

EMERGENCE OF CARBAPENEM RESISTANT *RAOULTELLA ORNITHINOLYTICA*
ISOLATES CO-HARBORING *BLANDM-1* AND *BLAOXA-48*Atul Khajuria*¹ and Ashok Kumar Praharaj²¹PhD Scholar, Department of Microbiology, Armed Forces Medical College, Pune 411040, India.²MD, PhD Microbiology, Professor and Head, Department of Microbiology, AIIMS, Bhubaneswar-751019 Odisha, India.

*Corresponding Author: Atul Khajuria

PhD Scholar, Department of Microbiology, Armed Forces Medical College, Pune 411040, India.

Article Received on 21/03/2019

Article Revised on 11/04/2019

Article Accepted on 02/05/2019

ABSTRACT

Objective: To examine the distribution, emergence and spread of genes encoding beta-lactamase resistance in *R.ornithinolytica* recovered from hospitalized patients in a tertiary care hospital. **Methods:** A prospective study was conducted in an 1800 bedded tertiary care centre in Pune, India from October 2013 to October 2017. A total of 13 isolates were recovered from clinical specimens of hospitalized patients admitted to the Medical and Surgical intensive care units (one isolate per patient). Polymerase chain reaction (PCR) assays and sequencing was used to determine the presence of beta-lactamase encoding genes and conjugation experiments were performed to determine the transferability. Isolate relatedness were determined by REP PCR, ERIC PCR and RAPD. **Results:** A total of 13 *R.ornithinolytica* isolates were recovered, largest proportion of specimens were from UTIs 38.5%, followed by 23% in IAIs, 15.4% in both BSIs, and SSTIs and 7.7% in RTIs, respectively. Among 13 tested isolates, 84.6% isolates showed MIC >4µg/ml against imipenem and meropenem. Of the total number of samples, males contributed 38.5% while females contributed 61.5%. Of the total samples, highest 61.5% were from ICU Surgery followed by 38.5% in ICU Medical. Majority of Carbapenem resistant *R.ornithinolytica* 45.6% were from UTIs, followed by 18.2% in both BSIs, and SSTIs, and 9% in both IAIs, and RTIs, respectively. MHT, DDST, CDST, and MBL(IP/IPI) E-test was positive in 84.6% isolates. 100% *R.ornithinolytica* isolates retained susceptible to polymixin and colistin. Conjugation experiments indicated that *bla_{NDM-1}*, *bla_{OXA-48}*, *bla_{SHV-12}*, *bla_{SHV-28}*, *bla_{CTX-M-15}*, *bla_{CTX-M-14}* were transferable via plasmid. **Conclusion:** This study highlights prevalence of *bla_{OXA-48}* & *bla_{NDM-1}* producing *R.ornithinolytica* along with other β-lactamases genes carried on a single or multiple plasmids that serve as a driving force for the horizontal spread of carbapenem resistance.

Raoultella ornithinolytica is an encapsulated Gram-negative aerobic bacillus belonging to new genus of the Enterobacteriaceae family formerly known as *Klebsiella ornithinolytica*. In 2001, *Raoultella* was excised from *Klebsiella*, the bacterium was reclassified as *Raoultella* based on genetic approaches.^[1-3] *R.ornithinolytica* is known to inhabit aquatic environments and can be found in hospital environments. *R.ornithinolytica* isolates that could have been erroneously reported as *Klebsiella* are sometimes be underestimated due to difficulties in its identification that may have led to uncertainty in its incidence, epidemiology and its pathogenic role in human infections.^[1-5] The pathogenic potential of *R.ornithinolytica* isolates in human disease has become increasingly important. Recently, clinical reports of *Raoultella* in clinical microbiology laboratories have been increased, probably through the introduction of accurate molecular techniques to identify. Many cases of infection caused by *R.ornithinolytica* have been reported so far.^[1-5] However, the clinical features, antimicrobial susceptibility, and clinical outcomes of *R.ornithinolytica* infection have not yet been well elucidated. There are

very few studies in Medical literature, regarding MBL detection among *R.ornithinolytica* in India and abroad as compared to other members of family *Enterobacteriaceae*.^[3-8] In this prospective study conducted in a tertiary care centre in Pune, seventeen *R.ornithinolytica* isolates were recovered from various clinical specimens, among them 13 were found to be Imipenem, Meropenem, Ertapenem (carbapenem) resistant based upon Minimum inhibitory concentrations (MIC) of antibiotics as determined by VITEK-2 AST-GN25 and AST-GN280 susceptibility cards. MICs were further determined by the E-test (bioMérieux, Marcy l'Etoile. MHT, DDST, CDST and MBL (IP/IPI) E-test was performed to detect Carbapenemase as well as Metallo-beta-lactamase production as described previously.^[2] In the present study, some important cases associated with *bla_{NDM-1}* producing *R.ornithinolytica* are reported below herein- Case 1-A 67-year-old male patient was transferred to our tertiary care centre; he was presented with hypotension, sensory impairment, and poor general condition, poor peripheral perfusion, desaturation and crackling rales. He was undergoing

chemotherapy with methotrexate (20 mg/week) and prednisone (150mg/day) for acute myeloid leukemia. Two days after his admission, *R.ornithinolytica* isolate was obtained from his blood culture and a tracheal aspirate also yielded the same organism, the patient was treated with carbapenem and an amino glycoside. Concurrently, a hypermucoviscous *R.ornithinolytica* strain was recovered from a second tracheal aspirate culture (135 CFU/ml). The isolate was resistant to all beta-lactams including carbapenems, aminoglycosides, fluoroquinolones, trimethoprim sulfamethoxazole, doxycycline, fosfomycin, but remained susceptible to colistin, and tigecycline. MHT, MBL E-test and *bla_{NDM-1}* was positive.

Case 2- A 68-year-old female patient was transferred from a nursing home to our ICU with symptoms of pneumonia, respiratory failure and shock, clinical examination revealed bronchopneumonia in the right lower lung field. Tracheal aspirate culture yielded carbapenem-resistant *R.ornithinolytica*, and the patient received fosfomycin, and cefepime. Due to progressive leukocytosis, unstable hemodynamics, and increased O₂ demand, colistin was prescribed instead of cefepime. The patient recovered well and was discharged alive from the ICU after 14 days. The isolate was resistant to all beta-lactams, carbapenems, aminoglycosides, fluoroquinolones, trimethoprim sulfamethoxazole, doxycycline, fosfomycin, but remained susceptible to colistin, and tigecycline. MHT, MBL E-test and *bla_{NDM-1}* was found to be positive.

Case 3-A 62-year-old female patient had bilateral mediastinal lymph nodes and multiorgan metastasis, catheter-related bacteremia was suspected, and blood cultures grew carbapenem-resistant *R.ornithinolytica* positive for MHT, MBL E-test and *bla_{NDM-1}*.

Case-4 A 57-year-old female who had been on peritoneal dialysis for the past six years because of end-stage renal disease secondary to diabetes mellitus was admitted for kidney transplant. The patient at the time of admission complained of dysuria, fever and chills. On physical examination, her general condition was moderate and she had a blood pressure of 150/70 mmHg, a heart rate of 115 beats/min, a respiratory rate of 21 breaths/min, and his body temperature was 38 °C. She had suprapubic tenderness, white blood cell count was 19,400 cells/ μ L (absolute neutrophil count, 16,811 cells/ μ L), and urinalysis showed a high number of white blood cells and erythrocytes. Her biochemical test results were as follows: plasma creatinine 12.8 mg/dL, blood urea nitrogen 92 mg/dL, sodium 139 mEq/dL, potassium 5.2 mEq/dL, and C-reactive protein (CRP) 110 mg/L (reference range, 0–5 mg/L). The peritoneal fluid cell count was 15/mm³. Two peripheral blood cultures, urinary culture and a peritoneal fluid culture sent to microbiology lab. The blood cultures and peritoneal fluid culture turned out negative but her urine culture yielded *bla_{OXA-48}* & *bla_{NDM-1}* producing *R.ornithinolytica* resistant to all beta-lactams, aminoglycosides, fluoroquinolones,

trimethoprim sulfamethoxazole, doxycycline, fosfomycin, but remained susceptible to colistin. The patient was treated with empirical broad-spectrum intravenous colistin for 10 days.

Case-5 was a 63-year-old man who underwent coronary bypass surgery, without complications. 8TH days after the surgery, the patient presented with dysuria and urgency without fever. A urinary tract infection was diagnosed. The urine analysis showed 800,000 leukocytes/mL, and the culture yielded *bla_{OXA-48}* & *bla_{NDM-1}* producing *R.ornithinolytica*.

Case-6 was a 58-year-old woman who underwent cardiac surgery, owing to surgical complications; she was admitted to the ICU. 7th day post-discharge patient presented with dysuria and pelvic discomfort. The urine analysis showed 739,000 leukocytes/ mL, and the culture yielded *bla_{OXA-48}* & *bla_{NDM-1}* producing *R.ornithinolytica* resistant to all beta-lactams, aminoglycosides and fluoroquinolones, but remained susceptible to colistin.

Case-7 was a 58-year-old man with chronic renal failure on peritoneal dialysis who was admitted to the hospital with a lower UTI. The urine analysis showed 562,000 leukocytes/mL, and the culture yielded *bla_{OXA-48}* & *bla_{NDM-1}* producing *R.ornithinolytica* resistant to all beta-lactams, aminoglycosides and fluoroquinolones, but remained susceptible to colistin.

A urinary catheter was not used (Case 4-7) in later four patients. Treatment for all patients included colistin for 10 days, without adverse effects. Urine analyses were conducted, and cultures were collected at the end of the treatment and again one week later. No leukocyturia was evident, and all cultures remained negative. Upon admission, the urine cultures of all four patients showed colony counts >100,000 CFU/mL (colony-forming units per milliliter), with growth of *R.ornithinolytica* identified by using the Vitek 2 Compact® automated system. The MHT, MBL E-test and PCR for both *bla_{OXA-48}* & *bla_{NDM-1}* genes were positive for all urinary isolates. Antibiogram of 11 *bla_{NDM-1}* producing *R.ornithinolytica* as a donor strains and its Transconjugants shown in Table-1. REP PCR, ERIC PCR and RAPD PCR showed that Isolates were clonally different. *bla_{NDM-1}* gene was located on IncA/C and IncHI2 while IncL/M associated with *bla_{OXA-48}*. *bla_{CTX-M-15}* was located on IncY, IncP and IncT whereas *bla_{TEM-1}* was associated with IncFIA. *bla_{SHV-12}* located on IncFIB while *bla_{SHV-28}* associated with IncFIC. **Plasmid Typing and characterization of MBL gene shown in Table-2.**

Table-1: Showing Antibiogram of 11 *R.ornithinolytica* bla_{NDM-1} producers selected as a donor strains for conjugation studies.

ISOLATE	IPM	MEM	ETP	ATM	CST	TGC	CAZ	CTX	FOX	FEP	CPZ	PIT	AMP	CIP	LVX	MOX	GEN	AMK	TOB
BACT 721	32	32	32	64	0.5	1	128	128	64	64	128	>128	256	4	8	4	>16	64	>16
TC 721	8	16	8	0.5	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	0.5	1	1	2	2	4
BACT 732	32	32	32	64	0.25	2	128	128	64	64	128	>128	256	8	8	8	>16	64	>16
TC732	16	16	8	0.5	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	0.5	0.5	0.5	2	2	2
PC-RO-18	32	32	32	64	0.5	2	128	128	64	64	128	>128	256	4	4	4	>16	64	>16
TCRO18	8	8	8	0.5	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	1	1	1	2	2	1
PC-RO-23	32	32	32	64	0.25	0.75	256	256	64	64	256	>128	256	4	8	8	>16	64	>16
TCRO23	16	16	8	1	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	0.25	1	1	2	4	1
ETB –521	32	32	32	64	0.25	1	128	128	64	64	128	>128	256	8	8	8	>16	64	>16
TCRO521	16	16	8	0.5	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	0.5	1	1	4	4	2
IAI -RO-8	32	32	32	64	0.25	0.5	128	128	64	64	128	>128	256	4	4	4	>16	64	>16
TC -RO-8	16	16	8	1	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	1	0.5	1	2	4	4
UC-RA05	32	32	32	64	0.125	2	256	256	64	64	256	>128	256	4	8	8	>16	64	>16
TCRO5	16	16	8	1	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	0.5	0.25	1	2	2	2
UC-RA-13	64	64	32	64	0.5	0.5	128	128	64	64	128	>128	256	4	8	4	>16	64	>16
TCRA13	8	8	8	0.5	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	0.25	1	0.5	1	2	2
UC-RA-25	32	32	32	64	0.25	1	128	128	64	64	128	>128	256	8	4	8	>16	64	>16
TCRA25	16	32	8	0.5	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	1	1	0.5	1	2	2
UC-RA-34	64	64	32	64	0.125	0.5	128	128	64	64	128	>128	256	4	8	4	>16	64	>16
TCRA34	8	8	8	1	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	0.5	1	0.25	1	2	2
UC-RA-56	32	32	32	64	0.5	0.75	256	256	64	64	256	>128	256	4	4	8	>16	64	>16
TCRA56	8	8	8	1	<0.25	<0.5	>64	>64	>64	>64	>64	>64	>64	1	0.5	1	1	2	2

Imipenem (IMP), meropenem (MEM), ertapenem (ETP), azetronam (ATM), colistin (CST), tigecycline (TGC); ampicillin(AMP), ceftazidime (CAZ), Cefotaxime (CTX), Cefoxitin (FOX), cefepime (FEP), ceftriaxone (CRO), cefoperazone (CPZ),Piperacillin-Tazobactam (PIT), ciprofloxacin (CIP), levofloxacin (LVX), moxifloxacin (MXF), gentamicin (GEN), amikacin (AMK), and tobramycin (TOB),

Table-2: Showing Plasmid Typing and characterization of beta-lactamase genes present in *R.ornithinolytica*.

ISOLATE	MBL	Plasmid	Transfer	Other ESBL gene present			Plasmid type			Transfer	Class D	Plasmid
BACT 721	NDM-1	A/C	Transferable	TEM-1	CTXM-15	SHV-28	FIA	T	FIC	Transferable	*	*
BACT 732	NDM-1	HI2	Transferable	TEM-1	CTXM-15	SHV-12	FIA	T	FIB	Transferable	*	*
PC-RO-18	NDM-1	A/C	Transferable	TEM-1	CTXM-15	SHV-28	FIA	P	FIC	Transferable	*	*
PC-RO-23	NDM-1	HI2	Transferable	TEM-1	CTXM-15	SHV-12	FIA	P	FIB	Transferable	*	*
ETB –521	NDM-1	HI2	Transferable	TEM-1	CTXM-15	SHV-28	FIA	Y	FIC	Transferable	*	*
IAI -RO-8	NDM-1	A/C	Transferable	TEM-1	CTXM-15	SHV-28	FIA	P	FIC	Transferable	*	*
UC-RA05	NDM-1	HI2	Transferable	TEM-1	CTX-M-15	SHV28	FIA	T	FIC	Transferable	OXA-48	L/M
UC-RA-13	NDM-1	HI2	Transferable	TEM-1	CTX-M-15	SHV28	FIA	T	FIC	Transferable	OXA-48	L/M
UC-RA-25	NDM-1	A/C	Transferable	TEM-1	CTX-M-15	SHV12	FIA	T	FIB	Transferable	OXA-48	L/M
UC-RA-34	NDM-1	A/C	Transferable	TEM-1	CTX-M-15	SHV12	FIA	T	FIB	Transferable	OXA-48	L/M
UC-RA-56	NDM-1	A/C	Transferable	TEM-1	CTX-M-15	SHV12	FIA	T	FIB	Transferable	OXA-48	L/M

The presence of a urinary catheter is considered a complicating factor probably due to the biofilm production that may be related with microbiological failure, catheter absence might have contributed to the successful treatment of these four patients.

Furthermore, colistin might be able to eradicate biofilms caused by *R.ornithinolytica*. Over the past decade, *R.ornithinolytica* has emerged as an infrequent, but important causal agent of human infections. To our knowledge, our study reports *bla*_{OXA-48} & *bla*_{NDM-1} producing *R.ornithinolytica* resistant to all beta-lactams, aminoglycosides, fluoroquinolones, trimethoprim sulfamethoxazole, doxycycline, fosfomycin, but remained susceptible to colistin. There are few reports regarding Carbapenem resistance *bla*_{NDM-1} producing *R.ornithinolytica* from India and especially from Indian sub continent.^[1-5] Our findings are similar to the findings of other studies,^[1-5] since this variation in genes is due to the fact that carbapenems are recently being introduced for their use in hospitals, previously being only used for nonfermenters, and due to their consistent use along with aminoglycosides and other antibiotics in ICUs, there is an increase in development of resistance against *R.ornithinolytica* due to irrational and inappropriate use of higher antibiotics in medical settings and their selective pressure along with factors such as lengthy ward-stay, debilitating clinical condition especially among immunosuppressed, immunocompromised, impaired immunity in ICU patients, and frequent exposure to medical interventions such as mechanical ventilation, tracheostomy, catheters, surgery or severe burns also aid in development and dissemination of resistant genes. In this study, we found have that *bla*_{NDM-1} gene located on IncA/C plasmids was immediately bracketed by a truncated insertion sequence ISAb125 upstream and the bleomycin resistance gene *ble*_{MBL} downstream using a PCR mapping approach and sequencing and demonstrated that these plasmids are highly conserved, particularly in regions encoding proteins involved in stability and conjugal transfer. We also have found that IncHI2 type broad-host-range plasmid carrying insertion sequence ISAb125 fragment containing the 145 promoter region, the *bla*_{NDM-1} gene, the bleomycin resistance gene *ble*_{MBL}, and a truncated transposon Tn125. This study provides an insight into the acquisition and emergence of *bla*_{OXA-48} & *bla*_{NDM-1} producing *R.ornithinolytica*, and emphasizes its transmission capability through plasmids. It is a cause for great concern as treatment options are virtually exhausted.

REFERENCES

1. Drancourt M., Bollet C, Carta A, and Rousselier P. Phylogenetic analyses of *Klebsiella* species delineate *Klebsiella* and *Raoultella* gen. nov., with description of *Raoultella ornithinolytica* comb. nov., *Raoultella terrigena* comb. nov. and *Raoultella planticola* comb. nov. *Int J Syst Evol Microbiol*, 2001; 51: 925–932.

2. Khajuria A, Praharaj AK, Grover N, Kumar M. First report of *bla*_{NDM-1} in *Raoultella ornithinolytica*. *Antimicrob Agents Chemother*, 2013; 57: 1092-3.
3. Haruki Y, Hagiya H, Sakuma A, Murase T, Sugiyama T, and Kondo S. Clinical characteristics of *Raoultella ornithinolytica* bacteremia: a case series and literature review. *J Infect Chemother*, 2014; 20: 589–591.
4. Castanheira, M., Deshpande, L.M., DiPersio, J.R., Kang, J., Weinstein, M.P., and Jones, R.N. First descriptions of *bla*_{KPC} in *Raoultella* spp. (*R. planticola* and *R. ornithinolytica*): report from the SENTRY Antimicrobial Surveillance Program. *J Clin Microbiol*, 2009; 47: 4129–4130.
5. Ponce-Alonso M, Rodríguez-Rojas L, Del Campo R, Cantón R, and Morosini MI. Comparison of different methods for identification of species of the genus *Raoultella*: report of 11 cases of *Raoultella*-causing bacteraemia and literature review. *Clin Microbiol Infect*, 2015.
6. Boattini, M., Almeida, A., Cardoso, C., Cruz, C.S., Machado, C., Vesza, Z. et al. Infections on the rise: *Raoultella* spp., clinical and microbiological findings from a retrospective study, 2010–2014. *Infect Dis (Lond)*, 2016; 48: 87–91.
7. Seng P, Boushab BM, Romain F, Gouriet F, Bruder N and Martin C. Emerging role of *Raoultella ornithinolytica* in human infections: A series of cases and review of the literature. *International Journal of Infectious Diseases*, 2016; 45: 65-71.
8. Walckenaer E, Poirel L, Leflon-Guibout V, et al. Genetic and Biochemical Characterization of the chromosomal class A β -lactamases of *Raoultella* (formerly *Klebsiella*) *planticola* and *Raoultella ornithinolytica*. *Antimicrob Agents Chemother*, 2004; 48: 305-312.