

First Report on a Hyperepidemic Clone of KPC-3-Producing *Klebsiella pneumoniae* in Israel Genetically Related to a Strain Causing Outbreaks in the United States[∇]

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A highly epidemic carbapenem-resistant clone of KPC-3-producing *Klebsiella pneumoniae* emerged in Israel in 2006, causing a nationwide outbreak. This clone was genetically related to outbreak strains from the United States isolated in 2000 but differed in KPC-carrying plasmids. The threat of the global spread of hyperepidemic, extensively drug-resistant bacterial strains should be recognized and confronted.

KPC-type enzymes in carbapenem-resistant *Klebsiella pneumoniae* strains were first reported in 2001 in North Carolina (16), and until 2005, the geographical distribution of these enzymes in *K. pneumoniae*, with both KPC-2 and KPC-3, was limited to the eastern United States (1, 12, 15). In the New York area, KPC-producing strains have become a frequently encountered nosocomial pathogen (2, 4). The first case of KPC-producing *K. pneumoniae* outside the United States occurred in France, where a patient who had been hospitalized in New York carried the strain with him (9). KPC-producing *K. pneumoniae* strains have since been reported from Israel (6), Colombia (13), China (14), and Greece (8).

Carbapenem resistance in *Klebsiella pneumoniae* carrying *bla*_{KPC-2} was first observed in the Tel Aviv Sourasky Medical Center in late 2005. In February 2006, we noted a sharp increase in *bla*_{KPC}-carrying, carbapenem-resistant *K. pneumoniae* strains in our hospital, mainly possessing *bla*_{KPC-3} (6). These extensively drug-resistant (XDR) isolates caused difficult-to-treat infections and had an adverse impact on patients' outcomes (11). Despite infection control efforts that limited the spread of KPC-2-producing clones of *Klebsiella* in our hospital, KPC-3-producing *K. pneumoniae* isolates continued to appear rapidly during 2006, in our hospital and in other hospitals throughout the country (10).

To characterize the extent of the nationwide occurrence of KPC-3-producing, carbapenem-resistant *K. pneumoniae* in Israel, 100 single-patient isolates collected during 2006 in eight hospitals and five chronic care centers with wide geographical

distribution in the north, center, and southern parts of the country were sent to our lab for further study. Six to eight isolates were included from each institution based on their resistance to at least one carbapenem antibiotic (imipenem and/or meropenem) determined using the Vitek2 automated system (bioMérieux, Marcy l'Etoile, France) with an AST-GN09 card. The presence of *bla*_{KPC} in all isolates was verified using PCRs followed by sequencing to determine the type of KPC gene (6). MICs of imipenem, meropenem, and ertapenem were determined by agar dilution according to the method of the Clinical and Laboratory Standards Institute (3). Susceptibility testing for tigecycline was performed via Etest according to the manufacturer's instructions (AB Biodisk, Solna, Sweden). Interpretive criteria for tigecycline MICs were defined based on the U.S. Food and Drug Administration breakpoint criteria for *Enterobacteriaceae* (susceptible, ≤ 2 $\mu\text{g/ml}$; intermediate, 4 $\mu\text{g/ml}$; resistant, ≥ 8 $\mu\text{g/ml}$). The genetic relatedness of all carbapenem-resistant *K. pneumoniae* strains was determined by pulsed-field gel electrophoresis (PFGE) analysis. DNA was prepared as described previously (6), and chromosomal restriction fragments obtained after SpeI or ApaI cleavage were documented and compared.

Sites of isolation included urine ($n = 46$), blood ($n = 19$), respiratory tract ($n = 15$), wounds ($n = 15$), and other ($n = 5$). The isolates were resistant to nearly all antimicrobial agents, including all cephalosporins, beta-lactam/beta-lactamase inhibitor combinations, trimethoprim-sulfamethoxazole, and fluoroquinolones; MICs of the carbapenems varied (Table 1). The majority of the isolates (98%) were susceptible only to gentamicin and colistin, while the other 2% were gentamicin resistant and kanamycin susceptible or resistant to both aminoglycosides. Among 20 isolates tested for tigecycline, MICs were at 1 to 3 $\mu\text{g/ml}$. PFGE analysis revealed that all isolates belonged to the same genetic clone, indistinguishable from the KPC-3-producing clone described previously in our hospital (6) (Fig. 1A). This finding suggested the nationwide spread of an epidemic carbapenem-resistant *K. pneumoniae* strain, designated clone Q.

PCR and sequencing revealed the presence of *bla*_{KPC-3} in all isolates.

When the Israeli outbreak started, the only country in which

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TABLE 1. Carbapenem-resistant *K. pneumoniae* isolates included in this binational study

Country	Yr of isolation	No. of isolates	MIC ($\mu\text{g/ml}$) ^a		
			Imipenem	Meropenem	Ertapenem
Israel	2006–2008	100	16, 32 (<0.25–128)	64, 128 (2–256)	128, 256 (<0.5–>256)
United States ^b	2000–2006	26	4, 64 (<0.25–128)	8, 64 (<0.5–256)	16, 128 (<0.25–>256)

^a MIC₅₀, MIC₉₀ (MIC range).

^b Outbreak isolates from various states, including New York, Delaware, Maryland, Pennsylvania, New Jersey, and Arizona.

KPC-producing *K. pneumoniae* had been reported was the United States. Therefore, we compared the hyperepidemic strain isolated in Israel to a collection of 26 KPC-producing, carbapenem-resistant *K. pneumoniae* outbreak isolates from the United States. These U.S. isolates originated from patients with diverse infections who were hospitalized in five different states between 2000 and 2006 (Table 1).

Testing for genetic relatedness between the Israeli hyperepidemic XDR clone and the American isolates revealed that nine (35%) of the 26 KPC-3-producing U.S. isolates had PFGE profiles identical or highly similar to each other (Fig. 1B). These nine isolates represented outbreaks that occurred in a New York medical center in 2000 (15) and in New Jersey and Arizona in 2006. Four of these nine isolates were indistinguishable by PFGE from the Israeli epidemic clone, and the remaining five isolates were closely related to that clone (with differences of up to three bands). The other 17 isolates, representing an outbreak during 2004 in a New York medical center and clinical isolates identified in Delaware, Maryland, New Jersey, and Pennsylvania during 2006, differed from the common PFGE type by more than seven bands.

Plasmid analysis was performed to determine whether the highly genetically related Israeli-American *K. pneumoniae* clones carried the same KPC-3-encoding plasmid. Ten isolates were analyzed: five isolates from Israel (from 2006 to 2008) and five isolates from the United States (from 2000 to 2006). Plasmid DNA was purified using a NucleoBond PC100 plasmid

Midi kit (Macherey-Nagel GmbH, Duren, Germany), and transformation was performed into *Escherichia coli* GeneHogs (Invitrogen Corp., Dorset, United Kingdom). Transformants were selected on Luria-Broth agar plates with ampicillin (100 $\mu\text{g/ml}$), and selected colonies were subjected to *bla*_{KPC} PCR screening to confirm the acquisition of a KPC-encoding plasmid. KPC-encoding plasmids were purified from 10 transformants originating from the 10 genetically related Israeli and American isolates (Fig. 2A). All five Israeli isolates carried the same plasmid with an apparent molecular mass of 100 kb. The genetically related U.S. isolates carried various plasmids ranging in size from 165 kb to 38 kb, which differed from the typical KPC-3-encoding Israeli plasmid (Fig. 2B).

Transfer of the KPC-3-encoding plasmids to the susceptible *E. coli* GeneHogs recipient strain raised the MICs of extended-spectrum cephalosporins and aztreonam by more than 60-fold, rendering resistance. MICs of carbapenems, although increased (from 0.012 $\mu\text{g/ml}$ to 0.016 to 0.5 $\mu\text{g/ml}$ for meropenem and ertapenem, and from 0.094 $\mu\text{g/ml}$ to 0.125 to 1.0 $\mu\text{g/ml}$ for imipenem), were not in the resistant range. This observation may be due to the presence of additional mechanisms of carbapenem resistance in these strains, such as porin alterations (17).

Our finding of the occurrence of closely related *K. pneumoniae* strains carrying *bla*_{KPC-3} on different plasmids in two continents is intriguing.

This report on the international occurrence of a clone of

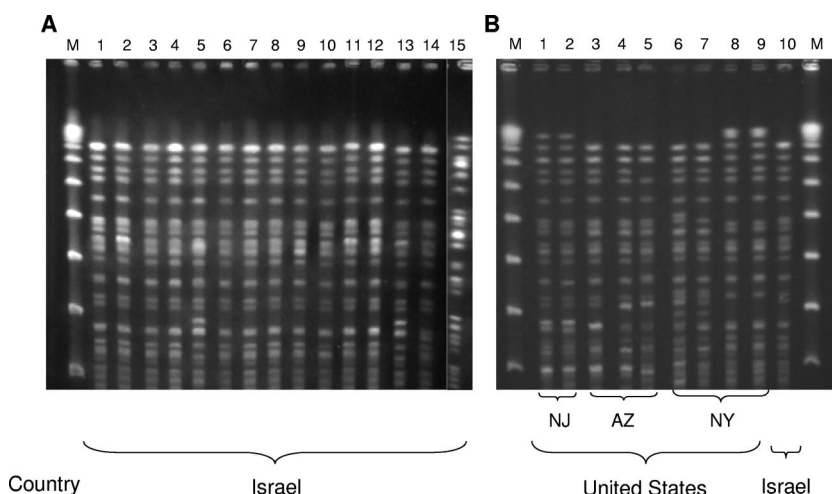


FIG. 1. (A) PFGE (SpeI restricted) of 14 representative KPC-3-producing, carbapenem-resistant *K. pneumoniae* isolates from various Israeli hospitals (lanes 1 to 14) demonstrates a nationwide epidemic clone different from a KPC-2-producing clone reported previously in our hospital (6) (lane 15). (B) Genetic relatedness of the predominant Israeli epidemic clone (lane 10) with nine KPC-3-producing *K. pneumoniae* isolates involved in outbreaks in the United States, from New Jersey (NJ; lanes 1 and 2), Arizona (AZ; lanes 3 to 5), and New York (NY; lanes 6 to 9). M, lambda ladder (New England Biolabs).

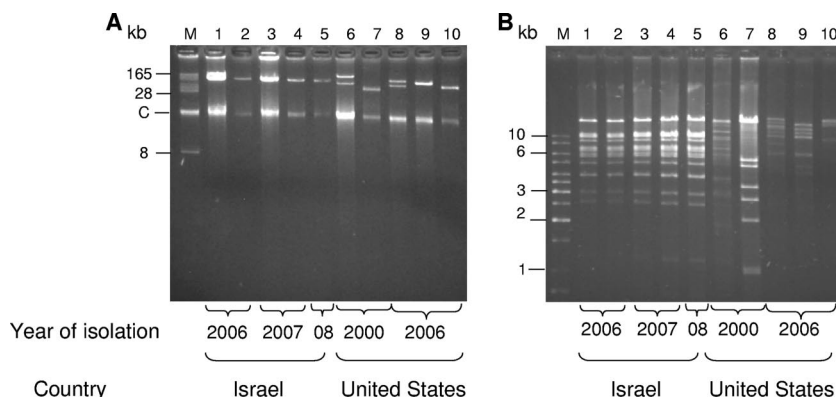


FIG. 2. A comparison of *bla*_{KPC-3}-encoding plasmids (A) and their EcoRI restriction patterns (B) from five Israeli *K. pneumoniae* isolates (lanes 1 to 5 in each panel) and five of the nine genetically related *K. pneumoniae* U.S. isolates (lanes 6 to 10 in each panel). Analysis was performed on plasmids isolated from *E. coli* GeneHogs transformants carrying the *bla*_{KPC-3}-encoding plasmids. Lanes 1 to 5, representative Israeli isolates from 2006 to 2008; lanes 6 to 10, representative U.S. isolates from 2000 and 2006; lanes 6 and 7, two isolates from a New York medical center; lanes 8 to 10, one isolate from New Jersey and two isolates from Arizona. For panel A, lane M is a BAC-Tracker supercoiled DNA ladder (Epicentre Biotechnologies); for panel B, lane M is a GeneRuler 1-kb DNA ladder (Fermentas Life Sciences).

XDR *bla*_{KPC-3}-producing *K. pneumoniae*, which has caused outbreaks in the United States and Israel, emphasizes the potential for transmission of highly resistant, gram-negative pathogens that are difficult, and in some cases impossible, to treat with currently available antimicrobial agents. This situation is exacerbated by the lack of novel agents in the antimicrobial pipeline for treating these pathogens. Our finding of similar strains in the United States and Israel raises the possibility of a clonal XDR strain of *K. pneumoniae*, akin to what has been described for the 300.0114 strain of methicillin-resistant *Staphylococcus aureus* (5) and the NAP1/027 strain of *Clostridium difficile* (7). This issue will require further investigation, but it may help inform future studies directed at characterizing the molecular mechanisms, environmental factors, and selection pressures which promote the spread of this XDR hyperepidemic strain. In the meantime, currently established infection control measures, such as hand hygiene, isolation precautions, and judicious antimicrobial use, should be employed throughout the world to limit the emergence and transmission of KPC-producing organisms.

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