

A Pilot Study on Aetiology of Acute Lower Respiratory Tract Infections Among Children Hospitalized of Respiratory Illness at a Rural Hospital in South Coastal Karnataka

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Aetiological diagnosis can significantly impact the clinical management and outcome of acute lower respiratory tract infections (LRTI) in children. There is a paucity of data on etiological agents of acute LRTI among children in Karnataka, especially in Udupi district. Present study provides an insight into the pathogens associated with acute LRTI among children in Udupi district of south coastal Karnataka. A cross sectional study was performed at a rural hospital in south coastal Karnataka, A total of 50 children clinically diagnosed for acute LRTI and admitted in paediatric ward were enrolled for the study. Nasopharyngeal/throat swab specimens were collected, and nucleic acid was extracted, and Multiplex real-time PCR was performed for detection of bacterial and viral aetiology. *S. pneumoniae* was detected in 16% (8/50), followed by ? Respiratory syncytial virus (RSV) 14% (7/50), *H. influenzae* 8 % (4/50) and *M. pneumoniae* 2% (1/50). Mixed infection was detected in 28% (14/50) of children. *S. pneumoniae* and *H. influenzae* was the most prevalent co-infection and was detected in 10% (5/50) followed by *H. Influenzae* and RSV (4%, 2/50) co-infection. *S. pneumoniae* and RSV were the most predominant bacterial and viral pathogens respectively associated with LRTIs among paediatric population in present study. Further we found very high number of cases with mixed infections which signifies the urgent need of much elaborate studies for elucidating the clinical significance of these infections as well as for better understanding of epidemiology of LRTI among children in this region.

Keywords: Haemophilus influenzae; Influenza Virus A;
Respiratory Syncytial Virus; Streptococcus pneumoniae.

The burden of respiratory tract infections among the paediatric population in India is growing faster day by day due to polluted environment, poor lifestyle, and inadequate care. Respiratory tract infections account for the highest number

of deaths among children every year in majority of developing countries like India. Pneumonia accounts for 17% of all deaths among the children below 5 years of age in India¹. Further 30-40 % of hospital admissions in developing countries are due

to lower respiratory tract infections (LRTI)². Thus, LRTI remains one of the major healthcare issues worldwide. Due to involvement of a wide range of pathogens, lack of specific clinical presentation, and inability of conventional diagnostic methods to provide timely and efficient diagnosis; establishing the aetiology of LRTI always remains a challenge. Recently developed molecular methods such as polymerase chain reaction (PCR) and real time PCR offer great promise as future diagnostics for respiratory tract infection providing rapid identification of pathogens with high sensitivity and specificity. It also permits identification of more than one pathogen in the same respiratory specimen that has previously been difficult to culture³. Aetiological diagnosis can significantly impact the clinical management and outcome of acute LRTI in children. There is a paucity of data on etiological agents of acute LRTI among children in Karnataka, especially in Udupi district. Present study provides an insight into the pathogens associated with acute LRTI among children in Udupi district of south coastal Karnataka.

MATERIAL AND METHODS

Study setting

A cross sectional study was conducted from August 2017 to November 2017 at Department of Microbiology, Kasturba Medical College, Manipal, in collaboration of Dr. TMA Pai Rotary Hospital, Karkala and Manipal Centre for Virus Research, Manipal Academy of Higher Education (Deemed to be University), after obtaining approval from Institutional Research and Ethical Committee (IEC-336/2017). A total of 50 children presenting with fever and cough and clinically diagnosed for acute LRTI as per Infectious Disease Society of America (IDSA) guidelines were admitted in paediatric ward and were enrolled for the present study. Samples were collected after obtaining the written consent from the parents of the enrolled participants.

Sample collection

Two nasopharyngeal/throat swab specimens were collected from each participant using nylon flocked swabs (COPAN, Italy) and then swabs were placed in tubes containing universal transport medium and viral transport medium.

Respective nasopharyngeal swabs were transported to the laboratory maintaining cold chain.

Molecular Detection of Bacterial Aetiology

DNA was extracted from samples using a commercial kit from Qiagen (Hilden, Germany) using manufacturer's guidelines and Multiplex real-time PCR was performed for detection of *Chlamydomphila pneumoniae*, *Mycoplasma pneumoniae*, *Legionella pneumophila*, *Bordetella pertussis*, *Bordetella parapertussis*, *Streptococcus pneumoniae*, and *Haemophilus influenzae* using Allplex™ Respiratory Panel 4 kit (Seegene, Seoul Korea) as per manufacturer's guidelines.

Molecular Detection of Viral Aetiology

Total nucleic acid was extracted using MagMax kit (Ambion Inc., USA) and tested using multiplex real time PCR assay (Respiratory 21, Fast-Track Diagnostics, Luxemburg) for Influenza Virus A and B, Respiratory Syncytial Virus, Adenovirus, Enterovirus, Parainfluenza Viruses 1-4, Human Coronaviruses and Human Boca virus.

Data collection

Demographic data, immunization status and history of present illness were collected by interviewing the parents of the patients. Nutritional status was assessed using weight for age National Center for Health Statistics (NCHS) centile charts.

Statistical analysis

Details of the enrolled children was recorded in an excel spread sheet and statistical analysis was done using SPSS 16 (IBM, USA). Association between categorical variables was evaluated by Chi-squared test. A *p* value < 0.05 was considered significant.

RESULTS

A total of 50 children clinically diagnosed for LRTI and hospitalized in general ward were enrolled for the present study, 30 among them were male whereas 20 were females. Median age and weight were found to be 31.5 months (Range: 4 months- 144 months) and 10kg (Range: 3.7-27.5 kg) respectively, A detailed description of demographic details and clinical findings of the enrolled children is given in Table 1.

A total of 34 (68%) children were found to have infection either bacterial/ viral or mixed. *S. pneumoniae* was the commonest (16%; 8/50)

organism, followed by Respiratory syncytial virus (RSV) 14% (7/50), *H. influenzae* 8% (4/50) and *M. pneumoniae* 2% (1/50). Mixed infection was detected in 28% (14/50) of children. *S. pneumoniae* and *H. influenzae* was the most prevalent co-infection and was detected in 10% (5/50) followed by *H. Influenzae* and RSV (4%, 2/50) co-infection. A detailed description of the pathogen detected in the present study is given in Fig 1.

On analysing the aetiology among different age groups, we found that RSV (28.57%, 4/14) infection was commonly observed in children between age group 0-1 years, whereas *S. pneumoniae* (24.14%, 7/29) was commonest in children between 1 year to 5 years. Further, mixed infections (33.33%, 12/36) were more prevalent among children above 1 year of age. Normal duration of hospital stay was found to be 5-7

Table 1. Details of children enrolled for the study

	Positive for viral or bacterial aetiology (n=34, %)	Normal (n=16)	Odds Ratio (95% CI) <i>p</i> value
Age			
Mean (SD) in months	45.15 (38.18)	35.05 (23.04)	-
Range in months	4-144	5-72	0.25
Gender			
Male	18(36)	12(24)	0.38 (0.1,1.4)
Female	16(32)	04(08)	0.13
Weight			
Mean (SD)	12.07(6.18)	11.13 (4.98)	-
Range in Kg	3.7-27.8	5.1-19	0.58
Housing Condition			
Kuccha	26(52)	15(30)	0.22 (0.02,1.9)
Pukka	8(16)	01(02)	0.13
Nutritional Status			
Normal	17(34)	9(18)	0.78 (0.24,2.57)
Malnourished, PEM, FTT, Obese	17(34)	7(14)	.67
Immunization status			
up to date	30(60%)	14(28)	1.07 (0.17,6.56)
Due	04(08)	02(04)	0.94
Fever			
Present	32(64)	15(30)	1.07 (0.09,12.71)
Absent	02(04)	01(02)	0.96
Cough			
Present	33(66)	14(28)	4.71(0.39,56.32)
Absent	01(02)	02(04)	0.18
Blocked runny nose			
Present	30(60)	13(26)	1.73 (0.34,8.85)
Absent	04(08)	03(06)	0.50
Difficult Breathing			
Present	24(48)	13(26)	0.55 (0.13,2.38)
Absent	10(20)	03(06)	0.42
Chest Indrawing			
Present	17(34)	12(24)	0.33 (0.09,1.24)
Absent	17(34)	04(08)	0.09
X-ray findings			
Normal	24(48)	13(26)	0.55 (0.13,2.38)
Suggestive of Pneumonia/Asthma	10(20)	03(06)	0.42

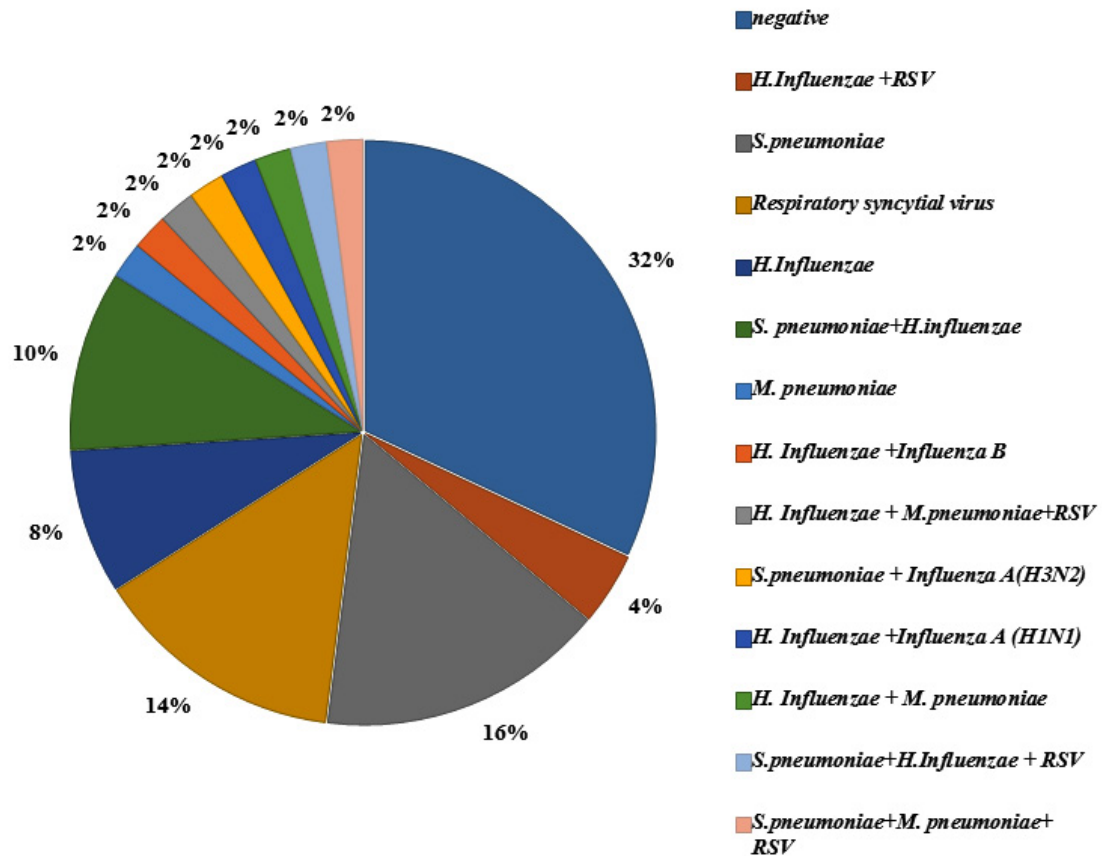


Fig. 1. Frequency of pathogens detected in the present study

days. Recovery rate was 100% as all hospitalized children with LRTI responded well to intravenous combination of beta lactams plus amino glycoside treatment.

DISCUSSION

Present study provides an insight into the infectious aetiology of lower respiratory tract infections among hospitalized children in Udupi district of south coastal Karnataka. Several studies in past have reported *S. pneumoniae* as the most common pathogen associated with LRTI in children; similar to their findings 16 % of the children in present study were also detected to have *S. pneumoniae* infection⁴⁻⁷. For PCR based diagnosis quantification of bacterial DNA load is important; a cut off value of C_T e³⁵ (e³⁵ 8000 copies/ ml) has been described previously for

distinguishing pneumococcal infection from asymptomatic colonization thus in present study a sample was considered positive only if the mean cycle threshold (C_T) value was less than 35^{8,9}.

RSV is the most frequently isolated virus in infants and children worldwide in LRTI. In the present study also, RSV remained the commonest viral agent accounting for 14% cases¹⁰⁻¹². Age dependant distribution of pathogens showed RSV (8%, 4/50) as a substantial threat in children below 1 year of age as maternal antibodies are the only means of protection during initial 6-12 months of their age¹³. Earlier studies have reported a gradual decrease in the prevalence of RSV with increase in age due to development of child's own immunity against RSV; similar pattern was also seen in the present study^{13,14}.

Mixed infections due to respiratory pathogens are reported frequently; 28% (14/50)

cases in present study were also found to have mixed infections, 12 % (6/50) among them had mixed bacterial infection. Several epidemiological studies have reported a positive correlation between colonization of *S. pneumoniae* and *H. Influenzae*, a recent study also showed increase in pneumococcal biofilm formation in presence of *H. influenzae*. In concordance to them 10% (5/50) cases in present study were also found to have co-infection of *S. pneumoniae* and *H. influenzae*¹⁵. A study showed that viral infection predisposes the host to bacterial infection by favouring bacterial attachment to the nasopharyngeal sites and promoting bacterial growth. 16% (8/50) cases in present study had both bacterial and viral mixed infections^{16, 17}.

Although few Indian studies has shown significant correlation when clinical, demographic and risk factors were compared between infected and non-infected cases of acute LRTI, no such significant correlation was seen in present study^{18,19}. The possible reason for discrepancy might be the low sample size in the present study which is one of the major limitations of this study. Further the prevalence of bacterial and viral pathogens majorly depends on the geographical area, seasonal variations and the pathogen panel included for PCR. Though we have included major pathogens responsible for acute LRTI among paediatric age group discrepancies may arise. Despite above limitations the present study provides important baseline information on aetiology of acute LRTI among paediatric population in Udupi district of south coastal Karnataka which can be useful for future researchers in formulating strategies for larger surveys or management in this region.

CONCLUSION

S. pneumoniae and RSV were the most predominant bacterial and viral pathogens respectively associated with LRTI among paediatric population in present study. Further we found very high number of cases with mixed infections which signifies the urgent need of much elaborate studies for elucidating the clinical significance of these infections as well as for better understanding of epidemiology of LRTI among children in this region.

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Conflict of interest

Authors declare no conflict of interest.

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