

## A ten year analysis of multi-drug resistant blood stream infections caused by *Escherichia coli* & *Klebsiella pneumoniae* in a tertiary care hospital

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Received October 11, 2010

**Background & objectives:** Extensive use of antibiotics has added to the escalation of antibiotic resistance. This study was undertaken to evaluate the association, if any between antibiotic use and resistance in a hospital setting, and also detect the predominant mechanism of antibiotic resistance in *Escherichia coli* and *Klebsiella pneumoniae* over a period of 10 years.

**Methods:** In a retrospective study of 10 years, a total of 77,618 blood culture samples from 2000 to 2009 from indoor patients were screened and those yielding *E. coli* and *K. pneumoniae* were included in the study. Antibiotic susceptibility records as well as the percentage of ESBL producers were noted. A total of 423 isolates of 2009 were also screened for AmpC and carbapenemase production. Antibiotic consumption data of 10 years were analysed.

**Results:** ESBL producing *E. coli* increased from 40 per cent in 2002 to 61 per cent in 2009, similarly there was a significant ( $P<0.05$ ) rise in resistance to cefotaxime (75 to 97%), piperacillin-tazobactam (55- 84%) and carbapenem (2.4-52%) in *K. pneumoniae*. A significant ( $P<0.05$ ) association was observed between resistance and consumption of carbapenem and piperacillin and tazobactam consumption in *K. pneumoniae*.

**Interpretation & conclusions:** Our study demonstrated a rise in consumption and resistance to broad spectrum antimicrobial agents and also established an association between consumption and resistance to these antibiotics. Over a period of 10 years, the emergence of pan-resistance in *K. pneumoniae* could be due to the production of carbapenemases whereas ESBL production was the common mechanism of resistance in *E. coli*. This study warrants a directed effort towards continued surveillance and antibiotic stewardship to minimize selection pressure and spread.

**Key words** Antibiotic resistance - blood stream infection - *Escherichia coli* - *Klebsiella pneumoniae*

The emergence of antibiotic resistance is a global public health problem. Gram-negative bacterial resistance is of particular importance as there is a dearth of novel antibiotics directed against these organisms. The clinical utility of carbapenems, the

agents of last resort against multi-drug resistant *Enterobacteriaceae* is under threat with the growing incidence of pan resistant isolates<sup>1,2</sup>. Studies on the trends of antimicrobial consumption and resistance as well as the common mechanisms of resistance of

multi-drug resistant *Escherichia coli* and *Klebsiella pneumoniae* are scanty. Moreover, the organisms encountered in various hospitals around the country is far from uniform<sup>3,4</sup>. The emergence of multi-drug resistant and pan-resistant *E. coli* and *K. pneumoniae* and the lack of consistent data prompted us to conduct this study to demonstrate a relationship if any, between antimicrobial use and resistance over a period of ten years with reference to the common mechanism of resistance in these isolates in a tertiary care hospital in north India.

### Material & Methods

This retrospective study was undertaken in a 650 bedded tertiary care hospital with an active transplant programme at New Delhi, India, to evaluate the emergence of antibiotic resistance in *E. coli* and *K. pneumoniae* over a period of ten years (2000-2009). The antibiotic consumption rate was also analysed. A total of 77,618 consecutive blood culture samples from in-patients admitted during the study period were evaluated and those yielding *E. coli* and *K. pneumoniae* were included in the study. All repeat isolates were excluded using speed-miner Hospital Information System (HIS) (Australia).

**Blood culture isolates and susceptibility testing:** BactT Alert 3D (bioMerieux, France) automated system was used to culture all the blood samples received. Blood culture bottles were incubated for seven days before being reported negative. Identification and antibiotic susceptibility testing of all these isolates were done using routine biochemical tests and standard Kirby-Bauer method for antibiotic susceptibility<sup>5</sup> and subsequently from 2002 using VITEK 1 and VITEK - II from 2005 (bioMerieux, France) and interpreted as per CLSI guidelines<sup>6</sup>. Blood culture samples yielding *E. coli* and *K. pneumoniae* were only included in the study. *E. coli* and *K. pneumoniae* were screened and confirmed for extended spectrum  $\beta$ -lactamase (ESBL) activity as per CLSI guidelines<sup>6</sup>. Besides the above method, antibiotic susceptibility was done subsequently by Vitek-2 which also confirmed the presence of ESBL.

**Carbapenemase detection:** All isolates of *E. coli* and *K. pneumoniae* in the year 2009 (423 isolates) were tested for the presence of carbapenemases using the modified Hodge test (MHT) as per the CLSI guidelines<sup>7</sup>. *K. pneumoniae* ATCC<sup>®</sup> BAA-1705 and ATCC<sup>®</sup> BAA-1706 were used as positive and negative controls, respectively. Since MHT is a phenotypic screen for the

presence of carbapenemase and does not differentiate between Class A KPC (*K. pneumoniae* carbapenemase) and Class B MBL (metallo- $\beta$ -lactamases), which are encountered in these isolates, all isolates were also screened for MBL production by using imipenem - EDTA combined disc diffusion test and MBL E-test (AB Biodisk, Solna, Sweden) method<sup>8</sup>.

**AmpC screening:** Since 2009 onwards all isolates were screened for Amp C  $\beta$ -lactamases. Though there are no CLSI guidelines for their detection, screening for AmpC was done using AmpC disc test method. This is a reliable, sensitive and economically viable method for diagnosing AmpC production in the laboratory<sup>9</sup>. All 423 isolates of 2009 were screened for AmpC production.

**Antibiotic consumption data:** The monthly antibiotic issued to the inpatients was obtained from HIS (Hospital Information System). The consumption of the antibiotics was finally expressed as number of DDDs (daily defined dosages)/100 days<sup>10</sup>. The following antibiotics were studied using the Anatomical Therapeutic Chemical (ATC) classification: cefotaxime (JOIDD01), amikacin (JOIGB06), ciprofloxacin (JOIMA02), piperacillin-tazobactam (JOICR05) and imipenem (JOIDH51) representing the third generation cephalosporins (3GC), aminoglycosides, fluoroquinolones,  $\beta$ -lactam and  $\beta$ -lactamase combination and carbapenems, respectively<sup>10</sup>. Antibiotic resistance data were analysed using a FoxPro based indigenously designed program till 2006 and after that speedminer program of HIS was used. The data were analysed in relation with antimicrobial resistance over a period of 10 years.

**Statistical analysis:** The data for each antibiotic prescription and resistance were first analyzed individually for studying the trend over time using the function  $Y=ae^{bt}$ . Taking log on both sides,  $\log(y) = \log(a) + b t$ , where b stands for the percentage growth rate per year. The growth rate was tested for its significance using t-test. The relationship between antibiotic resistance and consumption was further established using linear regression by fitting the linear function  $Y = a + bx$ , where x is the antibiotic consumption and y is the antibiotic resistance. Here b indicates the change in antibiotic resistance for each unit change in antibiotic consumption.

### Results

A total of 77,618 blood samples were screened over a period of 10 years, of which 13153 samples (16.9%)

rate of both *K. pneumoniae* and *E. coli* bacteraemia over a period of 10 years was 2.59 and 17.6 per cent, respectively which was statistically significant ( $P<0.05$ ) (Table I). *E. coli* was the predominant organism till 2006 and was subsequently replaced by *K. pneumoniae* (Table I).

Years	2000	2001	2002	2003	2004	2005	2006	2007	2008	2009	Rate of change (%)	R <sup>2</sup>	P value
Trends													
Isolates/1000 patient days													
<i>E.coli</i>	0.53	0.53	0.49	0.65	0.59	0.68	0.57	0.54	0.66	0.95	2.59	0.46	0.032
<i>K. pneumoniae</i>	0.14	0.57	0.48	0.735	0.424	0.62	0.51	0.68	1.6	1.4	17.6	0.63	0.006
Number of blood samples (N)	4380	3660	3385	4533	10700	12413	10865	5062	8608	14,012	12.9	0.510	0.020
Antibiotic consumption (DDD/100 BD)													
Total antibiotics	201.2	210.8	209.2	207.2	222.7	307.1	262.6	212.4	318.5	226.5	3.1	0.30	0.099
Cefotaxime (J01DD)	66.7	43.3	37.5	29.4	31.7	42.8	43.6	41.5	35	19	-6.5	0.37	0.062
Aminoglycosides (J01GB)	25	35.3	18.1	19.6	12.6	21.5	22.1	17.1	28.5	30.1	0.50	0.00	0.897
Fluoroquinolones (J01MA)	30	47	58.1	61.1	71.1	91.9	67.4	44.2	47.8	27.5	-0.90	0.005	0.843
Piperacillin+Tazobactam (J01CR05)			17.5	15.9	16.3	18	26.3	18.8	61.4	44	17.80	0.67	0.014
Imepenem (J01DH51)			1.1	5.2	5.7	5.3	8.2	7	11.6	9.9	24.30	0.67	0.013
<i>K. pneumoniae</i> (% resistance)													
Cefotaxime	75	80	94	89	81	89	98	95	95	97	2.4	0.61	0.008
Amikacin	70	70	76	90	77	77	89	76	59	45	-3.20	0.22	0.17
Ciprofloxacin	64	80	64	83	82	66	81	86	64	84	1.3	0.09	0.411
Piperacillin+tazobactam			55	71	51	64	67	60	88	84	5.40	0.49	0.055
Carbapenems			2.4	1.7	0	0	7	3	47	52	45.30	0.70	0.039
ESBL			38	35	42	48	46	43	44	40	1.8	0.19	0.293
<i>E. coli</i> (% resistance)													
Cefotaxime	64	67	73	71	79	77	88	83	75	83	2.70	0.65	0.005
Amikacin	57	63	68	87	75	69	58	67	57	14	-8.90	0.282	0.115
Ciprofloxacin	53	62	84	90	49	86	89	95	98	91	5.60	0.439	0.037
Piperacillin+tazobactam			53	67	47	37	30	27	10	42	-15.5	0.42	0.08
Carbapenem			3	2	0	0	2	6	10	6	19.30	0.54	0.096
ESBL			40	45	58	60	75	78	61	61	6.7	0.51	0.046

The consumption of the antibiotics is expressed as number of DDDs (daily defined dosages) / 100 bed days as per the guidelines for ATC classifications and DDD assignment by WHO Collaborating Centre for Drug Statistics and Methodology<sup>10</sup>

**Table II.** Relationship of antibiotic consumption and antibiotic resistance by linear regression analysis

Antibiotic resistance (Y)	Antibiotic consumption (X)							
	<i>K. pneumoniae</i>				<i>E. coli</i>			
	R <sup>2</sup>	b	P value	95% CI	R <sup>2</sup>	b	P value	95% CI
Fluoroquinolones	0.004	-0.021	0.867	-0.301, 0.259	0.005	0.048	0.846	-0.502, 0.598
3 <sup>rd</sup> gen cephalosprin	0.263	-0.146	0.129	-0.346, 0.053	0.206	-0.140	0.187	0.363, 0.084
BL-BLI	0.724	0.305	0.007	0.117, 0.493	0.489	-0.765	0.053	-1.545, 0.016
Carbapenem	0.560	1.127	0.041	-0.750, 2.004	0.163	0.344	0.427	-0.738, 1.426
Aminoglycoside	0.298	-0.365	0.102	-0.822, 0.091	0.235	-0.811	0.156	-2.005, 0.383
BL-BLI, beta lactam - beta lactam inhibitor								

In *K. pneumoniae* there was a significant increase in resistance to cefotaxime, carbapenems and piperacillin-tazobactam over the years. On the other hand, significant antibiotic resistance in *E. coli* was observed to cefotaxime and ciprofloxacin (Table I). Tigecycline was introduced in the hospital formulary from 2007 and in the same year a resistance of 14 per cent was observed which further increased to 20 per cent in 2009 in *K. pneumoniae*. However, in *E. coli* the resistance to tigecycline remained 1.7 per cent in 2008 and increased marginally to 3 per cent in 2009. All the isolates remained sensitive to colistin.

There was a significant ( $P<0.05$ ) rise in ESBL producers in *E. coli* from 40 to 61 per cent over a period of 10 years and the rate of change was 6.7 per cent. No such change was observed in *K. pneumoniae* during the study period. However, a decrease in ESBL producers in *K. pneumoniae* was noticed from 2006 onwards though not statistically significant (Table I).

Due to the rise in carbapenem resistance, all the isolates of *K. pneumoniae* and *E. coli* were screened for carbapenemase and Amp C production from 2009 to determine the mechanism of resistance. A total of 423 isolates (167 *E. coli* and 256 *K. pneumoniae*) were screened. More than half of the total 256 isolates of *K. pneumoniae* (130 isolates, 51%) were MHT positive thus carbapenemase producers. Of these, carbapenemase producers in *K. pneumoniae*, 103 (79%) were MBL producers with MBL E-test and imipenem - EDTA combined disc test positive. In *E. coli*, of the total of 167 isolates, 25(15%) were MHT positive and thus carbapenemase producers. Of these, seven (28%) were MBL producers and were MBL E-test and combined disc test positive. The remaining 18 (72%)

were all class A KPC producers *i.e.* these were only MHT positive and MBL E-test and combined disc test negative and were resistant to carbapenems. All the 423 isolates of 2009 were screened for AmpC production and 8 per cent of the isolates of both *K. pneumoniae* (20) and *E. coli* (13) were AmpC positive.

Though the total antibiotic consumption did not show a significant change over a period of 10 years but there was a significant increase in prescription of broad spectrum antibiotics like piperacillin and tazobactam (JOICR05) from 17.5 to 44 DDD/100 days and imipenem (JOIDH51) from 1.1 to 9.9 DDD/100 days ( $P<0.05$ ) at the rate of 17.8 and 24.3 per cent, in 10 years, respectively (Table I).

Table II demonstrates the linear regression analysis used to establish the strength of the relationship between consumption and its resistance. There was a significant association ( $P<0.05$ ) between rise in antibiotic resistance and increase use of piperacillin-tazobactam combination (JOICR05) and carbapenem (JOIDH51) in *K. pneumoniae*.

## Discussion

Our study highlights the extensive consumption of broad spectrum antimicrobial agents coupled with high resistance rates to these agents among the isolates surveyed. There was a significant rise in ESBL isolates with an increase in 3GC and quinolone resistance in *E. coli*. In *K. pneumoniae*, an increase in resistance to 3GC, piperacillin-tazobactam and carbapenem was observed. Increase in the incidence of multi-drug resistant *E. coli* has been reported in various studies<sup>11-13</sup>. In the SMART study conducted in Asia Pacific region, the ESBL rate in India amongst *E. coli* was also alarmingly high (79%)<sup>14</sup>.



A multi-centric study conducted in 2004-2006 reported the highest ESBL rates in *K. pneumoniae* from India (72%)<sup>15</sup>. In this study, the ESBL producers in *K. pneumoniae* also increased to a high of 74 per cent till 2005 but has shown a fall after that. Emergence of other mechanisms of resistance like AmpC and carbapenemase producers may have contributed to this phenomenon. It is a known fact that AmpC  $\beta$ -lactamases and carbapenemases when present along with ESBL may mask the phenotype of the latter<sup>16,17</sup>.

The increasing resistance to piperacillin-tazobactam combination and carbapenems seen in *K. pneumoniae* in the study period may be attributed to presence of carbapenemase producing isolates. Though AmpC producers along with porin loss may contribute to carbapenem resistance also but in our experience 8 per cent of *E. coli* and *K. pneumoniae* that were AmpC producers were all carbapenem sensitive. Other studies from India also have reported AmpC prevalence of 8-43 per cent<sup>16</sup>.

Carbapenemases both class A and class B (KPC and MBL) have shown a worldwide dissemination and probably are the major contributors of carbapenem resistance<sup>18</sup>. These KPCs and MBLs were the main contributors of pan-resistance in our study. MBL producers amongst *K. pneumoniae* isolates is on the rise worldwide and in India<sup>19</sup>. In this study, it was interesting to note that though there was an increase in ESBL producers amongst *E. coli*, the antibiotic pressure due to increased use of piperacillin-tazobactam and carbapenems did not highlight pan-resistance in *E. coli* as was seen in *K. pneumoniae*. This phenomenon is difficult to explain. Probably a preferential selection of pan-resistant *K. pneumoniae* could have occurred as a result of fitness cost under antibiotic pressure. Further genotypic evaluation is needed to know whether the high prevalence of MBL in *K. pneumoniae* is due to the presence of bla<sub>NDM-1</sub> which is now on the rise<sup>19</sup>. In a pilot study conducted at our hospital MDR *K. pneumoniae* were found to be blaNDM-1 positive using PCR method (unpublished).

Though there is a paucity of data on trends of antibiotic consumption from India, the available data suggest that it is higher than other developing nations of the world<sup>20,21</sup>. Rates are further lower in developed nations<sup>22</sup>. Use of carbapenem and piperacillin-tazobactam has increased significantly in 10 years. A study done by our group earlier (1995-2001) showed a significant increase in consumption of 3GC<sup>23</sup>. Rise in 3GC resistance due to emergence of ESBL producers

resulted in a shift to usage of piperacillin-tazobactam and carbapenems causing significant increasing consumption of these broad spectrum antibiotics in our study. Such increasing trend of carbapenem and piperacillin-tazobactam use was also seen in various studies conducted in hospitals world over<sup>11,24,25</sup>.

The increase in antibiotic resistance is due to several factors but the major cause appears to be excessive use of antibiotics<sup>26</sup>. Our study showed a significant association between increased use and resistance to carbapenems and piperacillin and tazobactam combination in *K. pneumoniae* isolates as has been shown earlier<sup>11,27</sup>.

Screening for carbapenemase activity by the laboratory is the first step towards early detection and continued surveillance of these pan-resistant isolates.

There are several limitations of this work. As this study is retrospective in nature, potential confounders such as changes in length of stay, shift of patients from the ICU to the ward and vice versa, case stratification, time of collection of blood samples could not be ascertained while observing the trends. Hence cases could not be differentiated from hospital acquired infections and otherwise. Genotypic identification of the common mechanisms of resistance in these MDR isolates of *Enterobacteriaceae* was also not done.

In conclusion, our study highlights alarming increase in resistance and antibiotic use and the emergence of MDR isolates amongst *E. coli* and *K. pneumoniae*, and emphasizes on prompt remedial actions to salvage the situation. Reducing consumption by judicious use is the first intervention in the direction of antibiotic surveillance. There is an urgent need for early detection of these isolates for better treatment outcomes.

### Acknowledgment

The authors thank Prof. Peter M. Hawkey, Professor of Clinical & Public Health Bacteriology and Honorary Consultant Microbiologist, Health Protection Agency, Heart of England NHS Trust UK, for his valuable suggestions, and Ms. Parul Takkar Chugh for statistical analysis.

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