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Citation for final published version:

Tanner, Windy D, Atkinson, Robyn M, Goel, Ramesh K., Toleman, Mark A., Benson, Lowell Scott, Porucznik, Christina A. and VanDerslice, James A. 2017. Horizontal transfer of the blaNDM-1 gene to Pseudomonas aeruginosa and Acinetobacter baumannii in biofilms. FEMS Microbiology Letters 364 (8), pp. 102-120. 10.1093/femsle/fnx048

Publishers page: https://doi.org/10.1093/femsle/fnx048

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- Title: Horizontal transfer of the bla_{NDM-1} gene to Pseudomonas aeruginosa and Acinetobacter
 baumannii within a biofilm
- **Running title:** Transfer of *bla*_{NDM-1} into *P. aeruginosa* and *A. baumannii*
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- 6 Benson, a Christina A. Porucznik, James A. VanDerslice

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ABSTRACT

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Horizontal gene transfer has contributed to the global spread of the bla_{NDM-1} gene. Studies 21 22 have demonstrated plasmid transfer of bla_{NDM-1} into various Gram-negative bacterial species, but 23 attempts to demonstrate transfer of bla_{NDM-1} plasmids into Pseudomonas aeruginosa and Acinetobacter baumannii have either been unsuccessful or only observed with a donor of the 24 same genus. There is evidence that plasmid transfer frequency may increase when conjugation 25 occurs within a biofilm versus between planktonic cells. To determine whether bla_{NDM-1} gene 26 27 transfer to P. aeruginosa or A. baumannii could be facilitated in a biofilm environment, one E. coli and two Klebsiella pneumoniae strains carrying NDM-1-encoding plasmids of different 28 29 incompatibility types were mated with an E. coli J53 strain to produce E. coli J53-bla_{NDM-1} transconjugant plasmid donors. Dual-species biofilms were then created using the <u>E. coli</u> J53 30 transconjugants and a P. aeruginosa or A. baumannii recipient strain and incubated for 24 or 72 31 hours. Transfer of an NDM-encoding plasmid from one E. coli J53- bla_{NDM-1} transconjugant into 32 P. aeruginosa was successful in a 72-hour biofilm, and transfer of NDM-encoding plasmids 33 34 from two E. coli J53-bla_{NDM-1} transconjugants to A. baumannii was successful in 24-hour 35 biofilms,

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Deleted: Biofilm transfer of NDM-encoding plasmids to these bacterial species has serious implications for community and healthcare environments.

INTRODUCTION

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53 Spread of the bla_{NDM-1} carbapenemase gene is a global public health concern 54 (Kumarasamy et al., 2010). The gene is typically mobile, commonly carried on plasmids of diverse sizes and incompatibility types that are capable of inter-species, inter-genus, and inter-55 family transfer. (Carattoli, 2013). Successful dissemination of the bla_{NDM-1} gene is more 56 2013). ¶ commonly attributed to conjugational transfer of NDM-1-encoding plasmids to other Gram-57 negative bacteria, rather than clonal spread (Carattoli, 2013, Johnson & Woodford, 2013). 58 Deleted: T 59 The bla_{NDM-1} gene is typically found in Enterobacteriaceae species, but has also been Deleted: , globally 60 detected in a variety of non-fermenting Gram-negative bacteria such as Aeromonas caviae, Deleted: -located or Deleted: frequently 61 Stenotrophomonas maltophilia, Acinetobacter baumannii, and several Pseudomonas species, including Pseudomonas aeruginosa (Walsh et al., 2011, Zhang et al., 2013). Dissemination of 62 the bland-1 gene in P. aeruginosa and A. baumannii isolates has now been reported globally in 63 both clinical and environmental samples, and treatment options have become significantly 64 including Deleted: ers limited (Chen et al., 2011, Chaudhary & Payasi, 2013). Many of these isolates likely acquired 65 the bla_{NDM-1} gene via intra- or inter-genus conjugational transfer of NDM-1-encoding plasmids; 67 however, successful inter-family transfer of the bla_{NDM-1} harboring plasmids to P. aeruginosa or Formatted: Subscript 68 A. baumannii has not yet been demonstrated in the laboratory (Potron et al., 2011, Janvier et al., Deleted: in nature 2013, Huang et al., 2015). 69 Deleted: plasmid Deleted: gene 70 Conjugation experiments using Gram-positive donors and recipients typically use surface mating approaches such as filter mating or biofilm formation (Roberts et al., 2001, Savage et al., 71 Formatted: Font: Italic 72 2013). However, because pili can assist in gene transfer in Gram-negative bacteria, broth mating

has been a commonly used method for conjugational transfer of NDM-encoding plasmids in

Gram-negative species (Potron et al., 2011, Sowmiya et al., 2012, Rahman et al., 2014).

Deleted: The rapid global increase in hospital- and community-acquired carbapenem-resistant infections is an urgent public health threat, and very few treatment options remain (Centers for Disease Control and Prevention, 2013). Carbapenem resistance mediated by carbapenemase enzymes is of particular concern, as the encoding genes are often found on mobile genetic elements that can be transferred horizontally to other bacterial species (Johnson & Woodford,

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Bacterial species such as *P. aeruginosa* and *A. baumannii*, can frequently be recovered from hospital or natural environments (Blanc *et al.*, 2007, Walsh *et al.*, 2011, Nutman *et al.*, 2016), and are often found in a sessile or biofilm state (Donlan, 2002, Gurung *et al.*, 2013). Studies have demonstrated that horizontal gene transfer can occur at higher frequencies in biofilms versus planktonic cells in Gram-negative bacterial species (Madsen *et al.*, 2012). The high density and close spatial proximity of the cells create an ideal environment for interspecies transfer of genetic information (Donlan, 2002, Madsen *et al.*, 2012). *P. aeruginosa* and *A. baumannii* are commonly associated with biofilm formation (Donlan, 2002, Qi *et al.*, 2016), which could potentially facilitate the transfer of *bla*_{NDM-1} to these bacteria. The objective of this study was to determine whether plasmid-borne *bla*_{NDM-1} genes originating in *Enterobacteriaceae* species could be transferred from an *E. coli* J53-*bla*_{NDM-1} transconjugant to *P. aeruginosa* or *A. baumannii* in a biofilm environment.

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MATERIALS AND METHODS

Donor and recipient organisms

Four NDM-1-producing *Enterobacteriaceae* strains with plasmids of different incompatibility types carrying the *bla*_{NDM-1} gene were used as the original plasmid donors: a *K. pneumoniae* donor (EKP) carrying the *bla*_{NDM-1} gene on a 100 kb plasmid (pEKP) belonging to the FII, L/M, or N2 incompatibility group; an *E. coli* donor (EEC) carrying the *bla*_{NDM-1} gene on a 150 kb plasmid (pEEC) belonging to the FII incompatibility group; a *K. pneumoniae* donor (CO-NDM) carrying the *bla*_{NDM-1} gene on a 130 kb plasmid (pCO-NDM) of unknown incompatibility type; and a *K. pneumoniae* donor (ATCC BAA-2146) carrying the *bla*_{NDM-1} gene on a 140 kb plasmid (pNDM-US) belonging to the A/C incompatibility group (Hudson *et al.*, 2014). EKP and EEC were recovered from environmental samples in Southeast Asia, and the

132 CO-NDM and ATCC strains were isolated from clinical samples. Azide-resistant E. coli J53 was **Deleted:** Details on the four NDM-1-producing *Enterobacteriaceae* strains used as *bla*_{NDM-1} donors in the first phase of the conjugation assays are given in Table 1. 133 used as a recipient for mating experiments with the original plasmid donor strains. Subsequent Deleted: an intermediate mating-out assays were performed using the E. coli J53-bla_{NDM-1} transconjugants as donors and 134 Deleted: . Deleted: as previously described (Potron et al., 2011) 135 rifampin-resistant P. aeruginosa and A. baumannii as recipients E. coli J53-bla_{NDM-1} Deleted: , and rifampin-resistant P. aeruginosa and A. baumannii were used as final recipients. transconjugants were used as donors to allow for comparison of transfer frequencies between 136 Formatted: Font: Italic 137 each of the NDM-encoding plasmid types, as previously described (Potron et al., 2011). 138 Broth conjugations into azide-resistant E. coli J53 139 Log phase Luria-Bertani (LB) broth cultures of each of the bla_{NDM-1} donors and the E. 140 coli J53 recipient were combined in a 10:1 donor-to-recipient ratio in fresh LB. Mating-out 141 assays were performed as described by Walsh et al. (Walsh et al., 2011) using LB rather than nutrient broth. Conjugation mixtures were incubated overnight at 30 and 37 degrees C, then 142 serially diluted and plated on LB agar containing 0.5 µg/mL meropenem and 100 µg/mL sodium 143 azide. 144 Transfer of bla_{NDM-1} into E. coli J53 was confirmed by PCR with a previously described 145 primer set (Poirel et al., 2011) and on CHROMagar OrientationTM (DRG International, 146 Springfield, New Jersey) containing 0.5 µg/mL meropenem. Putative NDM-1-positive E. coli 147 J53 transconjugants from the EEC donor were differentiated from the parent EEC strain by PCR 148 149 detection of the yja-A gene found in *E. coli* J53 but absent in the EEC strain (Clermont et al., Formatted: Font: Italic 150 2000). Three of the bla_{NDM-1} donors, EKP, EEC, and ATCC-BAA-2146 produced E. coli J53 Formatted: Font: Italic transconjugants, designated E. coli TcEKP, E. coli TcEEC, and E. coli TcNDM-US, 151 152 respectively. 153 Biofilm conjugations into rifampin-resistant P. aeruginosa and A. baumannii Deleted: and broth

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163 Biofilm mating-out assays were performed using NDM-1-positive E. coli J53 transconjugants 164 TcEKP, TcEEC, and TcNDM-US as <u>blandm-1</u> plasmid donors and P. aeruginosa and A. 165 baumannii as recipients. Using optical density measurements, log-phase donor and recipient LB 166 broth cultures were combined in a 1:4 donor-to-recipient ratio, with approximately 2.5 x 10⁷ cells 167 from an E. coli J53 transconjugant donor and 1.0 x 108 cells from the P. aeruginosa or A. baumannii recipient in 1 mL LB. 168 The 1 mL dual-species conjugation mixtures were placed in a 48-well plastic plate and 169 170 incubated for either 24 or 72 hours, allowing the culture to form a biofilm on the sides of the 171 wells. Conjugation mixtures were incubated at 30 degrees for the *P. aeruginosa* conjugations and 172 37 degrees for the A. baumannii conjugations. LB was exchanged every 24 hours to maintain nutrient levels. At the end of the incubation period the broth was again exchanged, and biofilms 173 were scraped from the well sides using a sterile metal scraper. The LB containing the biofilm 174 175 scrapings was pulse-vortexed to break apart the cells, and serial dilutions of the biofilm 176 suspension were plated on tryptic soy agar containing 75 µg/mL ticarcillin and 50 or 100 µg/mL 177 rifampin for A. baumannii or P. aeruginosa, respectively. Plates were incubated at 37 degrees C 178 for 48 hours. 179 Colonies from ticarcillin-rifampin selection plates were subcultured to CHROMagarTM with 0.5 µg/mL meropenem. P. aeruginosa or A. baumannii colony lysates from the 180 CHROMagarTM plates were tested for the *bla*_{NDM-1} gene by PCR, as described above. 181 Minimum inhibitory concentrations 182 183 Minimum inhibitory concentrations (MICs) of all bla_{NDM-1} donors, recipients, and transconjugants were determined by meropenem Etest® (bioMérieux Clinical Diagnostics, 184

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Marcy l'Etoile, France).

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Deleted: Cultures of each NDM-1-positive *E. coli* J53 transconjugant and the *P. aeruginosa* or *A. baumannii* recipient were combined in a 1:4 donor-to-recipient ratio in LB, as previously described (Potron *et al.*, 2011), with broth mating assays otherwise performed as detailed above.

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Deleted: When blaNDM-1 biofilm transfer to P. aeruginosa or A. baumannii was successful, NDM-1-positive E. coli J53 donors were used to create a "layered" biofilm to confirm biofilm transfer. Single species 24-hour P. aeruginosa and A. baumannii biofilms were formed and then rinsed with LB, followed by addition of an NDM-1-positive E. coli J53 cultures to each biofilm well. Following a 48-hour incubation, with daily LB exchange, biofilms were then harvested and plated as described above. ¶

217	Plasmid analysis	Deleted: and <i>bla</i> CMY-2 resistance
1040		Deleted: s
218	Genetic location (plasmid or chromosome) of the bla _{NDM-1} gene after conjugational	Deleted: bacterial
219	<u>transfer</u> was determined by a combination of Pulsed Field Gel Electrophoresis (PFGE) of S1	Deleted: Broth conjugations into azide-resistant E. coli J53¶
220	digested macro DNA of the various <u>plasmid</u> donor strains and transconjugants followed by	Figure 1 shows the overall sequence of the mating-out assays and results. Transfer rates of NDM-encoding plasmids from the ATCC BAA-2146 <i>K. pneumoniae</i> , EKP, and EEC donor
221	detection using ³² P labelled <i>bla</i> _{NDM-1} and <i>bla</i> _{CMY-2} probes using methods described by Patzer et al	strains to <i>E. coli</i> J53 are listed in Table 1. ¶ Broth and
222	2009 (Patzer et al., 2009). Probes were prepared by PCR using primers pairs: NDMF/R	Deleted: Conjugations
		Deleted: for
223	TGGCTTTTGAAACTGTCGCACC, CTGTCACATCGAAATCGCGCGA; CMY2F/R	Deleted: NDM-1-positive E. coli J53 and
224	AAATCGTTATGCTGCGCTCT, GACACGGACAGGGTTAGGAT, respectively.	Deleted: mating assays
224	AAATCOTTATOCTOCOCTCT, OACACOOACAOOOTTAOOAT, Tespectivery.	Deleted: 2
225	RESULTS	Deleted: bla _{NDM-1} -carrying
		Formatted: Font: Not Italic
226	Biofilm conjugations into P. aeruginosa and A. baumannii	Deleted: plasmid
227	Transfer frequencies of the blandm-1 plasmids from the E. coli J53-blandm-1	Deleted: was not observed in broth from any donor, but was successfully transferred
228	transconjugants to P. aeruginosa or A. baumannii are presented in Table 1. Transfer of the	Deleted: the ATCC BAA-2146 parental strain (via the
220	unisconfugation to 1. deruguiosa of 11. buminimum pre presented in Table 1. Transfer of the	Deleted: bla _{NDM-1} -J53 E. coli intermediate)
229	<u>bla_{NDM-1} plasmid pNDM-US</u> to <i>P. aeruginosa</i> from <u>TcNDM-US</u> was successful in a 72-hour	Deleted: and within the 48-hour layered biofilm
	THE CLASSIC COLUMN TO SECOND TO SECO	Deleted: ¶
230	biofilm, but was not detected in a 24-hour biofilm, Transfer of the bla _{NDM-1} plasmids pNDM-US	Deleted: to A. baumannii
231	and pEKP from the TcNDM-US and TcEKP to A. baumannii was successful in a 24-hour	Deleted: ATCC BAA-2146 parental strain (via
		Deleted: the <i>bla</i> _{NDM-1} -J53 <i>E. coli</i> intermediate
232	biofilm, but was not detected in a 72-hour biofilm.	Deleted:)
222	Minimum inhibitory concentrations	Formatted: Font: Not Italic
233	Minimum minibitory concentrations	Deleted: both
234	All parental <i>bla</i> _{NDM-1} donor strains had meropenem MICs greater than 32 μg/mL. All	Deleted: broth and
235	transconjugants had meropenem MICs of 24 µg/mL or greater.	Deleted: Transfer was also successful from the EKP parental strain plasmid (via the <i>bla</i> NDM-1-J53 <i>E. coli</i>
		Deleted: Transfer of <i>bla</i> _{NDM-1} -carrying plasmids to the <i>A</i> .
236	Plasmid analysis	Deleted: Meropenem MICs for all donor, recipient, and
237	\$1 PFGE and bla _{NDM-1} ³² P labeled probe showed that the ATCC BAA-2146 bla _{NDM-1}	Deleted: Plasmid sizes and incompatibility groups for eac
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238	donor harbored a NDM-encoding plasmid of the approximately 140 kb. An NDM-encoding	Formatted: Subscript
220	placemid of the same size was found in the D. gamusin as him a transcenius at D.	Formatted: Font: Italic
239	plasmid of the same size was found in the <u>P. aeruginosa-bla_{NDM-1} transconjugant, P. aeruginosa</u>	Formatted: Subscript
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286	TcATCC, and in the A. baumannii-bla _{NDM-1} transconjugant, A. baumannii TcATCC. Analysis of		Deleted: and ethidium bromide staining (Figure 2 (b)). Probing of the gel with a <i>blac</i> _{MY} -2 ³² P labeled probe (Figure 2
287	the EKP bla _{NDM-1} donor by S1 PFGE and bla _{NDM-1} ³² P labeled probe showed that the bla _{NDM-1}		(a) indicated that the <i>blac</i> _{MY-2} ere abeted probe (Figure 2 (a)) indicated that the <i>blac</i> _{MY-2} gene was associated with a 140kb plasmid in the <i>K. pneumoniae</i> ATCC donor only and probing of a replicate gel (Figure 2 (c)) indicated that the same 140kb plasmid was associated with the <i>bla</i> _{NDM-1} gene. Probed gels also indicated that transfer of the 140kb plasmid to <i>P. aeruginosa</i> and <i>A. baumannii</i> occurred as a result of mating the recipient strains with the <i>E. coli</i> J53- <i>bla</i> _{NDM-1} ATCC intermediate donor (Figure 2 (a-c), Figure 3).
288	gene, was located on a 100kb plasmid and a bland plasmid of the same size was found in the		
289	A. baumannii-bla _{NDM-1} transconjugant, A. baumannii TcEKP. S1 PFGE analysis did not show		
290	insertion of the <i>bla</i> _{NDM-1} gene into the chromosome of either <i>P. aeruginosa</i> or <i>A. baumannii</i> .		Interestingly, in one of the P . $aeruginosa$ transconjugants the $bla_{\text{CMY-2}}$ and $bla_{\text{NDM-1}}$ positive plasmid was slightly smaller
291	DISCUSSION		(130kb). The probed gel indicated that the <i>bla</i> _{CMY-2} and <i>bla</i> _{NDM-1} positive plasmid in <i>P. aeruginosa</i> was found in various multimeric forms.
292	We successfully transferred plasmids carrying the <i>bla</i> _{NDM-1} gene from <i>Klebsiella and E</i> .		Formatted: Font: Italic
			Deleted: isolate
293	<u>coli</u> donor strains into <u>E. coli</u> J53, and subsequently from the <u>E. coli</u> J53- <u>bla_{NDM-1}</u>		Deleted: revealed
294	transconjugants into <i>P. aeruginosa</i> and <i>A. baumannii</i> , Plasmid analyses by S1 PFGE and <i>bla</i> _{NDM} -		Deleted: that the
	<u> </u>		Deleted: resided
295	¹ ³² P labeled probe indicated that the <u>blandm-1</u> gene remained plasmid-located in the P.		Deleted: was associated with the same
	and the second of the second o		Deleted: 100kb plasmid in
296	<u>aeruginosa</u> and <u>A. baumannii <u>bla_{NDM-1}</u> transconjugants in our study. To our knowledge, plasmid</u>		Deleted: EKP
97	transfer of the bla _{NDM-1} gene into P. aeruginosa has not been observed experimentally, and		Formatted: Font: Italic
		ansusus M	Formatted: Font: Italic
298	plasmid transfer of <i>bla</i> _{NDM-1} into <i>A. baumannii</i> has previously only been <u>demonstrated by</u>	100000000000000000000000000000000000000	Deleted: s (Figure 3)
299	electroporation (Potron et al., 2011), or using agar surface mating techniques and an		Deleted: The S1 PFGE analysis also indicated that the <i>bla</i> _{NDM-1} gene in the CO NDM-1-producing <i>K. pneumoniae</i> isolate was plasmid-located, despite a lack of transfer into the
300	Acinetobacter donor (Huang et al., 2015). Agar and filter surface mating methods, like biofilm,		153 <i>E. coli</i> . The <i>bla</i> _{NDM-1} gene was associated with a 130kb plasmid in this isolate, which was <i>bla</i> _{CMY-2} negative (Figure 2).
301	provide a stable, spatially-structured environment for gene transfer; however, sites of nutrient		Deleted: Enterobacteriaceae
302	uptake and gas exchange differ, and less of the protective extracellular polymeric substance is	\\\\\	Deleted: via an NDM-1-positive <i>E. coli</i> J53 transconjuga
002	uptake and gas exchange differ, and less of the protective extracentual polymeric substance is		Moved (insertion) [2]
303	produced in agar colonies compared to biofilms (Davey & O'Toole, 2000, Stalder & Top, 2016).	- \\\\\	Deleted: While studies of clinical NDM-1-producing <i>P</i> .
304	Additionally, the plasmids donors in our assays were Enterobacteriaceae species and inter-	\\\\	Deleted: both
004	Additionally, the plasmids donors in our assays were Emeropacteriaceae species and inter-	\\\	Deleted: successful
305	family transfer via E. coli J53-bla _{NDM-1} transconjugants to P. aeruginosa and A. <u>baumannii</u> has	\\\	Deleted: using
			Deleted: mating techniques and
306	not previously been demonstrated with other NDM-1-encoding plasmids.		Deleted: (Shields <i>et al.</i> , 1985, Chen <i>et al.</i> , 2011, Jones <i>et</i>
307	NDM-1 mating assays described in the peer-reviewed literature frequently use broth		Formatted: Font: Italic
			Deleted: Transfer of NDM-encoding plasmids into <i>either</i>
808	mating techniques (Potron et al., 2011, Zhang et al., 2013, Ou et al., 2014). We did not observe		Formatted: Font: Italic
			Deleted: .

358 plasmid transfer of bla_{NDM-1} to P. aeruginosa in planktonic broth cultures, similar to other studies 359 (Potron et al., 2011, Janvier et al., 2013), and only pEKP transferred in broth to A. baumannii 360 (data not shown). Transfer of pEEC to either recipient was not detected. It should be noted that transfer of pEEC, or higher transfer rates of pEKP and pNDM-US donors, might have been 361 observed if mating with the P. aeruginosa and A. baumannii recipients had been performed using 362 363 the original parent bla_{NDM-1} donor, rather than via an *E. coli* 153 transconjugant. Mating was 364 performed using an E. coli J53- bla_{NDM-1} transconjugant to enable comparison of transfer frequency of the different NDM-1 plasmid types and comparison with previous attempts to 365 366 transfer blandmin plasmids into P. aeruginosa and A. baumannii (Potron et al., 2011). Prior 367 studies have been unable to detect transfer of NDM-encoding plasmids to P. aeruginosa or A. baumannii from Enterobacteriaceae donors by conjugation under broth conditions. This work 368 demonstrates that conjugative inter-family transfer of these plasmids can be successful when 369 370 mating occurs in a biofilm environment. 371

Funding Information

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- 373 This work was supported by the Health Studies Fund of the Department of Family and
- 374 Preventive Medicine of the University of Utah.

Conflict of Interest

376 The authors have no conflicts of interest to declare.

Acknowledgements

- 378 We wish to thank the Utah Public Health Laboratory for their assistance and the use of their
- facility for part of the PCR analyses of these isolates. We are also grateful to the Colorado
- 380 Department of Public Health & Environment Laboratory Services Division for providing the

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Moved up [2]: While studies of clinical NDM-1-producing *P. aeruginosa* isolates have found the *bla*_{NDM-1} gene located on the chromosome in *P. aeruginosa* (Janvier *et al.*, 2013, Jovcic *et al.*, 2014), the gene remained plasmid-located in both the *P. aeruginosa* and *A. baumannii* transconjugants in our study.

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Deleted: The plasmid was of identical size (140 kb) to the ATCC donor in one *P. aeruginosa* transconjugant but a second transconjugant was slightly smaller (approx. 130kb). This is likely due to a deletion event, which appears to be a common event during plasmid transfer of NDM-encoding plasmids (Kumarasamy *et al.*, 2010). Interestingly, in both *P. aeruginosa* transconjugants the NDM-encoding plasmids were found in multiple forms which are most likely multimeric forms of the same plasmid, which was also partially visible in the donor strain (Figure 2). Multiple copies of the *bla*NDM-1 gene on the *P. aeruginosa* chromosome have been reported previously (Jovcic *et al.*, 2014), and similar multimeric plasmid forms have also been seen in NDM-encoding plasmids in *Acinetobacter* species (Jones *et al.*, 2014).¶

In the Û.S., healthcare-acquired multidrug-resistant *Acinetobacter* and *P. aeruginosa* are responsible for an estimated 7300 and 6700 infections each year, respectively (Centers for Disease Control and Prevention, 2013). Biofilmforming *P. aeruginosa* and *A. baumannii* have been implicated in healthcare facility outbreaks traced to environmental sources (Hota *et al.*, 2009, Doidge *et al.*,

459	CO-NDM donor strain, and Dr. Mark Fisher for providing the clinical A. baumannii recipient
460	strain.
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