

Supplementary Materials

Title Genetic background and antibiotic resistance profiles of *K. pneumoniae* NDM-1 strains isolated from UTI, ABU and the GI tract, from one hospital in Poland, in relation to strains nationally and worldwide

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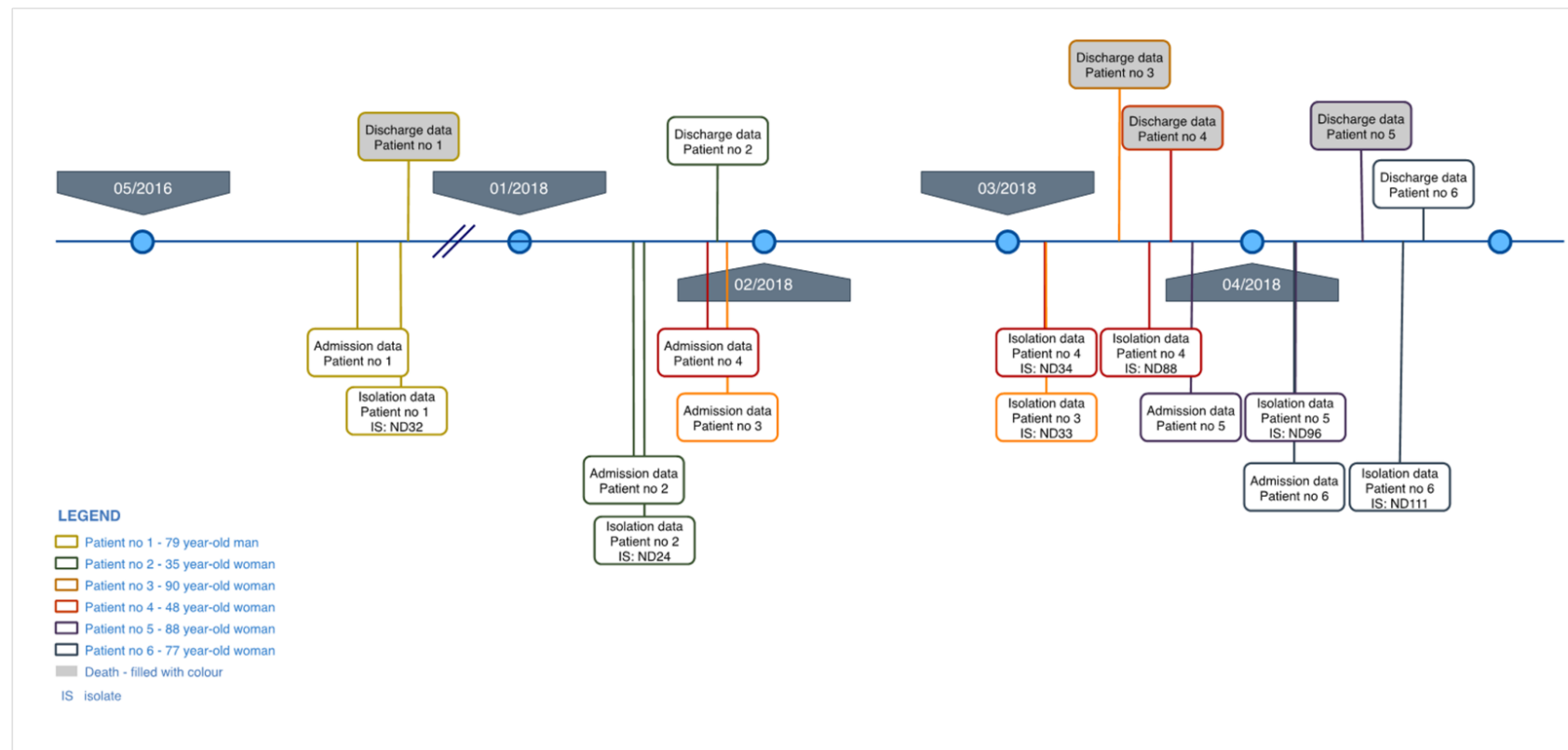


Figure S1. Timeline of events in the epidemiologically linked cases of New Delhi metallo- β -lactamase 1-producing *Klebsiella pneumoniae*. Dates are given as month/year. Abbreviations: IS, isolate.

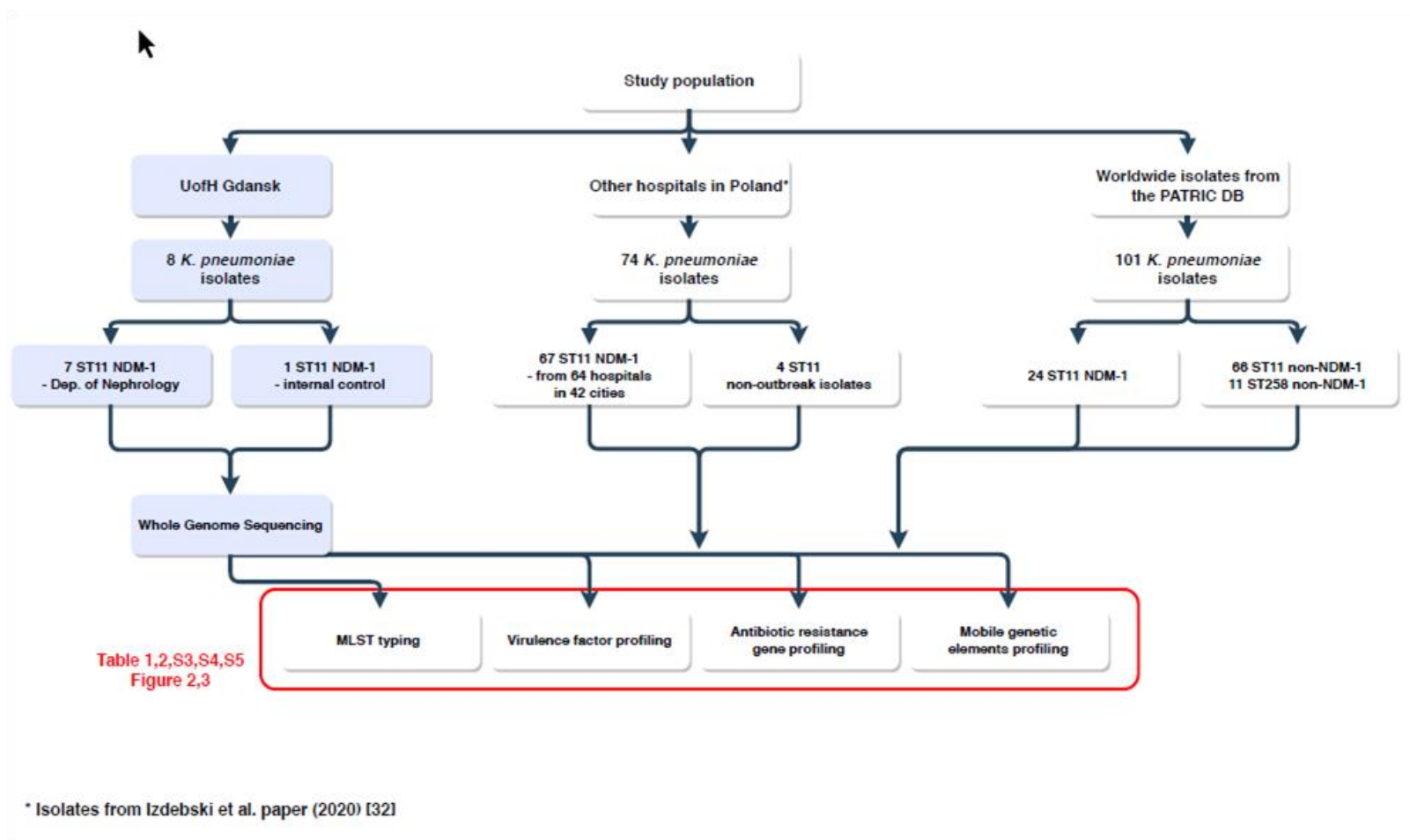


Figure S2. Diagram showing the groups of the analyzed strains in this work.

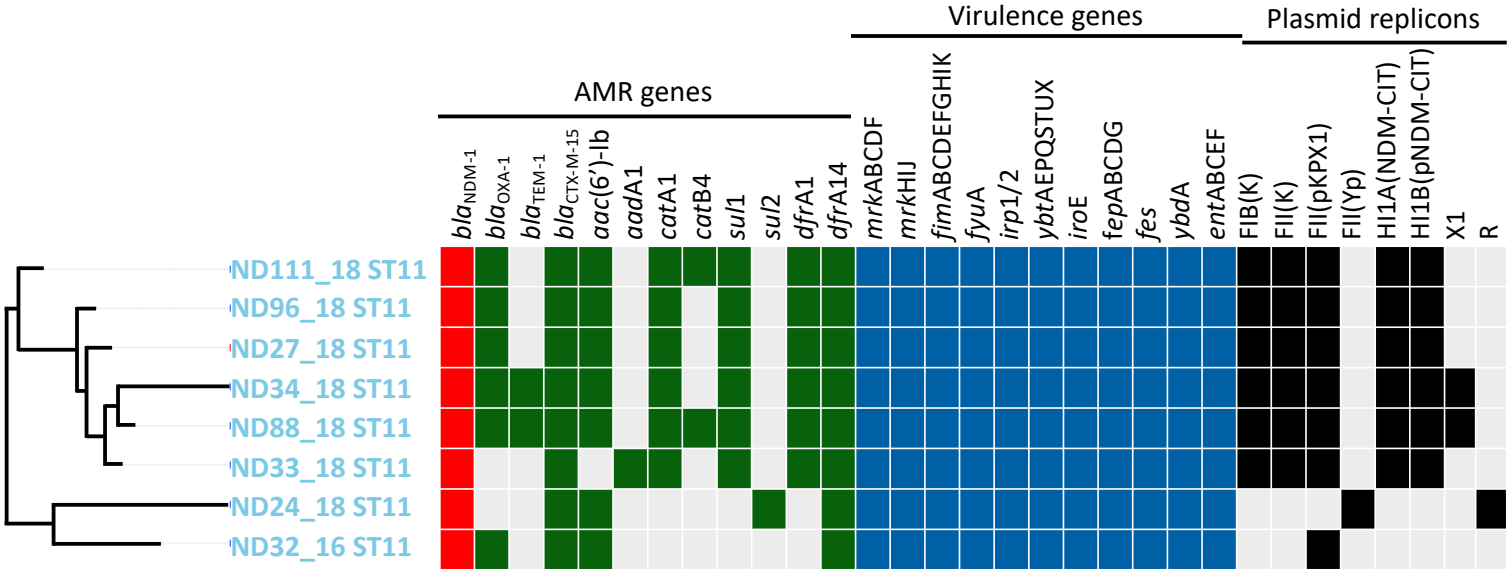


Figure S3. The core-gene phylogenetic tree for eight *K. pneumoniae* isolates from this study. The sequence type (ST) is indicated for each isolate, following the isolate name. In the heatmap the presence of the *bla*_{NDM-1} gene is indicated by red (present) or gray (absent) and the presence/absence profile of the genotype for genes encoding antimicrobial resistance (green – present, gray – absent), genes encoding virulence determinants (blue – present, gray – absent) and plasmid replicons is indicated.

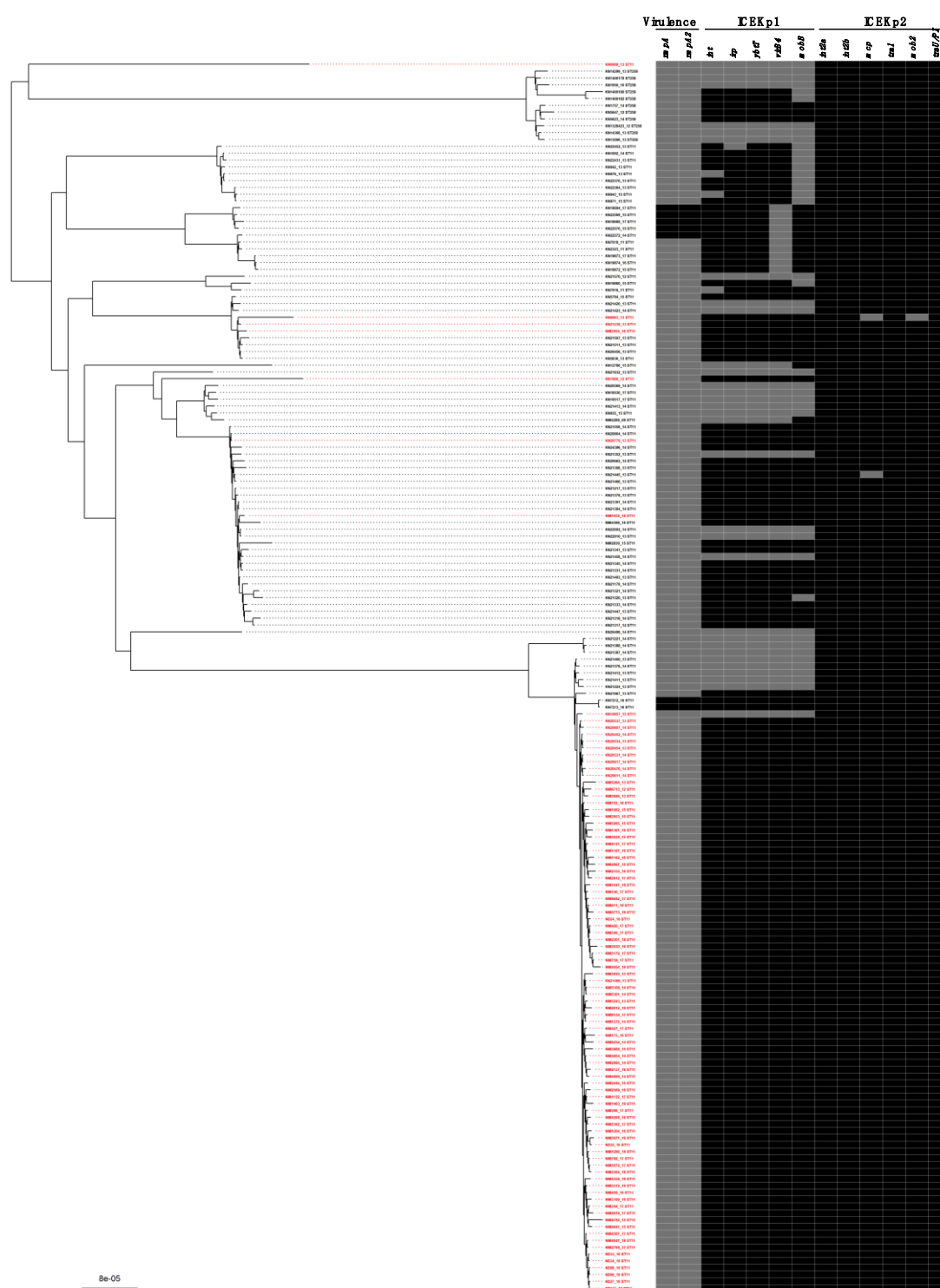


Figure S4. Occurrence of integrative conjugation elements (ICEKp1 and ICEKp2) in clinical isolates of *K. pneumoniae*. Genome core phylogenetic tree generated for 180 *K. pneumoniae* isolates. The color of the isolate corresponds to the presence or absence of the blaNDM-1 gene (red – present, black – absent). The sequence type (ST) is indicated for each isolate after the isolate name. The presence and coexistence of ICEKp1 and ICEKp2 (marker genes) is shown in the heatmap on the right (black – present, gray – absent). The virulence related genes *ompA* and *ompA2* were also included in the analysis to determine their correlation with the presence / absence of ICEKp1 and ICEKp2. The scale bar shown in the lower left corner shows the average number of substitutions per locus.

Table S1. Strain metadata. Characteristics of the single isolates, including collection details, genome assembly statistics, genomic features.

As- sem- bly/Sa- mple Name	Bi- osample accession	Coun- try	Hospi- tal	Year	Mon- th	Sour- ce	Graf- t	Infection Status	Age	Sex	ST	# con- tigs	Largest contig	Total length	GC (%)	N50	N75	L50	L75	# N's per 100 kbp	num_se- qs	sum_ len	min_l en	avg_le n	max_l en
ND27	SAMN17- 168596	Po- land	Torun	2018	5	urine	0	infection	ND	ND	11	138	483179	5966290	56.84	239522	101300	9	18	0.00	720,384	162,044,155	36	224.9	251
ND96	SAMN17- 168597	Po- land	Gdansk	2018	4	urine	0	infection	ND	K	11	113	804016	5959156	56.81	239630	100790	9	17	0.00	651,043	150,425,106	36	231.1	251
ND88	SAMN17- 168598	Po- land	Gdansk	2018	3	urine	1	coloniza- tion	47	K	11	136	483261	6006090	56.72	239522	99546	10	20	0.00	645,02	143,147,694	36	221.9	251
ND32	SAMN17- 168599	Po- land	Gdansk	2016	5	urine	0	infection	ND	M	11	98	483251	5441546	57.37	203937	101300	9	17	0.00	523,802	116,651,684	36	222.7	251
ND111	SAMN17- 168600	Po- land	Gdansk	2018	4	urine	0	coloniza- tion	ND	K	11	103	804016	5953882	56.82	241563	101300	8	17	0.00	651,012	142,164,105	36	218.4	251
ND24	SAMN17- 168601	Po- land	Gdansk	2018	1	urine	1	coloniza- tion	34	K	11	95	507788	5537205	57.30	258257	142128	8	16	0.00	710,264	168,737,035	36	237.6	251
ND33	SAMN17- 168602	Po- land	Gdansk	2018	3	anal swab	0	coloniza- tion	ND	K	11	112	491481	5956422	56.82	248278	153777	9	16	0.00	800,507	187,207,483	36	233.9	251
ND34	SAMN17- 168603	Po- land	Gdansk	2018	3	anal swab	1	coloniza- tion	ND	K	11	117	491539	5999106	56.73	258283	129570	9	17	0.00	524,443	121,940,935	36	232.5	251

Table S2. (EXCEL format). Metadata for all additional bacterial isolates included in the study from PATRIC database and Izdebski et al. paper [32].**Table S3. (EXCEL format)** The matrix of the core genome SNPs differences determined against the reference genome from the same ST.**Table S4.** *K. pneumoniae* isolates included in this study from PATRIC database—basic epidemiological data and resistomes based on sequencing WGS.

Isolates Name	Source	Year of isolation	Country	Host health	S T	Plasmid replicon profiles	Acquired antimicrobial resistance genes ^a							
							β-lactams	aminoglycosides ^b	fluoroquinolones ^b	macrolide, lincosamide, streptogramin antibiotics	phenicols	sulphonamides	tetracycline	trimethoprim
KN2323	urine	2011	China	ND	1	ColRNAI, FII(pHN7A8), L, R	blaKPC-2	<i>aadA2</i>	-	-	<i>catA2</i>	-	-	-
KN7016	blood	2011	China	sepsis	1	FIB(K), FII(K), I1-I(Gamma), R	blaCTX-M-55	-	<i>qnrB4</i> , <i>qnrS1</i>	<i>mph(A)</i>	-	<i>sul1</i>	-	-
KN7018	urine	2011	China	UTI	1	ColRNAI, FII(pHN7A8), L, N, R	blaKPC-2 blaCTX-M-65, blaTEM-1	<i>aac(6′)-Ib-cr</i> , <i>aadA16</i> , <i>aadA2</i>	<i>aac(6′)-Ib-cr</i>	-	<i>catA2</i>	<i>sul1</i>	<i>tet(A)</i>	<i>dfrA27</i>
KN21570	urine	2012	Germany	infection	1	FIB(K), FII, FII(K), L, R	blaCTX-M-15, blaOXA-48	<i>aac(3)-IVa</i> , <i>aph(3′′)-Ib</i> , <i>aph(4)-Ia</i> , <i>aph(6)-Id</i> , <i>aadA16</i> , <i>aadA2</i>	<i>qnrS1</i>	-	<i>catA2</i>	<i>sul1</i> , <i>sul3</i>	-	<i>dfrA27</i>

KN2 0456	urine	2013	Greece	infection	1 1	Col440I, FIB(K), FII(K), L, N	blaCTX-M-15, blaOXA-1,-48	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	<i>mph(A)</i>	-	<i>sul1</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1217	urine	2013	Slovakia	infection	1 1	FIA(HI1), FIB(K), FII(K), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1224	urine	2013	Czech Republic	infection	1 1	Col(pHAD28), C, FIB(K), FII(K), R	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aph(6)-Id</i>	<i>aac(6')-Ib-cr</i>	-	-	<i>sul1</i>	<i>tet(D)</i>	<i>dfrA22</i>
KN2 1306	urine	2013	Estonia	infection	1 1	FIB(K), FII(K)	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	-	-	<i>sul1</i>	-	-
KN2 1320	urine	2013	Poland	infection	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1341	urine	2013	Poland	infection	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1352	urine	2013	France	infection	1 1	FIB(K), FII(K), R	blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	<i>mph(A)</i>	<i>catA1</i> , <i>catB3</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 1378	urine	2013	Slovakia	infection	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i>	-	-	-	-	-	-

KN2 1400	urine	2013	Slova kia	coloniz ation	1 1	FIB(K), FII(K), R	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-Ila</i> , <i>aph(3'')-Ib</i> , <i>aph(6)-Id</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 1403	urine	2013	Slova kia	coloniz ation	1 1	FIB(K), FII(K), FIA(HII1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-Ila</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1411	urine	2013	Czec h Repu blic	infectio n	1 1	Col440I, FIB(K)	blaCTX-M-15, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-Ila</i>	<i>aac(6')-Ib-cr</i>	-	-	-	<i>tet(D)</i>	-
KN2 1412	urine	2013	Czec h Repu blic	infectio n	1 1	FIB(K), R	blaCTX-M-15, blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-Ila</i> , <i>aph(3'')-Ib</i> , <i>aph(6)-Id</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 1420	urine	2013	Czec h Repu blic	infectio n	1 1	FIB(K), FII(K)	blaOXA-1	<i>aac(6')-Ib-cr</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	<i>mph(A)</i>	<i>catA1</i> , <i>catB3</i>	<i>sul1</i>	-	-
KN2 1445	urine	2013	Eston ia	infectio n	1 1	FIB(K), FII(K), R	blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	<i>mph(A)</i>	<i>catA1</i> , <i>catB3</i>	<i>sul1</i>	-	<i>dfrA12</i>

KN2 1447	urine	2013	Poland	infection	1 1	FIA(HI1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1466	urine	2013	Estonia	colonization	1 1	FIB(K), FII(K), M1, R	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	-	<i>catA1</i> , <i>catB3</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 1507	urine	2013	Malta	colonization	1 1	Col440I, FIB(K), L, N	blaOXA-1,-48	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	<i>mph(A)</i>	-	-	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1511	urine	2013	Malta	infection	1 1	Col440I, FIB(K), FII(K), L, N	blaOXA-1,-48	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aph(3')-Ia</i>	<i>aac(6')-Ib-cr</i>	-	-	-	-	<i>dfrA14</i>
KN2 1932	urine	2013	Italy	infection	1 1	Col(BS512), ColRNAI, FIB(K), FII(K), R	blaTEM-1	<i>aac(3)-IVa</i> , <i>aph(4)-Ia</i> , <i>aadA2</i>	-	-	-	<i>sul3</i>	<i>tet(A)</i>	-
KN2 1967	urine	2013	Spain	infection	1 1	FIB(K), FII(K), R	blaCTX-M-15, blaOXA-1,-48	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB1</i>	<i>mph(A)</i>	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 2010	urine	2013	Hungary	infection	1 1	FIA(HI1), FIB(K), FII(K), R	blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 2370	urine	2013	USA	ND	1 1	ColRNAI, FIB(K), FII(K)	blaKPC-2, blaCTX-M-15,	<i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1</i> , <i>sul3</i>	-	<i>dfrA12</i>

							blaOXA-9, blaTEM-1							
KN2 2384	urine	2013	USA	ND	1 1	ColRNAI, FIB(K), FII(K), R	blaKPC-2, blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1, sul3</i>	-	<i>dfrA12</i>
KN2 2431	urine	2013	USA	ND	1 1	ColRNAI, FIB(K), FIB(pQil), FII(K), R, X3	blaKPC-2, blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aac(3)-IVa</i> , <i>aph(3')-Ia</i> , <i>aph(4)-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul1, sul3</i>	-	<i>dfrA12</i>
KN2 2452	urine	2013	USA	ND	1 1	ColRNAI, FIB(K), M1, R	blaOXA-2	<i>ant(2'')-Ia</i>	-	-	-	<i>sul1</i>	-	-
KN5 636	stool	2013	Greece	ND	1 1	ColI440I, C, FIB(K), FII(K), L	blaCTX-M-15, blaOXA-10, blaOXA-1,-48, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>ant(2'')-Ia</i> , <i>aph(3'')-Ib</i> , <i>aph(3')-Ia</i> , <i>aph(6)-Id</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	<i>mph(A)</i>	-	<i>sul2</i>	<i>tet(A)</i> , <i>tet(G)</i>	<i>dfrA12</i>
KN9 42	urine	2013	USA	ND	1 1	ColRNAI, FIB(K), FII(K), R, X1	blaKPC-2, blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1, sul2</i> , <i>sul3</i>	-	<i>dfrA12</i>
KN9 76	urine	2013	USA	ND	1 1	ColRNAI, FIB(K), FII(K), R, X1	blaKPC-2, blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1, sul2</i> , <i>sul3</i>	-	<i>dfrA12</i>

KN2 0499	urine	2014	Spain	infection	1 1	FIB(K), N, R	blaTEM-1	<i>aac(6′)-Ib-cr, aadA16, aph(3′′)-Ib, aph(6)-Id, aadA2</i>	<i>aac(6′)-Ib-cr, qnrB19, qnrB6</i>	-	-	<i>sul1, sul3</i>	-	<i>dfrA27</i>
KN2 0569	urine	2014	Portugal	ND	1 1	ColRNAI, R, X1	blaCTX-M-1, blaOXA-1	<i>aac(6′)-Ib-cr, aph(3′)-Ia</i>	<i>aac(6′)-Ib-cr, qnrB4</i>	<i>mph(A)</i>	<i>catB3</i>	<i>sul1</i>	-	-
KN2 0662	urine	2014	Portugal	ND	1 1	R	blaOXA-1	<i>aac(6′)-Ib-cr, aph(3′)-Ia</i>	<i>aac(6′)-Ib-cr, qnrB4</i>	<i>mph(A)</i>	<i>catB3</i>	<i>sul1</i>	<i>tet(A)</i>	-
KN2 0804	urine	2014	Romania	infection	1 1	R	blaOXA-1	<i>aac(6′)-Ib-cr, aadA2</i>	<i>aac(6′)-Ib-cr, qnrB4</i>	<i>mph(A)</i>	<i>catB3</i>	<i>sul1</i>	<i>tet(A)</i>	-
KN2 1178	urine	2014	Belgium	ND	1 1	FIB(K), FII(K), FIA(HI1), L, R	blaCTX-M-15, blaOXA-48,-9, blaTEM-1	<i>aac(6′)-Ib, aac(3)-IIa, aadA2</i>	-	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 1221	urine	2014	Slovakia	infection	1 1	ColRNAI, FIB(K), FII(K), R, C	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6′)-Ib-cr, aac(3)-IIa, aph(3′′)-Ib, aph(6)-Id</i>	<i>aac(6′)-Ib-cr</i>	-	<i>catA1</i>	<i>sul2</i>	<i>tet(D)</i>	-
KN2 1316	discharge from the lower respiratory	2014	Poland	colonization	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(3)-IIa, aadA2</i>	-	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12, dfrA14</i>

tury tract													
KN2 1317	urine	2014	Polan d	infectio n	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	-	-	<i>catA1</i>	<i>sul1</i>	- , <i>dfrA14</i>
KN2 1321	urine	2014	Polan d	infectio n	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	- , <i>dfrA14</i>
KN2 1331	discha rge from the lower respira tory tract	2014	Polan d	infectio n	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	-	<i>catA1</i>	<i>sul1</i>	- , <i>dfrA14</i>
KN2 1333	urine	2014	Polan d	infectio n	1 1	FIB(K), FII(K), FIA(HI1), R, M2	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IId</i> , <i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	- <i>dfrA12</i>
KN2 1345	urine	2014	Polan d	infectio n	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	- <i>dfrA12</i>
KN2 1376	urine	2014	Slova kia	coloniz ation	1 1	FIB(K), FII(K), R	blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i>	-	<i>catA1</i>	<i>sul1</i>	- <i>dfrA12</i>
KN2 1384	urine	2014	Slova kia	coloniz ation	1 1	Col440I, FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	- , <i>dfrA14</i>

KN2 1387	urine	2014	Slova kia	infectio n	1 1	FIB(K), FII(K), R	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aph(3'')-Ib</i> , <i>aph(6)-Id</i>	<i>aac(6')-Ib-cr</i>	-	-	<i>sul2</i>	<i>tet(D)</i>	-
KN2 1388	urine	2014	Slova kia	infectio n	1 1	FIB(K), FII(K), R	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aph(3'')-Ib</i> , <i>aph(6)-Id</i>	<i>aac(6')-Ib-cr</i>	-	<i>catA1</i>	<i>sul2</i>	<i>tet(D)</i>	-
KN2 1391	urine	2014	Slova kia	coloniz ation	1 1	FIB(K), FII(K), FIA(HI1), R	blaCTX-M-15, blaOXA-1,-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN2 1413	urine	2014	Franc e	coloniz ation	1 1	FIB(K), FII, M2, R	blaCTX-M-3, blaOXA-1, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aph(3'')-Ib</i> , <i>aph(3')-Ia</i> , <i>aph(6)-Id</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	-	<i>catB3</i>	<i>sul1</i>	-	-
KN2 1423	urine	2014	Czec h Repu blic	infectio n	1 1	FIB(K), FII(K), R	blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aph(3')-Ia</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	<i>mph(A)</i>	<i>catA1</i> , <i>catB3</i>	<i>sul1</i>	-	-
KN2 1458	urine	2014	Polan d	infectio n	1 1	FIB(K), FII(K)	-	<i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 1556	discha rge from the woun d	2014	Germ any	coloniz ation	1 1	FIB(K), FII(K), R	blaOXA-1	<i>aac(6')-Ib-cr</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB4</i>	<i>mph(A)</i>	<i>catA1</i> , <i>catB3</i>	<i>sul1</i>	-	<i>dfrA12</i>

KN2 2002	urine	2014	Hungary	infection	1 1	FIA(HI1), FIB(K), FII(K), R	blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aac(6')-Ib</i> , <i>aac(3)-IIa</i> , <i>aadA2</i>	<i>qnrB4</i>	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN2 2572	blood	2014	China	bloodstream infection	1 1	ColRNAI, FII(pHN7A8), HI1B(pNDM-MAR), I1-I(Gamma), R, repB	blaKPC-2, blaCTX-M- 14b,blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aadA16</i>	<i>aac(6')-Ib-cr</i>	-	<i>catA2</i>	<i>sul1</i>	-	<i>dfrA27</i>
KN2 4396	urine	2014	USA	ND	1 1	ColRNAI, pKPC- CAV1193	blaKPC-3, blaTEM-1	<i>aadA2</i>	<i>qnrS1</i>	-	-	<i>sul1</i>	-	-
KN1 002	urine	2014	USA	ND	1 1	ColRNAI, FIB(K), FIB(pQil), R	blaKPC-2, blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1, sul3</i>	-	<i>dfrA12</i>
KN1 2786	urine	2015	Nigeria	ND	1 1	ColKP3, FIB(K), FIB(pNDM-Mar), FII(K), R, X3	blaCTX-M-15, blaOXA-1,-181, blaTEM-1	<i>aac(6')-Ib-cr</i> , <i>aac(3)-IIa</i> , <i>aph(3'')-Ib</i> , <i>aph(6)-Id</i>	<i>aac(6')-Ib-cr</i> , <i>qnrB2</i> , <i>qnrS1</i>	<i>mph(A)</i>	<i>catA1</i>	<i>sul1</i>	<i>tet(A)</i>	<i>dfrA15</i>
KN1 9860	blood	2015	Brazil	infection	1 1	ColRNAI, FIB(K), FII(K), N	blaKPC-2, blaCTX-M-2, blaOXA-2, blaTEM-1	<i>aac(3)-IIa</i> , <i>aadA2</i>	-	-	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN1 9872	stool	2015	China	ND	1 1	FIB(K), FIB(pKPHS1), FII(pHN7A8), R	blaKPC-2, blaCTX-M-14 blaTEM-1	-	<i>qnrS1</i>	-	-	<i>sul1</i>	<i>tet(A)</i>	<i>dfrA1</i>
KN2 2569	blood	2015	China	bloodstream	1 1	ColRNAI, FII(pHN7A8),	blaKPC-2, blaCTX-M-147	<i>aadA2</i>	-	-	-	-	-	-

				infectio n		HI1B(pNDM-MAR), R, repB								
KN2 2570	blood	2015	China	bloodst ream infectio n	1 1	ColRNAI, FII(pHN7A8), HI1B(pNDM-MAR), repB	blaKPC-2, blaCTX-M-65, blaTEM-1	<i>aadA2</i>	-	-	-	-	-	-
KN5 794	urine	2015	Thailand	ND	1 1	Col8282, FIB(K), FII(K), I1-I(Gamma), R	blaCTX-M-15, blaOXA-1, blaTEM-1	<i>aac(6')-Ib- cr, aac(3)- IId, aph(3')- Ia, aadA17</i>	<i>aac(6')-Ib-cr, qnrB4</i>	<i>mph(A)</i>	<i>catA1, catB3</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN9 35	discharge from the wound	2015	USA	ND	1 1	FIA(HI1), FIB(K), FII(K), M1, R	blaCTX-M-15, blaOXA-1	<i>aac(6')-Ib- cr, aac(3)- IIa, ant(2'')- Ia, aadA2</i>	<i>aac(6')-Ib-cr, qnrS1</i>	-	-	<i>sul1</i>	<i>tet(A)</i>	<i>dfrA12, dfrA14</i>
KN9 43	urine	2015	USA	ND	1 1	ColRNAI, FIB(K), FII(K), R	blaKPC-2, blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aph(3')-Ia, aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1, sul3</i>	-	<i>dfrA12</i>
KN9 71	discharge from the wound	2015	USA	ND	1 1	ColRNAI, FIB(K), FII(K), R	blaKPC-2, blaCTX-M-15, blaOXA-9, blaTEM-1	<i>aph(3')-Ia, aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1, sul3</i>	-	<i>dfrA12</i>

KN1 9874	stool	2016	China	ND	1 1	FIB(K), FIB(pKPHS1), FII(pHN7A8), R	blaKPC-2, blaCTX-M-65 blaTEM-1	-	<i>qnrS1</i>	-	-	<i>sul1</i>	<i>tet(A)</i>	<i>dfrA1</i>
KN7 212	urine	2016	India	bloodst ream infection	1 1	ColKP3, FIB(pQil), FII(K), HI1B(pNDM- MAR), R, repB	blaCTX-M-15, blaOXA-232, blaTEM-1	<i>aph(3'')-Ib</i> , <i>aph(6)-Id</i>	<i>qnrB1</i>	-	-	<i>sul2</i>	-	-
KN7 213	blood	2016	India	bloodst ream infection	1 1	ColKP3, FIB(pQil), FII(K), HI1B(pNDM- MAR), R, repB	blaCTX-M-15, blaOXA-232, blaTEM-1	<i>aph(3'')-Ib</i> , <i>aph(6)-Id</i>	<i>qnrB1</i>	-	-	<i>sul2</i>	-	-
KN1 8517	urine	2017	USA	infection	1 1	C	-	-	-	-	-	<i>sul2</i>	-	-
KN1 8530	blood	2017	USA	infection	1 1	C, R	blaOXA-10	<i>aadA2</i>	-	-	-	<i>sul1, sul2</i>	<i>tet(D)</i>	<i>dfrA14</i>
KN1 8689	stool	2017	China	ND	1 1	ColRNAI, FII(pHN7A8), HI1B(p NDM-MAR), R, repB	blaKPC-2, blaCTX-M-65, blaTEM-1	<i>aadA2</i>	<i>qnrS1</i>	-	<i>catA2</i>	<i>sul2</i>	<i>tet(A)</i>	<i>dfrA14</i>
KN1 8694	stool	2017	China	ND	1 1	ColRNAI, FII(pHN7A8), N, R, repB	blaKPC-2, blaCTX-M-3, blaTEM-1	<i>aadA2</i> , <i>armA</i>	-	<i>mph(E)</i> , <i>msr(E)</i>	<i>catB8</i>	-	-	<i>dfrA12</i>
KN1 9873	stool	2017	China	ND	1 1	FIB(K), FIB(pKPHS1), FII(pHN7A8), R	blaKPC-2, blaCTX-M-90 blaTEM-1	-	<i>qnrS1</i>	-	-	<i>sul1</i>	<i>tet(A)</i>	<i>dfrA1</i>
KN1 950	urine	2010	Norway	ND	2 5 8	ColRNAI, FIB(K), FII(K), FII(Yp), I2(Delta)	blaKPC-3, blaOXA-9	<i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>

KN1 3284 23	urine	2012	USA	ND	2 5 8	ColRNAI, FIB(K), FII(K), R, X3	blaKPC-2,	<i>aac(6')-Ib</i> , <i>aac(3)-IVa</i> , <i>aph(4)-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul1, sul3</i>	-	<i>dfrA12</i>
KN5 647	urine	2012	Greece	ND	2 5 8	ColRNAI, FIB(K), FIB(pQil), FII(K), C, X3	blaKPC-2, blaOXA-10,-9	<i>aac(6')-Ib</i> , <i>ant(2'')-Ia</i> , <i>aph(3'')-Ib</i> , <i>aph(3')-Ia</i> , <i>aph(6)-Id</i> , <i>aadA2</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul2</i>	<i>tet(A)</i> , <i>tet(G)</i>	<i>dfrA12</i> , , <i>dfrA14</i> , , <i>dfrA23</i>
KN1 3096	urine	2013	USA	ND	2 5 8	ColRNAI, FIB(K), FIB(pQil), FII(K)	blaKPC-2, blaOXA-9	<i>aac(6')-Ib</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN1 4299	urine	2013	USA	ND	2 5 8	ColRNAI, X3	blaTEM-1	<i>aac(6')-Ib</i>	-	-	-	-	-	-
KN1 4368	urine	2013	USA	ND	2 5 8	ColRNAI, FIB(K), FIB(pQil), FII(K), X3	blaKPC-2, blaOXA-9	<i>aac(6')-Ib</i> , <i>aph(3')-Ia</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN1 757	urine	2014	Australia	UTI	2 5 8	ColRNAI, FIB(K), FII(K), FIB(pQil), X3	blaKPC-2, blaOXA-9	<i>aac(6')-Ib</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul1</i>	-	<i>dfrA12</i>
KN5 623	stool	2014	Greece	ND	2 5 8	ColRNAI, FIB(K), FII(K), M1, X3	blaKPC-2,	<i>aac(6')-Ib</i> , <i>aph(3')-Ia</i> , <i>aadA2</i>	-	<i>mph(A)</i>	-	<i>sul1</i>	-	<i>dfrA12</i>

KN1 4001 78	urine	ND	USA	ND	2 5 8	ColRNAI, FIB(K), FII(K), FII(Yp), X3	blaKPC-3, blaOXA-9, blaTEM-1	<i>aph(3'')-Ib</i> , <i>aph(6)-Id</i> , <i>aadA2</i>	-	<i>mph(A)</i>	<i>catA1</i>	<i>sul1, sul2</i>	-	<i>dfrA12</i> , <i>dfrA14</i>
KN1 4001 80	urine	ND	USA	ND	2 5 8	ColRNAI, FIA(pBK30683), FIB(K), X3	blaKPC-3, blaOXA-9	<i>aph(3'')-Ib</i> , <i>aph(6)-Id</i>	-	<i>mph(A)</i>	-	<i>sul2</i>	-	<i>dfrA14</i>
KN1 4001 82	urine	ND	USA	ND	2 5 8	FIA(pBK30683), FIB(K), X3	blaKPC-3,	<i>aph(3')-Ia</i>	-	<i>mph(A)</i>	-	-	-	<i>dfrA14</i>

^aOnly acquired resistance genes are shown, as identified by ResFinder 3.1 [34]. ^bThe *aac(6')-Ib-cr* gene shown in the 'aminoglycosides' and 'fluoroquinolones' columns is the same gene, conferring resistance to both classes of antimicrobials.

Table S5. Occurrence of integrative conjugation elements (ICEKp1 and ICEKp2) in clinical isolates of *K. pneumoniae* sequenced in this study and from PATRIC database and from Izdebski et al. paper [32]. (0 – absence, 1 - presence).

[illegible]

KN20454 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20456 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20470 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20499 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN20531 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20534 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20537 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20569 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN20607 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20611 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20617 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20662 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20779 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN20804 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21178 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21217 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21221 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21224 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21238 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21306 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21316 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21317 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21320 ST11	0	0	1	1	1	1	0	1	1	1	1	1	1	1
KN21321 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21331 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21333 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21341 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21345 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21352 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21376 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21378 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21384 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21387 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21388 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21391 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21400 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21403 ST11	0	0	1	1	1	1	1	1	1	1	1	1	1	1
KN21411 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21412 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21413 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21420 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1
KN21423 ST11	0	0	0	0	0	0	0	1	1	1	1	1	1	1

[illegible]

[illegible]

[illegible]

Table S6. *K. pneumoniae* isolates (non NDM-1) from PATRIC database included in the study – serotypes data and virulomes (0 – absence, 1 - presence)

[illegible]

KN21221	11	1	1	1	0	0	0	1	1	1	1	1
KN21224	11	1	1	1	0	0	0	1	1	1	1	1
KN21306	11	1	1	1	1	1	1	1	1	1	1	1
KN21316	11	1	1	1	1	1	1	1	1	1	1	1
KN21317	11	1	1	1	1	1	1	1	1	1	1	1
KN21320	11	1	1	1	1	1	1	1	1	1	1	1
KN21321	11	1	1	1	1	1	1	1	1	1	1	1
KN21331	11	1	1	1	1	1	1	1	1	1	1	1
KN21333	11	1	1	1	1	1	1	1	1	1	1	1
KN21341	11	1	1	1	1	1	1	1	1	1	1	1
KN21345	11	1	1	1	1	1	1	1	1	1	1	1
KN21352	11	1	1	1	0	0	0	1	1	1	1	1
KN21376	11	1	1	1	0	0	0	1	1	1	1	1
KN21378	11	1	1	1	1	1	1	1	1	1	1	1
KN21384	11	1	1	1	1	1	1	1	1	1	1	1
KN21387	11	1	1	1	0	0	0	1	1	1	1	1
KN21388	11	1	1	1	0	0	0	1	1	1	1	1
KN21391	11	1	1	1	1	1	1	1	1	1	1	1
KN21400	11	1	1	1	0	0	0	1	1	1	1	1
KN21403	11	1	1	1	1	1	1	1	1	1	1	1
KN21411	11	1	1	1	0	0	0	1	1	1	1	1
KN21412	11	1	1	1	0	0	0	1	1	1	1	1
KN21413	11	1	1	1	0	0	0	1	1	1	1	1
KN21420	11	1	1	1	0	0	0	1	1	1	1	1
KN21423	11	1	1	1	0	0	0	1	1	1	1	1
KN21445	11	1	1	1	1	1	1	1	1	1	1	1
KN21447	11	1	1	1	1	1	1	1	1	1	1	1
KN21458	11	1	1	0	0	0	0	1	1	1	1	1
KN21466	11	1	1	1	1	1	1	1	1	1	1	1
KN21507	11	1	1	1	1	1	1	1	1	1	1	1
KN21511	11	1	1	1	1	1	1	1	1	1	1	1
KN21556	11	1	1	1	1	1	1	1	1	1	1	1
KN21570	11	1	1	1	0	0	0	1	1	1	1	1
KN21932	11	1	1	1	0	0	0	1	1	1	1	1
KN21967	11	1	1	0	1	1	1	1	1	1	1	1

