

Klebsiella pneumoniae ST147 Coproducing NDM-7 Carbapenemase and RmtF 16S rRNA Methyltransferase in Minnesota

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A cquired 16S rRNA methyltransferase (16S-RMTase) has emerged as a mechanism of high-level aminoglycoside resistance among Gram-negative pathogens worldwide (1). RmtF is a 16S RMTase which was identified in a single *Klebsiella pneumoniae* isolate on La Réunion Island in 2011 (2). The isolate also produced NDM-1 and was resistant to all β -lactams and aminoglycosides tested. Subsequently, *rmtF* was identified in 34 of 140 aminoglycoside-resistant *Enterobacteriaceae* isolates collected in India (3). Twenty of these 34 isolates also carried *bla*_{NDM-1}. Here, we report the identification of a *K. pneumoniae* strain coproducing NDM-7 and RmtF and an *Escherichia coli* strain producing NDM-7 from a patient who was admitted to a hospital in Minnesota.

In 2012, *K. pneumoniae* was isolated from the intra-abdominal abscess of a 69-year-old man who had been referred from India to a tertiary hospital in Minnesota. He had presented with abdominal pain and jaundice and had undergone endoscopic retrograde pancreatic cholangiography with biliary stenting twice while in India. In Minnesota, he underwent laparoscopic pancreaticoduo-denectomy, and a diagnosis of adenocarcinoma of the pancreatic head was made. Six days postoperatively, he developed a complicated intra-abdominal infection for which he underwent percutaneous drainage of a fluid collection. The culture grew an *E. coli*

isolate (EC1) that was resistant to ceftriaxone and susceptible to carbapenems. One week into treatment with piperacillin-tazobactam, another drain was put in place, and a repeat culture grew carbapenem-resistant *K. pneumoniae* (KP1). Six weeks later, the drains were adjusted, and repeat cultures grew carbapenem-resistant *K. pneumoniae* (KP2) and carbapenem-resistant *E. coli* (EC2). He initially received colistin, vancomycin, metronidazole, and caspofungin after KP1 was isolated, but treatment was limited by renal toxicity; therapy was completed with vancomycin, tigecycline, meropenem, and caspofungin. He ultimately made a complete recovery, and the abdominal drains were removed.

KP2 and EC2 were available for further analysis; they were resistant to all β -lactams tested, including carbapenems tested by broth microdilution (Sensititre GNX2F; Trek Diagnostics, Oak-

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TABLE 1 MICs of K. pneumoniae and E. coli isolates, their transformants, and control strains
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	MIC (µg/ml)						
Drug	K. pneumoniae KP2	E. coli TOP10 transformants for KP2			E. coli	<i>E. coli</i> TOP10 transformant	E. coli
		pCTX-M-15	pNDM-7-KP2	pRmtF	EC2	for EC2 (pNDM-7-EC2)	TOP10
Imipenem	>8	≤ 1	2	≤1	>8	2	≤1
Ertapenem	>4	≤0.25	4	≤0.25	4	4	≤0.25
Doripenem	>2	≤0.12	>2	≤0.12	>2	>2	≤0.12
Meropenem	>8	≤ 1	4	≤ 1	2	4	≤ 1
Ceftazidime	>16	>16	>16	≤ 1	>16	>16	≤1
Cefotaxime	>32	>32	>32	≤ 1	>32	>32	≤ 1
Ticarcillin-clavulanate	>128/2	128/2	>128/2	≤16/2	>128/2	>128/2	≤16/2
Piperacillin-tazobactam	>64/4	$\leq 8/4$	>64/4	$\leq 8/4$	>64/4	>64/4	$\leq 8/4$
Cefepime	>16	>16	16	≤ 2	>16	16	≤ 2
Aztreonam	>16	>16	≤ 2	≤ 2	16	≤ 2	≤2
Gentamicin	$>\!\!8$	≤1	≤ 1	> 8	$>\!\!8$	≤ 1	≤1
Tobramycin	> 8	≤ 1	≤ 1	> 8	8	≤ 1	≤1
Amikacin	>32	8	≤ 4	>32	≤ 4	≤ 4	≤ 4
Ciprofloxacin	>2	≤0.25	≤0.25	≤0.25	>2	≤0.25	≤0.25
Levofloxacin	8	≤ 1	≤ 1	≤ 1	$>\!\!8$	≤ 1	≤ 1
Doxycycline	8	≤ 2	≤ 2	≤ 2	$>\!\!8$	≤2	≤2
Minocycline	16	≤ 2	≤ 2	≤ 2	8	≤ 2	≤2
Tigecycline	1	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25
Trimethoprim-sulfamethoxazole	>4/76	>4/76	≤/0.5/9.5	≤0.5/9.5	>4/76	≤0.5/9.5	$\leq 0.5/9.5$
Colistin	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25
Polymyxin B	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25	≤0.25

wood Village, OH) and interpreted by Clinical and Laboratory Standards Institute guidelines (Table 1) (4). KP2 was additionally resistant to all aminoglycosides tested. The sequence types (STs) of KP2 and EC2 were ST147 and ST617, respectively (5, 6). Both isolates were positive for *bla*_{NDM-7} and *bla*_{CTX-M-15} by PCR and sequencing of the entire genes (7). NDM-7 is a variant of NDM-1 with a two-amino-acid substitution that appears to have improved hydrolytic efficiency of carbapenems compared with NDM-1 (8). KP2 was also positive for *rmtF* by PCR and sequencing of an internal fragment of the gene. The plasmids of KP2 and EC2 were extracted by the standard alkaline lysis method and used to transform competent E. coli TOP10 cells to obtain transformants harboring plasmids with these genes using lysogenic agar plates containing ampicillin or gentamicin (9). As a result, KP2 vielded three different transformants demonstrating resistance to cephalosporins, carbapenems, and aminoglycosides, each with bla_{CTX-M-15}, bla_{NDM-7}, or *rmtF*, whereas EC2 yielded a transformant with *bla*_{NDM-7} (Table 1). The replicon could be determined only for the *bla*_{CTX-M-15}-carrying plasmid of KP2, which was assigned to IncR (10). Nonetheless, the identification of $bla_{\rm NDM-7}$, a relatively infrequent *bla*_{NDM} allele (11), in both KP2 and EC2 suggested that the bla_{NDM-7}-carrying plasmids in these two strains may share an origin. The emergence of K. pneumoniae coproducing NDM metallo-B-lactamase and RmtF 16S-RMTase in the United States highlights the continuous threat of global dissemination of highly resistant enteric organisms by means of travel.

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REFERENCES

1. Wachino J, Arakawa Y. 2012. Exogenously acquired 16S rRNA methyltransferases found in aminoglycoside-resistant pathogenic Gramnegative bacteria: an update. Drug Resist Updat. 15:133–148. http://dx .doi.org/10.1016/j.drup.2012.05.001.

- Galimand M, Courvalin P, Lambert T. 2012. RmtF, a new member of the aminoglycoside resistance 16S rRNA N7 G1405 methyltransferase family. Antimicrob. Agents Chemother. 56:3960–3962. http://dx.doi.org/10.1128 /AAC.00660-12.
- Hidalgo L, Hopkins KL, Gutierrez B, Ovejero CM, Shukla S, Douthwaite S, Prasad KN, Woodford N, Gonzalez-Zorn B. 2013. Association of the novel aminoglycoside resistance determinant RmtF with NDM carbapenemase in Enterobacteriaceae isolated in India and the UK. J. Antimicrob. Chemother. 68:1543–1550. http://dx.doi.org/10.1093/jac/dkt078.
- 4. Clinical and Laboratory Standards Institute. 2014. Performance standards for antimicrobial susceptibility testing; twenty-fourth informational supplement (M100-S24). Clinical and Laboratory Standards Institute, Wayne, PA.
- Diancourt L, Passet V, Verhoef J, Grimont PA, Brisse S. 2005. Multilocus sequence typing of *Klebsiella pneumoniae* nosocomial isolates. J. Clin. Microbiol. 43:4178–4182. http://dx.doi.org/10.1128/JCM .43.8.4178-4182.2005.
- Wirth T, Falush D, Lan R, Colles F, Mensa P, Wieler LH, Karch H, Reeves PR, Maiden MC, Ochman H, Achtman M. 2006. Sex and virulence in *Escherichia coli*: an evolutionary perspective. Mol. Microbiol. 60: 1136–1151. http://dx.doi.org/10.1111/j.1365-2958.2006.05172.x.
- Doi Y, O'Hara JA, Lando JF, Querry AM, Townsend BM, Pasculle AW, Muto CA. 2014. Co-production of NDM-1 and OXA-232 by *Klebsiella pneumoniae*. Emerg. Infect. Dis. 20:163-165. http://dx.doi.org/10.3201 /eid2001.130904.
- Gottig S, Hamprecht AG, Christ S, Kempf VA, Wichelhaus TA. 2013. Detection of NDM-7 in Germany, a new variant of the New Delhi metalloβ-lactamase with increased carbapenemase activity. J. Antimicrob. Chemother. 68:1737–1740. http://dx.doi.org/10.1093/jac/dkt088.
- 9. Sambrook J, Russell DW. 2001. Molecular cloning: a laboratory manual, 3rd ed. Cold Spring Harbor Laboratory Press, Cold Spring Harbor, NY.
- Carattoli A, Bertini A, Villa L, Falbo V, Hopkins KL, Threlfall EJ. 2005. Identification of plasmids by PCR-based replicon typing. J. Microbiol. Methods 63:219–228. http://dx.doi.org/10.1016/j.mimet.2005.03.018.
- Rahman M, Shukla SK, Prasad KN, Ovejero CM, Pati BK, Tripathi A, Singh A, Srivastava AK, Gonzalez-Zorn B. 2014. Prevalence and molecular characterisation of New Delhi metallo-β-lactamases NDM-1, NDM-5, NDM-6 and NDM-7 in multidrug-resistant Enterobacteriaceae from India. Int. J. Antimicrob. Agents 44:30–37. http://dx.doi.org/10 .1016/j.ijantimicag.2014.03.003.