

Superbugs causing infections at a tertiary care hospital and the return of pre-antibiotic era!

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Submitted: 15-06-2015

Revised: 05-10-2015

Published: 10-11-2015

ABSTRACT

Aims and Objectives: Increase in the incidence of *Escherichia coli* and *Klebsiella pneumoniae* carrying New Delhi metallo beta lactamase-1 (NDM-1) gene are called superbugs is of great concern as presence of *bla*_{NDM-1} gene makes *E.coli* and *K.pneumoniae* highly resistant to most of currently available antibiotics. This study was planned to observe the burden of *bla*_{NDM-1} gene carrying *E. coli* and *K.pneumoniae* at a tertiary care hospital in northern India. **Materials and Methods:** A total of 1709 *E. coli* and 327 *K. pneumoniae* nonrepetive isolates were taken from various clinical samples received in a tertiary care hospital in northern India Lucknow during the period from May 2012 to April 2013. Carbapenemase production was phenotypically detected in all the carbapenem resistant isolates by modified Hodge test. Metallo- β -lactamase production was detected by using meropenem and imipenem discs with and without EDTA and *bla*_{NDM-1} gene was detected by polymerase chain reaction. **Results:** Over all metallo β - lactamase production was found in 82% and 88.89% of carbapenem resistant *E.coli* and *K. pneumoniae* respectively. Out of 366 carbapenem resistant isolates, 102 were found positive for *bla*_{NDM-1} gene out of which 89 were *E.coli* and 13 were *K. pneumoniae*. **Conclusions:** With limited treatment options left for this crisis situation like the pre-antibiotic era; it is an alarm for rational antibiotic therapy usage and intensive education programs.

Key words: New Delhi metallo beta lactamase-1, *Escherichia coli*, *Klebsiella pneumoniae*. Metallo- β -lactamase

Access this article online

Website:

<http://nepjol.info/index.php/AJMS>

DOI: 10.3126/ajms.v7i2.12750

E-ISSN: 2091-0576

P-ISSN: 2467-9100

INTRODUCTION

During the last two decades, appearance of β -lactam resistance specially to 3rd and 4th generation cephalosporin in members of enterobacteriaceae family arose as a serious problem to global health.¹ Currently carbapenem are the therapeutic drug of choice for treating serious infections caused by *Escherichia coli* and *Klebsiella pneumoniae*.² One of the common mechanism of carbapenem resistance in *E.coli* and *K. pneumoniae* is production of carbapenemase enzyme. Recently a novel β -lactamase NDM-1 has been reported in multidrug resistant *E.coli* and *K.pneumoniae*. NDM-1 gene in Enterobacteriaceae isolates were reported from patients in India, Pakistan, Bangladesh and the UK in 2008–2009.³ NDM-1 gene is being reported worldwide, however exact burden of NDM-1 gene in world is not known as very few

studies has been done on NDM-1 gene and epidemiological data has not been known so far.

It is necessary to detect these strains as soon as possible and suitable infection control interventions should be implemented for all patients infected or colonized with MBL producing bacteria.

MATERIAL AND METHODS

Study isolates

A total of 1709 *E.coli* and 327 *K.pneumoniae* isolates were taken from the culture of 20,674 various clinical samples (urine, pus, blood, cerebrospinal fluid, endotracheal aspirate, catheter tips etc) received in the bacteriology laboratory of the Microbiology department during one year period from

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May 2012 to April 2013. All the carbapenem resistant isolates included in the study were further identified by following accepted laboratory criteria.⁴

Antimicrobial susceptibility testing

Antimicrobial susceptibility against isolated *E. coli* and *K. pneumoniae* for various antimicrobial agents using beta-lactam and non-beta lactam antibiotic discs were determined by Kirby-Bauer disc diffusion method and results were interpreted as per Clinical and Laboratory Standards Institute (CLSI) guidelines.⁵ Antibiotic discs were procured from Hi-Media, Mumbai, India, except imipenem, which was from BD Diagnostics, Franklin Lakes, NJ, USA. EUCAST guideline 2012 suggest MIC method for colistin drug susceptibility.⁶ Reference grade powder for imipenem and meropenem were obtained from Lac Zene Biosciences, Delhi, India. Minimum inhibitory concentration (MIC) values for imipenem, meropenem and colistin were determined by agar dilution method. Results were interpreted as standard protocol. All the isolates with reduced susceptibility to meropenem and/or imipenem on disc diffusion testing and/or MIC were screened for carbapenemase production. Phenotypic detection for carbapenemase production was done by modified hodge test.⁵ *K. pneumoniae* ATCC BAA-1705 as positive control and *K. pneumoniae* ATCC BAA-1706 as negative controls were used for MHT.

MBL production was detected in isolates with reduced susceptibility to imipenem and/or meropenem using combined disc diffusion test by imipenem (10 µg) and 0.1 M ethylenediaminetetraacetic acid (EDTA: 292 µg) as described previously by Franklin et al.⁷ This was complemented by the Modified Hodge test and the double disc synergy test with imipenem (10 µg) alone and imipenem (10 µg)+EDTA (750 µg) for all the isolates resistant to any carbapenem as performed earlier by Lee et al.⁸

Molecular detection of *bla*_{NDM-1} gene-

Template DNA from meropenem and/or imipenem resistant strain was prepared from freshly cultured by re-suspending 3-5 colonies in 100 µL of molecular grade water, and then by heating at 95°C for 10 minutes. The bacterial lysate was centrifuged at 10,000 rpm for 2 min, the supernatant was transferred to new eppendorf tube and used as DNA template for PCR amplification. *K. pneumoniae* ATCC BAA-2146 procured from Eurofins Genomics, MWG Biotech AG primer microbiologics KWIK STIC was used as positive control for NDM-1 gene. The resistance gene was amplified by polymerase chain reaction (PCR) by using previously published primers NDM-Fm 5'-GGTTTGGCGATCTGGTTTTC-3' and NDM-Rm-5'-CGGAATGGCTCATCACGATC-3'.⁹⁻¹¹ specific for *bla*_{NDM-1} in selected meropenem and/or

imipenem resistant strains.¹² The Amplified product was analysed on 2% (w/v) agarose gel containing (0.5 µl/ml) of ethidium bromide along with molecular weight marker (100 bp) electrophoretically. Constant current (80V) was maintained for 2 hr and analysed by 264 nm wavelength UV transilluminator and gel doc (alphamager3400HP) was used for documentation. A sharp band 621 bp of amplified DNA visualized, considered to be positive (Figure 1).

RESULTS

Out of 1709 *E. coli* and 327 *K. pneumoniae* isolates, 330 *E. coli* and 36 *K. pneumoniae* isolates were resistant to carbapenem. The patients suffering from these carbapenem resistant isolates having various risk factors [Table 1]. Percentage of *bla*_{NDM-1} positive isolates among carbapenem resistant *E. coli* and *K. pneumoniae* were 26.97%(89/330) & 36.11%(13/36) respectively. Majority of patients having infection with *bla*_{NDM-1} gene containing isolates were in 15-60 years of age group. Majority of the *bla*_{NDM-1} gene positive isolates were obtained from ICU and wards. All *bla*_{NDM-1} producing *E. coli* and *K. pneumoniae* were susceptible to colistin on MIC however only 53.9% *E. coli* and 61.5% *K. pneumoniae* were susceptible to tigecycline. For other antimicrobials, *bla*_{NDM-1} positive isolates showed variable susceptibility pattern [Table 2]. Out of total 89 *bla*_{NDM-1} positive *E. coli*, only 1 isolate was found moderately sensitive to meropenem on MIC, however it was resistant to imipenem. All isolates of *K. pneumoniae* with *bla*_{NDM-1} positive were resistant to both meropenem and imipenem by MIC method. Range of MIC value for meropenem, imipenem and colistin were 2-16 µg/ml, 4-16 µg/ml and 0.5-1 µg/ml respectively for *E. coli* and 4-16 µg/ml, 4-8 µg/ml and 0.5-1 µg/ml respectively for *K. pneumoniae*.

DISCUSSION

Enterobacteriaceae members are common causes of both health care and community acquired infections, raising the possibility of spread of resistant organisms

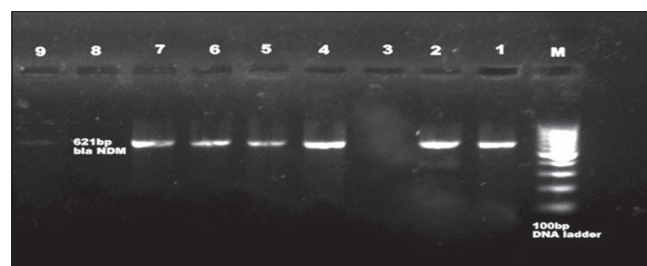


Figure 1: Representative gel picture showing amplification of *bla*_{NDM-1} gene (621bp). Lane M: 100 bp DNA ladder; Lanes 1 and 2: Positive control *Klebsiella pneumoniae* ATCC BAA-2146; Lane 3: Negative control; Lane 4-9: Clinical isolates

Table 1: Risk factors associated with carbapenems resistant isolates of *E. coli* and *K. pneumoniae*

Risk factors	CREC* (n=330)		CRPK** (n=36)	
	Number of patients	Percentage	Number of patients	Percentage
ICU stay >48 hrs	61	18.49	5	13.89
Duration of hospital stay >3 days	190	57.58	24	66.67
Exposure to antimicrobials within last 14 days				
BL+	197	59.70	19	52.78
AG++	206	62.42	24	66.67
FQ+++	162	49.1	8	22.22
AA++++	104	31.52	4	11.11
Presence of devices	192	58.18	23	63.89
• Shunt	33	10	6	16.67
• Prosthesis	6	1.82	4	11.11
• I V catheter	39	11.82	3	8.33
• ICD tube	15	4.55	0	0
• ET tube	46	13.94	4	11.11
• Tracheal tube	14	4.24	1	2.78
• Urinary catheter	53	16.06	7	19.44
• Drain	6	1.82	1	2.78
• Stent	10	3.03	0	0

+Betalactams, ++Aminoglycosides, +++Fluoroquinolones, ++++All antibiotics, *Carbapenem resistant *E. coli*, **Carbapenem resistant *K. pneumoniae*, #Intercostal drainage tube, ##Endotracheal tube

Table 2: Antibiotic resistant pattern of NDM-1 *Escherichia coli* and *Klebsiella pneumoniae*

Antibiotics	<i>Escherichia coli</i> (n=89)	<i>Klebsiella pneumoniae</i> (n=13)
Piperacillin/tazobactam	87 (97.75)	10 (76.9)
Ceftriaxone	89 (100)	12 (92.3)
Ceftazidime	89 (100)	12 (92.3)
Amikacin	72 (80.1)	11 (84.6)
Ciprofloxacin	81 (91)	11 (84.6)
Aztreonam	76 (85.4)	11 (84.6)
Meropenem	89 (100)	13 (100)
Imipenem	85 (95.5)	12 (92.3)
Ertapenem	89 (100)	12 (92.3)
Tigecycline	41 (46.1)	5 (38.5)

into the community.¹³ Our study gives an initial account on the presence of *bla*_{NDM-1} gene in a tertiary care hospital in northern India. Amongst all carbapenem resistant study isolates, *bla*_{NDM-1} gene was present in 89 isolates of *E. coli* (26.97%) and 13 *K.pneumoniae*. In a study done by Kumarasamy et al., in Chennai (India), *bla*_{NDM-1} gene was found in 25.33% *E.coli* and 23.33%⁹ isolates of *K.pneumoniae* in carbapenem resistant isolates. Over all *bla*_{NDM-1} gene was found in 5% of isolated *E.coli* and *K. pneumoniae*, showed almost similar finding with previous study done by Seema et al in which NDM-1 gene was found in 5.39% of total isolated *E.coli* and *K.pneumoniae*.¹⁴ Out of 102 *bla*_{NDM-1} positive isolates 76 (74.5%) were found in hospitalized patients. In our study, only 16 (15.68%) *bla*_{NDM-1} positive isolates were recovered from ICU patients. while in most of previous Indian studies showed that *bla*_{NDM-1} gene positive isolates are found predominantly in ICU patients however some of previous Indian studies have shown greater isolation from non-ICU wards.^{15,16} Eight isolates carrying *bla*_{NDM-1} gene were recovered from patients on ventilator.

All NDM-1-producing isolates of *E. coli* and *K. pneumoniae* were susceptible to colistin on MIC while in a study done by Kumarasamy et al., colistin showed variable sensitivity pattern to NDM-1 gene isolates.⁵ Reason may be different geographical distribution. Susceptible to tigecycline which are almost consistent with a study done by Seema et al, 2011¹⁴ in which tigecycline susceptibility was present in 46.2% and Kumarasamy et al, 2010³ in which it was 64% and 67%. In our study some of the *bla*_{NDM-1} harbouring isolates were found phenotypically susceptible to one or more carbapenems tested.

So all the carbapenem disc should be used for drug susceptibility with, and if possible molecular analysis should be done in hospital setting because false susceptibility testing may lead to treatment failure.

CONCLUSION

Present study shows higher rate of resistance to commonly used carbapenems in a tertiary care set up in northern India which may be due to irrational use of antibiotics and responsible genes can transfer from one bacteria to other. It is necessary to detect these strains as soon as possible and suitable infection control interventions should be implemented for all patients infected or colonized with MBL producing bacteria.

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Authors Contribution:

All the authors were equally involved in drafting and finalizing the manuscript. **RK** - Concept and design of study, statistically data analysis, supervised research work, manuscript preparation and editing, finalised manuscript; **SR** - Concept, collected data and literature search, did experimental work, helped in data analysis, helped in manuscript preparation and editing; **JA** - Concept of study, helped in molecular work, critical revision and editing of manuscript; **MS** - Concept of study, helped in molecular work, manuscript editing and review.

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Source of Support: Nil, **Conflict of Interest:** None.