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Identification and Antibiotic Susceptibility Testing of *Klebsiella* species from various Clinical specimens in a tertiary care hospital

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ABSTRACT

Klebsiella are Gram-negative, shorter and thicker rod shaped bacilli. On Mac-Conkey agar it shows mucoid colonies due to presence of polysaccharide capsule. Total 50 isolates of Klebsiella species collected from different clinical samples were studied biochemically. The different samples were urine 19 (38%), pus 15 (30%), blood 11 (22%), and sputum 5 (10%). Antibiotic susceptibility tests were done by Kirby Bauer disk diffusion method and the ESBL detection by double disc synergy method. *Klebsiellapneumoniae* were the most common 48 (96%), followed by Klebsiellaoxytoca 2 (4%) in our study. Antibiotic resistance to ampicillin were 100%, percentages to cephalosporins were high: cefozolin and cefuroxime 30 (60%), cefotaxime 27 (54%) and cefoperazone and ceftazidime were 28 (56%). Most of those 21 (42%) were from the infection associated in the hospital. Resistance to Aminoglycosides ranged from 60 to 70%. Resistance to co-trimoxazole and tetracycline were 25 (50%) and 27 (54%) respectively. Resistance to fluoroquinolones were 28 (56%). Resistance to carbapenem (meropenem) were 20 (40%) and to piperacillin sulbactam were 21 (42%). Resistance to third generation cephalosporins were 15 (30%) out of which 9 (60%) showed ESBL production using double disc synergy method

Keywords: Urinary tract infection, Community acquired pneumonia, ESBL, AST

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INTRODUCTION

Antimicrobial resistance has become now days a serious public health problem worldwide. Infections caused by resistant bacteria have increased morbidity and mortality than those caused by susceptible pathogens^{1,2}. Infections caused by resistant bacteria led to prolonged hospital stays, increased health care costs and in many cases to untreatable infections³. *Klebsiella* has a prominent capsule which is responsible for the mucoid appearance of the isolated colonies and enhances the virulence of the organism in-vivo. *Klebsiella pneumoniae* is the most important *Klebsiella* species from a medical stand point and most commonly isolated from the clinical specimens. Pneumonia caused by *Klebsiella* is frequently associated with necrotic destruction of the alveolar space, causing cavity formation and production of blood stained sputum⁴. It is also associated with long term care facilities causing nosocomial infections. *Klebsiella* also cause wound, soft tissue infection and urinary tract infection.

Beta-lactams antibiotics are the most frequently prescribed antibiotics. Emerging resistance to these antibiotics among gram negative bacilli limited their utility. Extended spectrum [beta]-lactamases (ESBL) producing *Klebsiella* spp. has been frequently implicated in outbreaks in intensive care units and neonatal intensive care units (NICUs)⁵. *Klebsiella pneumonia* are ubiquitously present and reported worldwide. In recent years, *K. pneumonia* has become important pathogens in nosocomial infections. The importance of *K. pneumonia* species in the ever increasing number of gram negative aerobic bacillary nosocomial infections in the United States ⁶ and India ⁷ has been well documented. Over 100 variations and point mutations in TEM gene were reported by DNA sequencing. These mutations are the most responsible factor for resistance to beta lactams in these isolates. [The SHV family of [beta]-lactamases has been derived from *Klebsiella* spp. SHV-1 is universally found in *K. pneumoniae*, evolved as a chromosomal gene in *Klebsiella* spp. and was later incorporated into a plasmid, which has spread to other entero-bacterial species.⁸. A total of 40 types of SHV type ESBL enzymes are already reported.

There are various reports on the prevalence of ESBL producing Klebsiella spp. from India. A recent study by Khan Erum et al describes occurrence of genetic variants in *K. pneumoniae* from clinical samples of various origin and reported that isolates having both TEM and SHV genes were more common than TEM and SHV alone.

Epidemic strains of cephalosporin resistant *K. pneumoniae* have been associated with increased morbidity and mortality in hospitalized patients ⁹. However, most *Klebsiellae* infections now occur in the hospital. Antibiotic – resistant strains have been responsible for a number of outbreaks of nosocomial infections in ICUS and neonatal nurseries .The most common clinical syndromes are pneumonia, UTI, abdominal infection, surgical site infection,

soft tissue infection and subsequent bacteremia. *K. pneumoniae* subspecies rhinoscleromatis is the causative agent of *rhinoscleroma* – a granulomatous, slowly progressive (over months to years) infection of the upper respiratory mucosa that causes necrosis and occasional obstruction of the nasal passages. *K.pneumoniae* subspecies ozenae has been implicated in chronic atrophic rhinitis and in rare cases of invasive disease in compromised hosts.

Hence, the present study was undertaken to isolate and identify *Klebsiella* spp. from various clinical samples and to know their antibiotic susceptibility pattern in an urban tertiary care centre.

MATERIALS AND METHOD:

A total of 50 isolates of *Klebsiella* species from various clinical samples received in the department of microbiology St. Johns medical college and hospital (SJMCH) were studied during the period Jan 2010 to Dec. 2010. The strains were sub cultured on MacConkey agar to obtain pure growth. Subsequently each isolate were preserved in nutrient agar deeps for further studies. Presumptively identified *Klebsiella* species with significant growth and pure culture were included while *Klebsiella* species with insignificant /mixed growth were excluded,

The presumptively identified *Klebsiella* species isolated from various clinical samples, biochemically were identified to the genus and species level as per the recommended standards. Antibiotic susceptibility testing of the isolates was performed by Kirby-Bauer disc diffusion method. The antibiotic concentration of disc used and zone size interpretation was in accordance with clinical laboratory standards institute (CLSI)¹⁰. Different antibiotics used were as Ampicillin, Piperacillin, Cefazolin, Cefuroxime, Cefotaxime, Cefaperazone, Cetazidime, Gentamicin, Netilmycin, Amikacin, Cotrimoxazole, Tetracycline, Ciprofloxacin, Meropenem and Piperacillin + Tazobactam

Muller Hinton agar media was prepared, autoclaved, allowed to cool in 50°C & poured on to Petri dishes to give a uniform depth of 4mm. 3-5 colonies of strain was inoculated into 2ml peptone water and incubated at 37°C for 4hrs. Then turbidity was adjusted to 0.5 Mac farlandstandard(1.5X10 cfu/m) with sterile normal saline.

Plates were inoculated within 15 min of preparation of the suspension so that the density did not change .A sterile cotton wool swab was dipped in to the suspension and surplus removed by rotation of the swab against the side of the tube above the fluid level .The medium was inoculated by even streaking of the swab over the entire surface of the plates in three directions. After the inoculums had dried, discs were applied with forceps and pressed gently to ensure even contact with the medium.

Controls:

ATCC Escherichia coli 25922 (American type culture collection) was used

ESBL DETECTION:

The strains found to be resistant to 3rd generation cephalosporins were done ESBL detection.

Method Used:

Double disc synergy method¹¹.

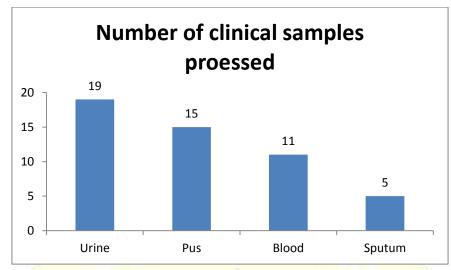
PROCEDURE:

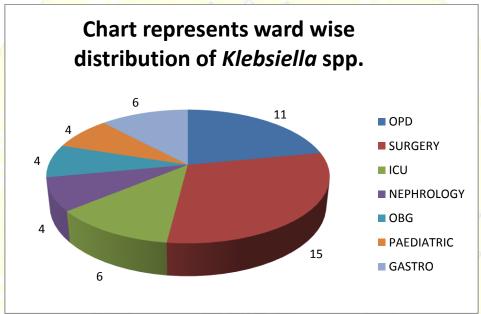
All the strains screened for resistance to third generation cephalosporins using three discs cefotaxime, ceftriaxone and ceftazidime by Kirby Bauer disc diffusion method. Those strains showing resistance to at least two of them (zone of inhibition less than 18 mm) were tested for ESBL — in duplicate by the Kirby Bauer dic diffusion in Mueller Hinton agar plates by Jarlier technique. In this technique, susceptibility disk containing amoxicillin and clavulanic acid was placed at the centre and cefotaxime, ceftriaxone and ceftazidime disks were placed at 15 mm distance (centre to centre) on a lawn culture of the organism using turbidity standard of 0.5 McFarland. An increased zone of inhibition towards the clavulanic acid disk was taken as positive for ESBL.

RESULTS AND DISCUSSION

Klebsiella pneumoniae were the most common 48 (96%), followed by Klebsiella oxytoca 2 (4%) in our study. Antibiotic resistance to ampicillin were 100%, percentages to cephalosporins were high: cefozolin and cefuroxime 30 (60%), cefotaxime 27 (54%) and cefoperazone and ceftazidime were 28 (56%). Most of those 21 (42%) were from the infection associated in the hospital. Resistance to Aminoglycosides ranged from 60 to 70%. Resistance to co-trimoxazole and tetracycline were 25 (50%) and 27 (54%) respectively. Resistance to fluoroquinolones were 28 (56%). Resistance to carbapenem (meropenem) were 20 (40%) and to piperacillin –sulbactam were 21 (42%).

Resistance to third generation cephalosporins were 15 (30%) out of which 9 (60%) showed ESBL production using double disc synergy method. Confirmation of ESBL production in all those strains using molecular method could not be done due to lack of infra-structure.





| Antibiotics | No. Of isolates found resistance | Percentage of resistance |
|---------------|----------------------------------|--------------------------|
| Ampicillin | 50 | 100 |
| Piperacillin | 17 | 34 |
| Cefazolin | 30 | 60 |
| Cefuroxime | 30 | 60 |
| Cefotaxime | 27 | 54 |
| Cefaperazone | 28 | 56 |
| Ceftazidime | 28 | 56 |
| Gentamicin | 30 | 60 |
| Netilmycin | 35 | 70 |
| Amikacin | 30 | 60 |
| Cotrimoxazole | 25 | 50 |
| Tetracycline | 27 | 54 |
| Ciprofloxacin | 28 | 56 |
| Meropenem | 20 | 40 |
| Zosyn | 21 | 42 |

Total Number Of Isolates Resistant To All 2 Or 3 Cephalosporin=15

ESBL DETECTION:

METHOD USED: double disc synergy method.

ESBL was detected in 9/15 strains (60%)

Klebsiella pneumoniae is considered an important pathogen causing nosocomial and community-acquired infections. It is often associated with the production of extended spectrum β-lactamases (ESBL) belonging to SHV and CTX-M families. These genes are frequently described as a part of complex integrons facilitating their horizontal transfer to other related as well as unrelated microbes. The present study was undertaken for the identification and antibiotic susceptibility testing of Klebsiella spp. from various specimens in a tertiary care hospital. ESBL producing isolates of Klebsiella pneumonia have been reported from various hospitals in India and other countries ¹². The need for ESBL production is under challenge to set the break point for injectable cephalosporins and aztreonam that accurately discriminate which ESBL-producing isolates can and cannot be reliably treated with these drugs.

The majority of our isolates were acquired from urine(38 %), followed by pus (30%), then blood (22%) and sputum (10%) signifying the true infection. 6 isolates (12%) were from the ICU putting them at high risk for ventilator associated pneumonia and 10?(%) showed bacteremia with gram negative rods. 4?(%) cases showed community acquired bronchopneumonia associated with chronic obstructive lesions. Out of 50 strains, 48 (96%) were due to the *Klebsiella pneumoniae* and 2 strains were *K. oxytoca*. Both those *K. oxytoca* strains were isolated as nosocomial pathogens, one from the pus wound and the other from the urine. Both the strains though similar to the *K. pneumoniae* biochemically, but formed indole and fermented dulcitol.

CONCLUSION:

Klebsiella isolates have been steadily increasing over the past years and they have been important source of transferable drug resistance. Indiscriminate use of third generation cephalosporins to treat gram negative bacterial infections is partly responsible for the emergence of resistance to beta-lactam antibiotics. Strict adherence to the hospital antibiotic policy and good infection control practices can play a significant role in reducing the increasing drug resistance. Epidemiological surveillance studies should provide useful information base to guide practice and policies on rational use of anti-infective agents and to eradicate the source of environmental.

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