
66-75	21	19	2
76-85	5	3	2
86-95	2	1	1
Total	100	71	29

Table.2 Common symptoms observed in the study group

Symptom	No. of cases	Percentage (%)
Cough with expectoration	98	98%
Fever	92	92%
Chest pain	57	57%
Dyspnea	60	60%

Table.3 Associated risk factors noted in the study group

Risk factor	No. of cases	Percentage %
Smoking	55	55%
Alcohol	30	30%
Diabetes mellitus	20	20%
COPD	11	11%
Asthma	3	3%
Heart disease	3	3%

Table.4 Culture positives in sputum (n=100)

Sputum culture	No. of samples	Percentage %
Positive	52	52%
Negative	48	48%

Table.5 Total no. of isolates in sputum culture n=52

Isolates	No.	Percentage %
<i>Klebsiellapneumoniae</i>	27	51.9
<i>Staphylococcus aureus</i>	14	26.9
<i>Escherichia coli</i>	4	7.6
<i>Pseudomonasaeruginosa</i>	3	5.7
<i>Streptococcus pneumoniae</i>	3	5.7
<i>Streptococcus pyogenes</i>	1	1.9
Total	52	100

Table.6 Distribution of isolates according to age

Age	No Pts	<i>K.pneumoniae</i>	<i>Staph aureus</i>	<i>E.coli</i>	<i>Pseudo Monas</i>	<i>S. pneumonia</i>	<i>S. pyogenes</i>	Total isolates
15-25	8	3	-	-	-	-	-	3
26-35	13	2	1	1	-	-	-	4
36-45	12	3	1	1	-	-	-	5
46-55	13	2	2	-	1	1	-	6
56-65	26	8	5	1	1	1	1	17
66-75	21	8	2	1	1	-	-	12
76-85	5	-	3	-	-	1	-	4
86-95	2	1	-	-	-	-	-	1
Total	100	27	14	4	3	3	1	52

Table.7 Studies showing the most common affected sex

Author	Year	Most common in	M:F ratio
WattanathumA <i>et al.</i> ,	2003	Males	1.6:1
Basheer shah <i>et al.</i> ,	2010	Males	1.3:1
Madhulata CK <i>et al.</i> ,	2013	Males	2.7:1
Rohinikumar <i>et al.</i> ,	2015	Males	1.7:1
Present study	2017	Males	2.4:1

Table.8 Occurrence of Clinical symptoms in various studies

Author	Year	Fever (%)	Cough + expectoration (%)	Chest pain (%)	Dyspnoea (%)
Irfan M <i>et al.</i> ,	2009	77.5	72	23	46
Shah BA <i>et al.</i> ,	2010	95	99	75	-
Madhulata CK <i>et al.</i> ,	2013	75	99	37	45
Rohinikumar <i>et al.</i> ,	2015	91	81	30	44
Present study	2017	92	98	57	60

Table.9 Sputum culture positivity in various studies

Author	Place	Year	Culture positive %	<i>K. pneumoniae</i> isolates (%)	<i>S. aureus</i> isolates (%)
Madhulata <i>et al.</i> ,	India	2013	54.5	44.7	2.6
Mythri <i>et al.</i> ,	India	2013	52.7	55.2	2.6
Priyanka Paul	India	2013	66.4	33.3	17.7
TripathiPurti <i>et al.</i> ,	India	2014	42	42	20.3
Rohini Kumar <i>et al.</i> ,	India	2015	46	19.5	-
Sunil Vijay	India	2016	77	36.7	22.2
Present study	India	2017	52	51.9	26.9

Fig.1&2

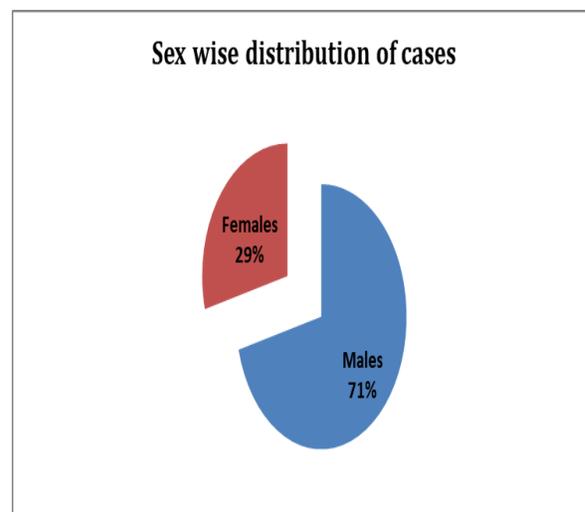
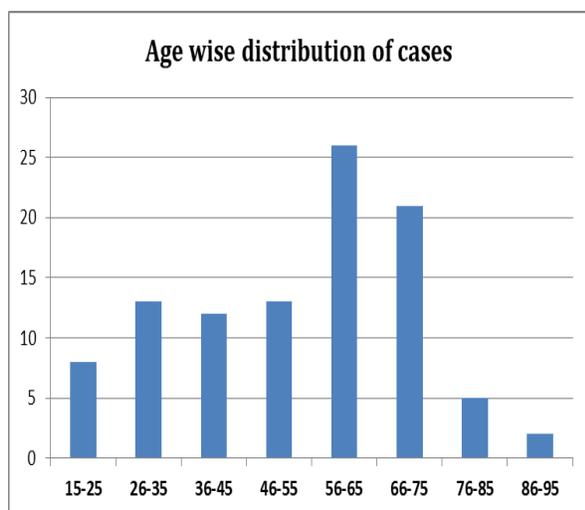


Fig.3

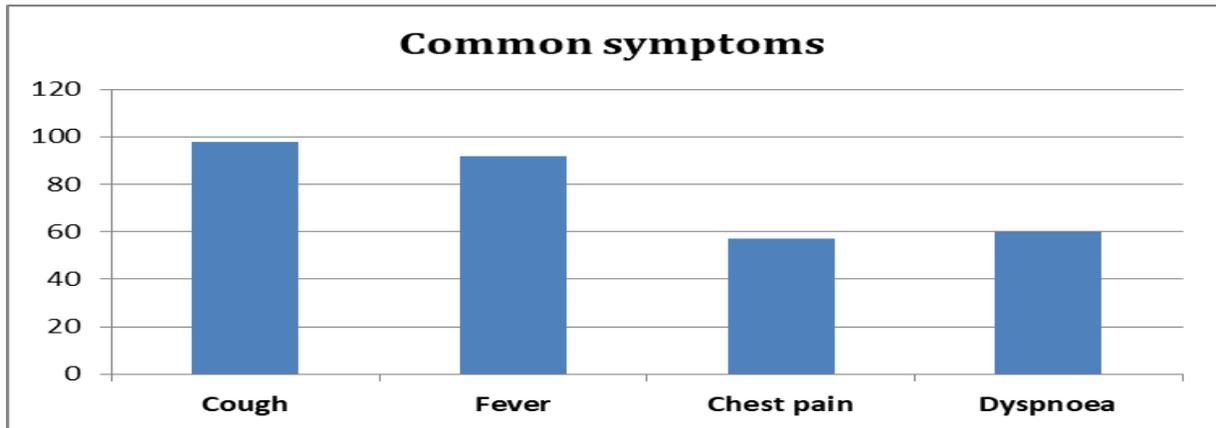


Fig.4 Associated Risk factors noted in the study

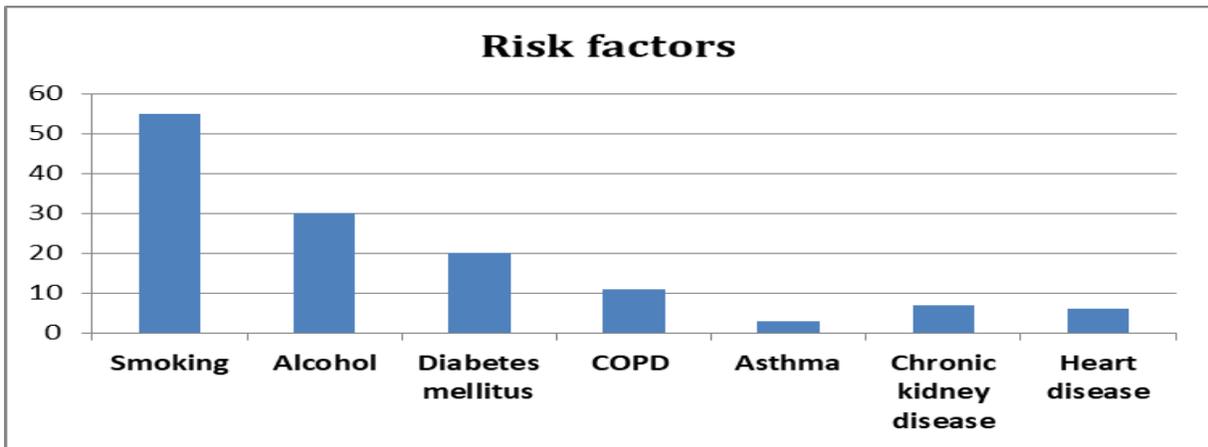


Fig.5 Culture positives in sputum

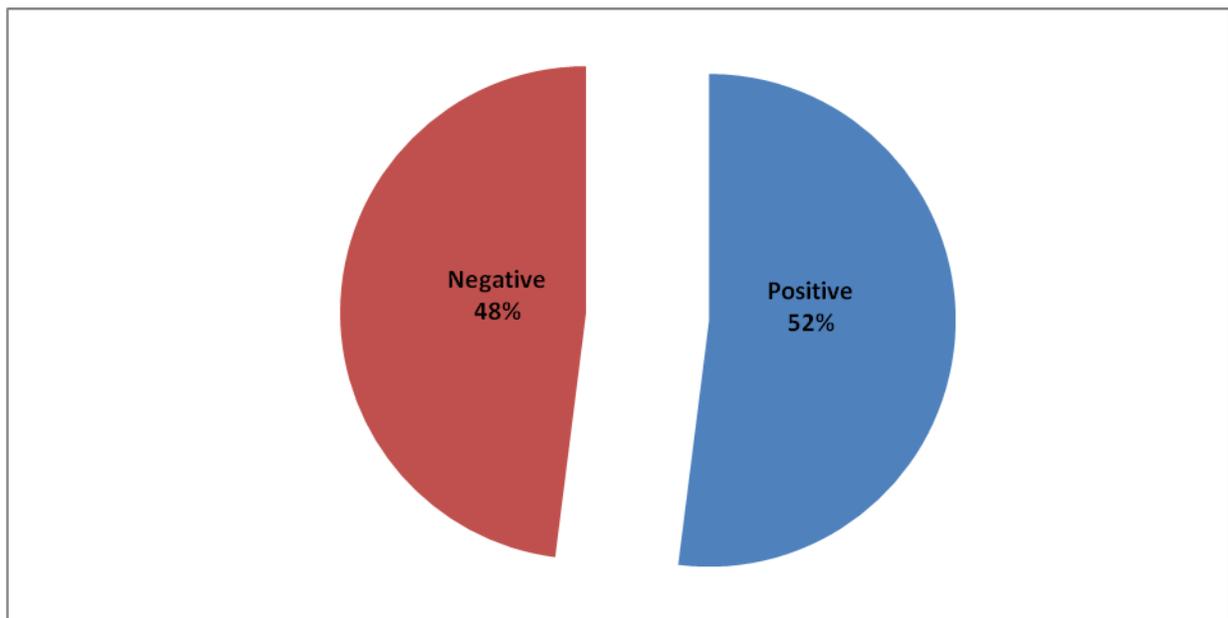


Fig.6 Total no of isolates in sputum culture

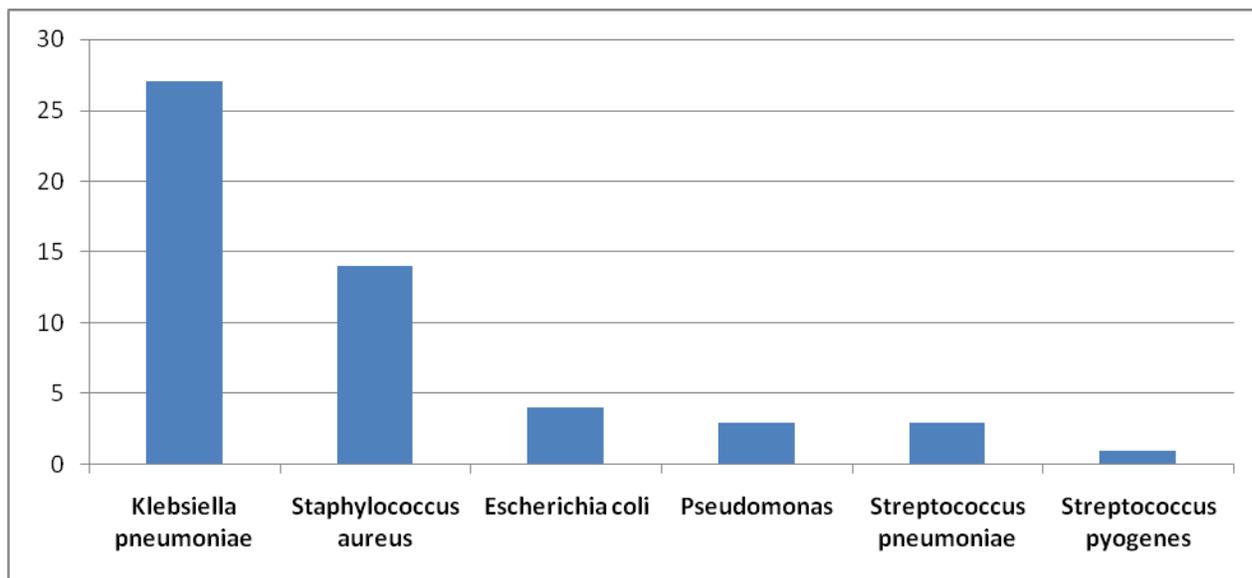
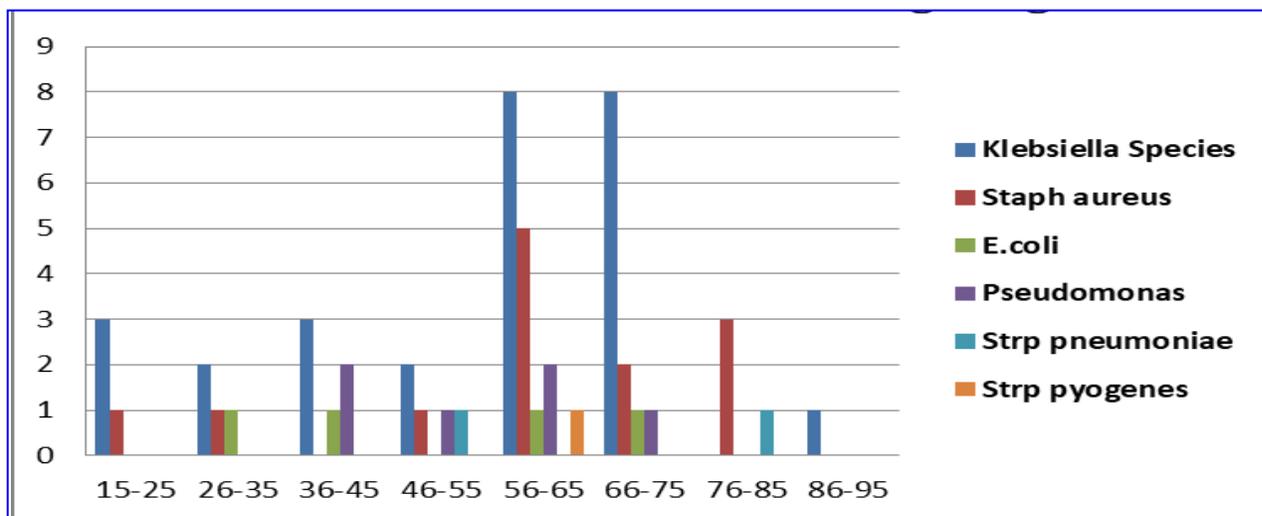


Fig.7 Distribution of isolates according to age



In the present study, sputum culture positivity was 52% and *Klebsiella pneumoniae* was most common pathogen isolated which correlated with Mythri *et al.*, High isolation of 77% culture positivity was reported by Sunil Vijay (2016) *Staphylococcus aureus* was second most common organism isolated in the present study which correlates with Sunil Vijay *et al.*, (2016). Whereas only 2.6% of *Staphylococcus aureus* was reported by

Madhulata *et al.*, (2013) and Mythri *et al.*, (2013).

In the present study aetiology remained unknown in 48% cases, which correlates with previous study, according to which, even with use of extensive laboratory testing and various invasive procedures etiological confirmation could be achieved in 45-70% according to studies conducted by Arabinca *et*

al., (2002) and Ewing S *et al.*, (2002). Though *Streptococcus pneumoniae* have been reported as the commonest organisms causing community acquired pneumonia, Indian studies over the last three decades have reported higher incidence of Gram negative organisms among culture positive pneumonia as per study conducted by Brown JS (2009). Increased incidence of *Klebsiella pneumoniae* may reflect the effects of different environmental conditions on transmission and host factors such as abnormal nutritional status, comorbidities or genetic background (Fig. 5, 6 and 7).

In the present study *Klebsiella pneumoniae* was the major pathogen. Majority (60%) of patients was above 45 years of age and habituated to smoking, or had COPD. Old age, smoking and underlying respiratory diseases such as COPD impair pulmonary defences and predispose to CAP caused by gram negative bacteria. Our hospital being a tertiary referral hospital, we receive community acquired pneumonia patients with wide range of severity, many of them carrying multiple co morbidities. These patients might have been exposed to antibiotics for treatment of respiratory or non-respiratory tract infections.

Summary

Males constitute a major proportion of patients affected by CA-Pneumonia.

People in the age group of 45-65 years were more affected by CAP.

The common risk factor observed was Smoking followed by Alcoholism and Diabetes mellitus.

Sputum culture was positive in 52% of patients.

Klebsiella pneumoniae (51.9%) was the most common organism isolated. Other Gram negative bacteria isolated were *Escherichia coli* (7.6%) and *Pseudomonas aeruginosa* (5.7%).

Among Gram positive cocci isolated, *Staphylococcus aureus* (26.9%) was the most common organism followed by *Streptococcus pneumoniae* (5.7%) and *Streptococcus pyogenes* (1.9%).

In conclusion, the present study was undertaken to know the prevalence of etiological microorganism of CAP and their antimicrobial susceptibility pattern, so that specific treatment can be advocated. Out of the 100 patients included in the study, 71 were males and 29 were females. Positive sputum culture was obtained in 52% and the major pathogen isolated was *Klebsiella pneumoniae* (51.9%) followed by *Staphylococcus aureus* (26.9%).

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How to cite this article:

Sarah Firdous and S. Jaya Prakash Rao. 2019. Study of Bacterial Isolates in Community Acquired Pneumonia. *Int.J.Curr.Microbiol.App.Sci.* 8(01): 644-654.
doi: <https://doi.org/10.20546/ijcmas.2019.801.072>