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### Full Length Research Article

## MOLECULAR DIAGNOSTIC VALUE OF PNEUMOCOCCAL PNEUMONIA AMONG EGYPTIAN CHILDREN

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#### ABSTRACT

Pneumonia is the most common invasive pneumococcal disease; however, only a small portion of cases can be confirmed by conventional techniques. The present study aimed at comparing the molecular detection of *Streptococcal pneumoniae* by the standard blood culture in community-acquired pneumonia among Egyptian children. Standard blood culture of *Streptococcal pneumoniae* and sequence-specific PCR amplification of autolysin gene were performed. There was a statistically significant increase in PCR diagnosed cases than that diagnosed by blood culture with higher PCR sensitivity. Also, there was a statistically significant decrease in culture diagnosed cases among those receiving empirical antibiotic treatment prior sampling. This study concluded that molecular detection of *Streptococcal pneumoniae* may be more accurate and rapid in diagnosis of pediatric community-acquired pneumonia below five years than the standard blood culture.

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#### INTRODUCTION

Pneumococcus is an important human pathogen that causes pneumonia which is the most common invasive pneumococcal disease (Johansson *et al.*, 2010). It accounts for up to 4 million deaths a year in developing countries in children below five years (Saad *et al.*, 2013); however, microbiologic confirmation of pneumococci is difficult. Diagnosis of pneumococcal Community-Acquired Pneumonia (CAP) has been made through conventional culture but yields are low (Mandell *et al.*, 2007) and available only 24–36 hours after blood collection, a delay that will be reflected in patient management (Charkaluk *et al.*, 2006). In the absence of a reliable and rapid diagnosis, empirical use of broad-spectrum antibiotics exposes the patients to adverse medication effects, and emergence of multidrug-resistant strains (Black *et al.*, 2010). Work on molecular diagnosis has been ongoing (Resti *et al.*, 2010). Autolysin (*lyt A*) gene, encoding autolysin, a well characterized virulence marker of pneumococci (Berry *et al.*, 1989), has restricted allelic variations. The *lyt A* gene therefore

makes an ideal target for specific identification of pneumococci in clinical and epidemiological studies (Llull *et al.*, 2006). So this study was conducted to compare the molecular detection of *lyt A* gene of pneumococci with the standard blood culture as a method for the etiological diagnosis of pneumococcal CAP among Egyptian children below five years.

#### MATERIALS AND METHODS

This study included 40 CAP cases admitted at the Department of Pediatrics, Benha University Hospital (with positive C-reactive protein and high total leukocyte count). Immune-compromised or lung cancer children were excluded. Approval of the Ethical Committee of the Faculty of Medicine, Benha University and informed consents from parents of children were obtained. The standard blood culture and molecular detection of pneumococci were done. The patients were exposed to full history taking, complete clinical examination and routine laboratory tests (complete blood count and CRP), culture and PCR diagnosis of pneumonia.

##### Sampling

Venous blood sample (5ml) was taken from each child; 3ml directly inserted into blood culture bottle, incubated at 37°C

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and 2ml immediately put in sterile vacutainer tube with EDTA, mixed, aliquoted (one part was used for CBC and the other was stored at -80°C for PCR).

**Blood culture**

Standard blood culture, aerobic and anaerobically subcultures were performed. *S. pneumoniae* were identified by accurate observation of its morphology, α-hemolysis of blood agar, catalase negativity, optochin susceptibility and bile solubility (Kellogg et al., 2001).

**PCR amplification of *lyt A* gene**

- Genomic DNA extraction from 400ul blood (with EDTA) using Genomic DNA purification kits (Fermentas, Germany) following the manufacturer instructions. The ratio between the absorbance values (optical densities) at 260 and 280 nm gives an estimate of DNA purity as reported by Mohamed et al. (2010).
- PCR amplification of autolysin gene using single-plex PCR in G-storm thermal cycler (England) with initial denaturation at 95°C (3min) then 35 cycles of denaturation, annealing and extension (95°C 1min, 55°C 1min and 72°C 2min respectively), then final extension at 72°C for 10min. Primers were provided by (Operon, inc Huntsville, Alabama Germany); forward primer 5'-CAA CCG TAC AGA ATG AAG CCG-3' and reverse primer 5'-TTA TTC GTG CAA TAC TCG TGC G-3'. Green PCR master mix 2x (Fermentas, Germany) was used in 50µl reaction; [25µl Taq PCR master mix 2x, 2.5µl forward primer, 2.5µl reverse primer, 5µl DNA template and nuclease free water up to 50µl]. Concentration of amplified DNA product by spectrophotometer= 50 ug/ml X A260 X dilution factor (Wilfinger et al., 1997). Bands were suspected to be seen at 308 bp.
- Gel electrophoresis of amplified products; 10µL reaction mixture and 5µL of 1000 base pair ladder were separated on 2% agarose gel with 0.3ug/ml ethidium bromide. The bands were visualized using UV transilluminator (254 nm) and imaged with a digital camera 8 mega pixel. The image was analyzed by computer software (Alpha In no Tech Gel Documentation System).

**Statistical analysis**

Qualitative data were expressed in number and percent. Appropriate statistical tests using SPSS version 8 were performed (Cohen's k test and Fisher exact probability test). ROC curve analysis was used to determine the diagnostic power of each test. Results were statistically significant at p < 0.05.

**RESULTS**

Standard blood culture and genomic PCR amplification of pneumococcal *lyt A* gene were done to determine the etiological diagnosis of pediatric CAP cases (♂:♀ 21:19). The study showed a statistically significant increase in CAP cases diagnosed by PCR (13/40) (p< 0.05), than that diagnosed by culture (3/40) (Table 1). None of the children negative for

pneumococcal infection by PCR was positive by standard blood culture (Table 1). PCR specificity was 100% compared to culture (Table 2). Area under the curve (AUC) for PCR diagnosis was 0.88 (76.5% sensitivity, 100% positive predictive value and 85.2% negative predictive value) in respect to 0.62 for culture AUC (23.5% sensitivity, 100% positive predictive value and 63.9% negative predictive value) (Table 2 and Fig.2). PCR diagnosed CAP cases were statistically increased in previous asthmatic patients (p<0.05) (Table 3). Culture positive cases were statistically decreased by empirical antibiotic treatment before diagnosis (p<0.01); however, PCR results were not affected (Table 3).

**Table 1. Standard blood culture and PCR in diagnosis of pediatric community-acquired pneumonia**

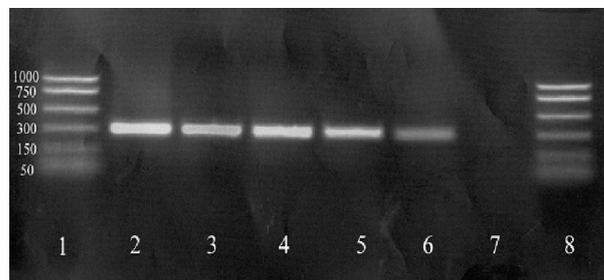
		Culture		Total	Cohen's k	P
		-ve	+ve			
PCR	-ve	n. 27	0	27	0.288	<0.01*
	%	100.0%	0.0%	100.0%		
	+ve	n. 10	3	13		
	%	76.9%	23.1%	100.0%		
Total	n.	37	3	40		
	%	92.5%	7.5%	100.0%		

-ve: negative, +ve: positive, n.: number, \*P<0.05 is significant

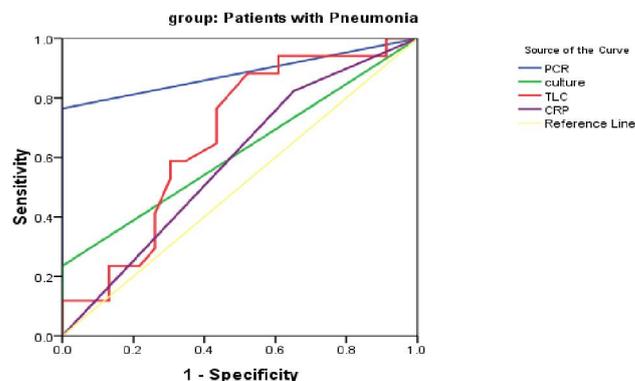
**Table 2. Comparison of different tests in diagnosis of pediatric community-acquired pneumonia**

Variable	Sensitivity	Specificity	PPV	NPV	AUC	95% CI	P
PCR (+ve)	76.5%	100.0%	100.0%	85.2%	0.88	0.76-1.00	<0.001*
Culture (+ve)	23.5%	100.0%	100.0%	63.9%	0.62	0.44-0.80	>0.05
TLC (13.7 x10 <sup>3</sup> /µl)	58.8%	69.6%	58.8%	69.6%	0.67	0.50-0.84	>0.05
CRP (72 ng/ml)	82.4%	34.8%	48.3%	72.7%	0.59	0.41-0.76	>0.05

+ve: positive, \*: P<0.05 is significant, NPV: negative predictive value, PPV: positive predictive value, AUC: area under the curve, CI: confidence interval, TLC: total leukocyte count, CRP: capsular reactive protein



**Fig. 1. Gel electrophoresis; DNA ladder in lanes 1 and 8 (its molecular weight 1000 base pairs), pneumococcal *lyt A* gene amplified products in lanes 2-6 (bands appeared at 308 bp) and a negative control in lane 7.**



**Fig. 2. ROC curve analysis for PCR, culture, total leukocyte count and C-reactive protein in diagnosis of pneumococcal pneumonia**

**Table 3. Percentage of cases with history of asthma, antibiotic treatment prior diagnosis and clinical diagnosis in relation to PCR and standard blood culture results among pneumonia cases**

		PCR			Total	Fisher's P	Culture		Total	Fisher's P
		-ve	+ve				-ve	+ve		
History of asthma	-ve	n.	25	8	33	0.027*	31	2	33	>0.05
		%	92.6%	61.5%	82.5%		83.8%	66.7%	82.5%	
	+ve	n.	2	5	7		6	1	7	
		%	7.4%	38.5%	17.5%		16.2%	33.5%	17.5%	
Antibiotic treatment prior diagnosis	-ve	n.	3	4	7	0.19	4	3	7	<0.01*
		%	11.1%	30.8%	17.5%		10.8%	100%	17.5%	
	+ve	n.	24	9	33		33	0	33	
		%	88.9%	69.2%	82.5%		89.2%	0%	82.5%	
Clinically	Lobar pneumonia	n.	3	3	6	0.37	5	1	6	>0.05
		%	50%	50%	15%		83.3%	16.7%	15%	
	Broncho-pneumonia	n.	24	10	34		32	2	34	
		%	70.6%	29.4%	85%		94.1%	5.9%	85%	

-ve: negative, +ve: positive, n.: number, \*P<0.05 is significant

## DISCUSSION

The progressive emergence of antibiotic resistance in *S. pneumoniae* requires rapid and accurate diagnosis (Lull *et al.*, 2006). Our work team chose CAP in children below five years because of more frequent episodes of respiratory tract infection with consequent disruption of normal lung structure and function, in addition to lack of immunization against *S. pneumoniae* in Egypt (Mishaan *et al.*, 2005). In the present study; there is a statistical significant increase in the number of cases diagnosed by PCR compared to culture (P<0.05) (Table 1). Target gene amplification has enabled the detection of low numbers of pathogens in clinical samples (Fredriksson-Ahomaa and Korkeala, 2003). There is also enhanced PCR specificity (100%) that is the same for culture (Table 1), as PCR-based detection depends on amplification of species-specific genes that are unique to pneumococci. Molecular assays targeting the autolysin gene do not cross-react with other streptococcal species (Blaschke, 2011). This study also shows enhanced PCR sensitivity (76.5%) that was diminished for culture (23.5%) (Table 2), this could be explained as PCR detection is not diminished with non viable organisms (McAvin *et al.*, 2001); however, culture is affected by the tendency of *S. pneumoniae* to autolyze when reaching the stationary phase of growth, antibiotic treatment prior to specimen collection and difficulty with adequate specimen collection (Blaschke, 2011). These results agreed with Peters *et al.*, (2009) who reported superiority of *lyt A* PCR over blood culture in pretreated patients.

The present study showed a statistically significant increase in PCR-diagnosed cases among patients with history of bronchial asthma (p<0.05) (Table 3); choking episodes that follow asthmatic attacks may predispose to aspiration pneumonia especially in 6-months to 3-years age (Patria and Esposito, 2013). In this study, there was a statistically significant decrease in culture results among those receiving antibiotics prior sampling; the only 3 cases who were culture positive were not receiving antibiotics prior sampling (p<0.01) (Table 3); so blood culture sensitivity was decreased by impaired micro-organism viability as explained by Ramirez and Melo-Cristino (2010). However, Zhang *et al.* (1995) attributed the decrease in culture results to sample preparation defects.

## Conclusion

Molecular detection of *S. pneumoniae* from blood of children with CAP may be more valuable in early rapid etiological

diagnosis of pneumococcal pneumonia. PCR was not affected by antibiotic treatment prior diagnosis or susceptibility to autolyze, but culture was.

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