

# Detection of *bla*<sub>NDM-1</sub> and *bla*<sub>NDM-5</sub> genes among Gram-negative bacteria isolated from human immunodeficiency virus patients in South India

Sir,

New Delhi metallo-β-lactamase (MBL)-1 (NDM-1) gene was first detected in extensively drug-resistant (XDR) *Klebsiella pneumoniae* from a Swedish patient of Indian origin.<sup>[1]</sup> Thereafter, NDM-1 emerged as a leading threat to the treatment of infections caused by Enterobacteriaceae. In this study, we aimed to study the positivity of *bla*<sub>NDM-1</sub> among Gram-negative bacteria (GNB) isolated from human immunodeficiency virus (HIV) patients attending YR Gaitonde Centre, Chennai, India. Antibiotic susceptibility of bacterial isolates was tested using Kirby–Bauer disc diffusion method.<sup>[2]</sup> Bacterial DNA extracted by boiling lysis method was used as template in polymerase chain reaction to detect the drug-resistant genes such as *bla*<sub>NDM-1</sub>, *bla*<sub>NDM-5</sub>,<sup>[3]</sup> extended-spectrum β-lactamases, Class 1 integron, Class 2 integron, sulfamethoxazole (*sul*), and trimethoprim (*dfr*). In the present study, 45.1% (78/173) of GNB isolated from HIV patients showed resistance to imipenem (IPM) which was highly noted among *Escherichia coli* (73.1%; *n* = 57). In the E-test, 66.7% of IPM-resistant isolates were positive to MBL production. Among MBL producers, 17.3% (*n* = 9; *P* < 0.001) showed positive for *bla*<sub>NDM-1</sub> gene, and among *bla*<sub>NDM-1</sub> isolates, 77.7% showed XDR profile and 22.2% multidrug-resistant (MDR) profile. Phylogenetic analysis using Molecular Evolutionary Genetics Analysis Version 7.0 (The Pennsylvania State University, University Park, Pennsylvania, United States) revealed that 8 *bla*<sub>NDM-1</sub> (KU695556) gene sequences had fallen into *bla*<sub>NDM-1</sub> clad. One *bla*<sub>NDM-1</sub> fell into *bla*<sub>NDM-5</sub> (KU695557) clad due to two amino acid substitutions such as valine instead of leucine (Leu) in

the 88<sup>th</sup> position and methionine instead of Leu in the 154<sup>th</sup> position. *bla*<sub>NDM</sub> positive isolates also exhibited co-positivity to other drug-resistant genes [Table 1].

Vignesh *et al.* in 2008 reported that IPM is the drug of choice against MDR bacteria and also that only 28% of GNB from HIV patients were resistant to IPM.<sup>[4]</sup> We found that 45% of GNB were resistant to IPM which indicates that IPM resistance rate has been increasing among HIV population. Another study demonstrated clonal similarity between *bla*<sub>NDM-1</sub> strains and difference in antibiotic resistance profiles based on 1–5 amino acid substitutions.<sup>[1]</sup> In our study, *bla*<sub>NDM</sub> isolates were found clonally different by random amplified polymorphic DNA analysis. In a study from Ecuador, *bla*<sub>NDM</sub>-positive *K. pneumoniae* from HIV patients exhibited co-positivity to *bla*<sub>CTX-M</sub> and *bla*<sub>SHV</sub> genes.<sup>[5]</sup> Our study reports first time the positivity of *bla*<sub>NDM-1</sub> and its variant *bla*<sub>NDM-5</sub> among GNB from HIV patients in South India. Continuous monitoring of *bla*<sub>NDM</sub> genes among GNB is needed due to XDR and MDR profiles which could help in the timely treatment of bacterial infections in HIV patients.

## Acknowledgments

This study was approved by the Institutional Review Board of YR Gaitonde Centre for AIDS Research and Education (approval number: YRG 209A), Voluntary Health Services Hospital Campus, Chennai, India.

## Financial support and sponsorship

Nil.

## Conflicts of interest

There are no conflicts of interest.

**Marimuthu Ragavan Rameshkumar<sup>1,2</sup>,  
Narasingam Arunagirinathan<sup>1,3</sup>, Purushothaman Indu<sup>1</sup>,  
Chinnambedu Ravichandran Swathirajan<sup>2</sup>,  
Sunil Suhas Solomon<sup>2,4</sup>, Ramachandran Vignesh<sup>2,5</sup>,  
Pachamuthu Balakrishnan<sup>2</sup>**

<sup>1</sup>Department of Microbiology and Biotechnology, Presidency College (Autonomous), Chennai, Tamil Nadu, India, <sup>2</sup>Infectious Diseases Laboratory, YR Gaitonde Centre for AIDS Research and Education, Voluntary Health Services Hospital Campus, Chennai, Tamil Nadu, India, <sup>3</sup>Central Research Laboratory, Meenakshi Academy of Higher Education and Research (Deemed to be University), Chennai, Tamil Nadu, India, <sup>4</sup>Department of Medicine, Johns Hopkins University School of Medicine, Baltimore, MD, USA, <sup>5</sup>Preclinical Department, Faculty of Medicine, Universiti Kuala Lumpur Royal College of Medicine Perak (UNIKL RCMP), Ipoh, Malaysia

**Table 1: Positivity of *bla*<sub>NDM-1</sub> and *bla*<sub>NDM-5</sub> along with other drug-resistant genes among Gram-negative bacteria isolated from HIV patients**

| Parameters                                   | <i>bla</i> <sub>NDM</sub> gene positive isolates                         |  |   |   |  |  |   |   |  |
|--|--|--|---|---|--|--|---|---|--|
|  | 8EC  | 10EC   | 62EC  | 124EC   | 137EC  | 142EC  | 144KO   | 156EC   | 161AB  |
| <i>bla</i> <sub>NDM</sub> positivity         | <i>bla</i> <sub>NDM-1</sub>  | <i>bla</i> <sub>NDM-5</sub>  | <i>bla</i> <sub>NDM-1</sub>   | <i>bla</i> <sub>NDM-1</sub>   | <i>bla</i> <sub>NDM-1</sub>  | <i>bla</i> <sub>NDM-1</sub>  | <i>bla</i> <sub>NDM-1</sub>   | <i>bla</i> <sub>NDM-1</sub>   | <i>bla</i> <sub>NDM-1</sub>  |
| Organism                                     | <i>E. coli</i>   | <i>E. coli</i>   | <i>E. coli</i>  | <i>E. coli</i>  | <i>E. coli</i>   | <i>E. coli</i>   | <i>K. oxytoca</i>   | <i>E. coli</i>  | <i>A. baumannii</i>  |
| Sample                                       | Urine  | Pus  | Urine   | Vaginal swab  | Blood  | Urine  | Urine   | Urine   | Pus  |
| Age and sex of persons with HIV              | 52/male  | 34/female  | 56/female   | 33/female   | 50/male  | 42/male  | 47/male   | 40/male   | 46/male  |
| CD4 cell count                               | 106 cells/mm <sup>3</sup>  | 351 cells/mm <sup>3</sup>  | 47 cells/mm <sup>3</sup>  | 47 cells/mm <sup>3</sup>  | 503 cells/mm <sup>3</sup>  | 15 cells/mm <sup>3</sup>   | 06 cells/mm <sup>3</sup>  | 145 cells/mm <sup>3</sup>   | 72 cells/mm <sup>3</sup>   |
| Phenotypic detection of MBL production       |  |  |   |   |  |  |   |   |  |
| CDM  |  |  |   |   |  |  |   |   |  |
| IPM  | 17 mm  | 12 mm  | 18 mm   | 4 mm  | 0 mm   | 12 mm  | 0 mm  | 15 mm   | 13 mm  |
| IPM + EDTA                                   | 31 mm  | 28 mm  | 28 mm   | 22 mm   | 12 mm  | 27 mm  | 12 mm   | 25 mm   | 24 mm  |
| E-test (MIC)                                 |  |  |   |   |  |  |   |   |  |
| IPM  | 8 µg   | 8 µg   | 4 µg  | >64 µg  | 12 µg  | >64 µg   | 3 µg  | 12 µg   | >64 µg   |
| IPM + EDTA                                   | >265 µg  | >265 µg  | >265 µg   | >265 µg   | >265 µg  | >265 µg  | 32 µg   | >265 µg   | >265 µg  |
| Molecular screening of drug resistance genes |  |  |   |   |  |  |   |   |  |
| Positivity of ESBL genes                     | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>CTX-M</sub>                  | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>CTX-M</sub>  | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>CTX-M</sub>   | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>OXA</sub>                                       | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>CTX-M</sub> and <i>bla</i> <sub>OXA</sub>                  | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>CTX-M</sub>  | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>CTX-M</sub> and <i>bla</i> <sub>OXA</sub>   | <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>CTX-M</sub> and <i>bla</i> <sub>OXA</sub> | <i>bla</i> <sub>TEM</sub>  |
| Integrans                                    | -  | -  | -   | Class 2 integron  | Class 2 integron   | -  | Class 1 Integron  | -   | -  |
| Sulfamethoxazole resistance                  | <i>sul1</i> , <i>sul2</i>  | <i>sul2</i>  | -   | <i>sul1</i> , <i>sul2</i>   | <i>sul1</i> , <i>sul2</i>  | <i>sul1</i> , <i>sul2</i>  | <i>sul1</i> , <i>sul2</i>   | <i>sul1</i> , <i>sul2</i>   | <i>sul1</i> , <i>sul2</i>  |
| Trimethoprim Resistance                      | -  | -  | -   | -   | -  | <i>dfrA7</i>   | <i>dfrA7</i>  | <i>dfrA7</i>  | <i>dfrA7</i>   |
| Antibiotic resistance profile                |  |  |   |   |  |  |   |   |  |
| Resistance to antibiotics                    | pip, amp, ctx, caz, cpd, cpz, cro, cxm, at, dox, cip, sxt, ipm, tet, lvx | amk, gen, pip, tzp, amp, amx, ctx, caz, cpz, cpd, cro, cxm, fox, at, cip, sxt, ipm, tet, fox, at, cip, sxt, tg, chl, etp, ipm, tet, lvx, nit | pip, tzp, amp, ctx, caz, cro, cpz, at, cip, sxt, ipm, ctx, cxm, fox, at, cip, sxt, cz, fox, ofx | amk, pip, tzp, amp, amx, ctx, caz, cro, cpm, fox, at, cip, sxt, tg, chl, ipm, lvx, ofx, nit | gen, pip, tzp, amp, amx, ctx, caz, cpd, cpz, cro, cxm, fox, at, dox, chl, cip, sxt, ipm, tet, nor, lvx | gen, pip, tzp, amp, amx, ctx, caz, cpd, cpz, cxm, fox, at, dox, chl, cip, sxt, ipm, tet, lvx, ofx, nit | amk, gen, pip, tzp, amp, amx, ctx, cpd, cpz, cro, cxm, fox, at, cip, sxt, ipm, tet, lvx | amk, gen, pip, tzp, amp, amx, ctx, cpz, cro, cxm, fox, at, chl, ipm, tet, lvx, nit    | amk, pip, tzp, amp, amx, ctx, cpz, cro, cxm, fox, at, chl, ipm, tet, lvx, nit, ofx |
| Type of resistance                           | MDR  | XDR  | MDR   | XDR   | XDR  | XDR  | XDR   | XDR   | XDR  |

*E. coli*=*Escherichia coli*; *K. oxytoca*=*Klebsiella oxytoca*; *A. baumannii*=*Acinetobacter baumannii*; IPM=Imipenem, EDTA=Ethylenediaminetetraacetic Acid; MIC=Minimum Inhibitory Concentration; MDR=Multi-drug-resistant; XDR=Extensively drug-resistant; Amk=Amikacin; Amp=Ampicillin; At=Aztreonam; Cpd=Cefpodoxime; Cpz=Cefoperazone; Ctx=Cefotaxime; Fox=Cefoxitin; Caz=Ceftazidime; Cro=Ceftriaxone; Cxm=Cefuroxime; Chl=Chloramphenicol; Cip=Ciprofloxacin; Dox=Doxycycline; Gen=Gentamicin; Ipm=Imipenem; Pip=Piperacillin; Tzp=Piperacillin-tazobactam; Tet=Tetracycline; Tmp=Trimethoprim; Sxt=Trimethoprim-sulfamethoxazole; Lvx=Levofloxacin; Nit=Nitrofurantoin; Ofx=Ofloxacin; Nor=Norfloxacin; MBL=Metallo-Beta-Lactamase; CDM=Combination disk method; ESBL=Extended-Spectrum Beta-Lactamase

**Address for correspondence:** Dr. Narasingam Arunagirinathan, Department of Microbiology and Biotechnology, Presidency College (Autonomous), Chennai - 600 005, Tamil Nadu, India. E-mail: n\_arunagiri@yahoo.co.in

## REFERENCES

- Liu Z, Wang Y, Walsh TR, Liu D, Shen Z, Zhang R, *et al.* Plasmid-mediated novel *bla*<sub>NDM-17</sub> gene encoding a carbapenemase with enhanced activity in a sequence type 48 *Escherichia coli* strain. *Antimicrob Agents Chemother* 2017;61. pii: e02233-16.
- Clinical and Laboratory Standards Institute. M100-S23 Performance Standards for Antimicrobial Susceptibility Testing; Twenty-Third Informational Supplement. Wayne, PA, USA: Clinical Laboratory Standards Institute; 2013.
- Teo J, Ngan G, Balm M, Jureen R, Krishnan P, Lin R, *et al.* Molecular characterization of NDM-1 producing *Enterobacteriaceae* isolates in Singapore hospitals. *Western Pac Surveill Response J* 2012;3:19-24.
- Vignesh R, Shankar EM, Murugavel KG, Kumarasamy N, Sekar R, Irene P, *et al.* Urinary infections due to multi-drug-resistant *Escherichia coli* among persons with HIV disease at a tertiary AIDS care centre in South India. *Nephron Clin Pract* 2008;110:c55-7.
- Romero-Alvarez D, Reyes J, Quezada V, Satán C, Cevallos N,

Barrera S, *et al.* First case of New Delhi metallo- $\beta$ -lactamase in *Klebsiella pneumoniae* from Ecuador: An update for South America. *Int J Infect Dis* 2017;65:119-21.

This is an open access journal, and articles are distributed under the terms of the Creative Commons Attribution-NonCommercial-ShareAlike 4.0 License, which allows others to remix, tweak, and build upon the work non-commercially, as long as appropriate credit is given and the new creations are licensed under the identical terms.

| Access this article online   |  |
|--|--|
| Quick Response Code:<br> | Website:<br><a href="http://www.jmsjournal.net">www.jmsjournal.net</a> |
|  | DOI:<br>10.4103/jrms.JRMS_627_18                                       |

**How to cite this article:** Rameshkumar MR, Arunagirinathan N, Indu P, Swathirajan CR, Solomon SS, Vignesh R, *et al.* Detection of *bla*<sub>NDM-1</sub> and *bla*<sub>NDM-5</sub> genes among Gram-negative bacteria isolated from human immunodeficiency virus patients in South India. *J Res Med Sci* 2018;23:112.