



DETECTION OF HLY-A GENE ENCODING ALPHA HEMOLYSIN IN URINARY ISOLATES OF ESCHERICHIA COLI FROM TERTIARY CARE HOSPITAL IN KANCHEEPURAM DIST

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ARTICLE INFO

Article History:

Received 20th December, 2016

Received in revised form 16th January, 2017

Accepted 6th February, 2017

Published online 28th March, 2017

Key words:

hlyA gene, alpha hemolysin, Escherichia coli, PCR, Urinary Tract Infections

ABSTRACT

Alpha hemolysin is a toxin secreted by Escherichia coli causes cell death by binding with the outer membrane, with subsequent oligomerization of the toxin monomer and water-filled channels. These are responsible for osmotic phenomena, cell depolarization, and loss of vital molecules, leading to its demise. A total of 20 clinical isolates of E. coli were screened for the presence of hlyA gene by PCR. We have observed 60% positivity among our isolates. There are several virulent factors that are associated with urinary tract infections, production of alpha hemolysin one amongst those factors also play a crucial role in pathogenesis of UTI.

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INTRODUCTION

Escherichia coli causes different infections that possess numerous virulence factors, including hemolysin production. *E. coli* -hemolysin (HlyA) produces large, clear zones of hemolysis around colonies on blood agar. The hemolysin is present in cell-free filtrates and is the best characterized member of the RTX (repeat in toxin) toxin family^[1]. HlyA lyses cells by the creating pores in the target cell membrane and affects erythrocytes, leukocytes^[3], and renal tubular cells^[2]. It also acts on polymorphonuclear granulocytes liberates leukotrienes, histamine, and ATP^[4] and is neutralized by specific antiserum.

Sublytic concentrations of this toxin induce various reactions in eukaryotic target cells which lead to cellular dysfunction^[5]. The hly operon required for synthesis and extracellular secretion of E. coli hemolysin which contains four structural genes arranged in the order hlyC, hlyA, hlyB, and hlyD^[6]. Gene hlyA encodes 110-kDa hemolysin protein (pro-HlyA) which represents an inactive precursor of the mature toxin. The conversion of pro-HlyA to the hemolytically active hemolysin (HlyA) takes place in the cytoplasm of E. coli and is mediated by HlyC gene^[2]. With this background our aims to detect the presence of hlyA gene among urinary isolates of E. coli.

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MATERIALS AND METHODS

Bacterial isolates

A total of 20 non repetitive urinary isolates of *Escherichia coli* were collected from Saveetha Medical College and Hospitals, Chennai. They were processed for a battery of standard biochemical tests and confirmed. Isolates were preserved in semisolid trypticase soy broth stock and were stored at 4 °C until further use.

Antibiotic susceptibility testing

Antibiotic susceptibility test was determined for these isolates to routinely used antibiotics such as ampicillin, amoxicillin, amikacin, norfloxacin, ceftazimide, cefotaxime, ciprofloxacin and gentamicin, imipenem as by Kirby Bauer disc diffusion method^[7].

Detection of hlyA gene in E.coli

Escherichia coli isolates were detected for the presence of hlyA gene by PCR analysis. Detection of the gene was carried out using primer as depicted in table 1. Bacterial DNA was extracted by boiling lysis method. 1 µL of DNA extract was used as template for PCR reaction. The reaction mixture contained 1mM of MgCl₂ 0.2mM dNTP mix and 0.8µM of hlyA gene with 0.5U of Taq polymerase (New England Biolabs) in a 1x PCR buffered reaction. A positive control of E.coli with hlyA gene was also included in this study. PCR amplification was carried out using thermal cycler (Eppendorf) with the following cycling condition. Initial denaturation at 98°C for 5 min and 30 cycles for 30s, 70°C for

30s and 68° C for 60s, followed by a final extension of 6 min at 75°C. PCR products were resolved in 2% agarose gel. A 100bp ladder was including in all the gel analysis^[8].

Table 1 Gene sequencing of hlyA gene

Primer	Primer sequence	Product size
hlyA	AAC AAG GAT AAG CAC TGT TCT GGC T ACC ATA TAA GCG GTC ATT CCC GTC A	172 bp

RESULTS

Sample wise distribution of clinical isolates of E.coli

Of the 20 clinical isolates of E.coli, 12/20 (60%) were from acute urinary tract infections and 8/20 (40%) were from chronic urinary tract infections. Figure 1 depicts the sample wise distribution of clinical isolates of E.coli.

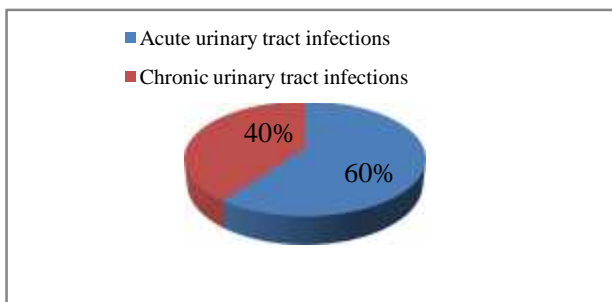


Figure 1 Sample wise distribution of urinary isolates of E.coli

Antibiotic susceptibility testing

In our isolates, we have found increased percentage 14/20 (70%) of isolates showed sensitivity to amikacin followed by gentamicin, which showed sensitivity of 9/20 (45%). 80- 90% of E.coli isolates showed resistance to cephalosporin group of drugs. 6/20 (30%) were found to be resistant to imipenem. However, we have observed an elevated level of resistance to other routinely used antibiotics. The detailed resistant pattern of E.coli isolates is shown in table 2.

Table 2 Showing antibiotic sensitivity pattern of E. coli

Antibiotics	Sensitivity(20) (%)	Intermediate (20) (%)	Resistant(20) (%)
Ampicillin	5	0	95
Amoxicillin	5	0	95
Ceftazidime	10	10	80
Cefotaxime	5	5	90
Amikacin	70	10	20
Gentamicin	45	20	35
Norfloxacin	15	15	70
Ciprofloxacin	20	5	75
Imipenem	70	0	30

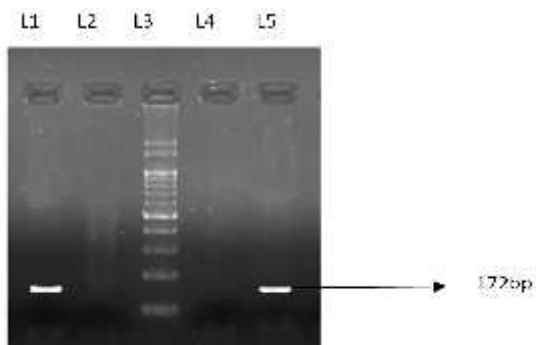


Figure 2 Representative gel picture showing positive for hlyA gene

Result of hlyA gene in E.coli

12/20 (60%) clinical isolates of urinary isolates of E.coli was found to harbor hlyA gene.

DISCUSSION

The most frequently occurring extraintestinal infections caused by E. coli are usually urinary tract infections. The -hemolysin is present in about 25 to 56% of isolated strains from urinary tract infections^[9], similar to the previous studies we also got increased percentage of E. coli isolates from urinary tract. Study done by Kerenyi and colleagues found that, the presence of the sheA gene in normal fecal strains is higher (85.4%) than in the isolates from urinary tract infection (47.1%). Wherein, the occurrence of hlyA gene in normal fecal isolates was 7.3%, similar to that previously reported^[10]. Similarly we also found 12/20 (60%) of our uropathogenic E.coli were showed positive for hlyA gene. Though it is acknowledged that the -hemolysin is an important virulence factor to the pathogenic profile of E. coli, the role played by the silent hemolysin PSheA in disease is still unknown.

CONCLUSION

From our study we have observed the presence of hlyA gene encodes for alpha hemolysin production. There are several virulent factors are associated with urinary tract infections, production of alpha hemolysin also plays a crucial role in pathogenesis of UTI. To conclude this more number of isolates with different sets of genes for other hemolysins needs to be detected.

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Please cite this article in press as:

Reshma Harikrishnan and Gopinath. P (2017), Detection Of Hly-A Gene Encoding Alpha Hemolysin In Urinary Isolates Of Escherichia Coli From Tertiary Care Hospital In Kancheepuram Dist, *International Journal of Current Advanced Research*, 6(3), pp. 2654-2656.
<http://dx.doi.org/10.24327/ijcar.2017.2656.0067>
