

Supporting Information

Colistin Resistance Mechanisms in Human and Veterinary *Klebsiella pneumoniae* Isolates

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Supplementary Table S1. Primers used for detection and Sanger sequencing of beta-lactamase genes and mutations in the genes *mgrB*, *pmrAB* and *phoPQ*.

Name	Sequence (5'→3')	Gene	Size [bp]	Annealing temperature [°C]	Ref.
CTX-M_F	TTAGGAARTGTGCCGCTGYA	<i>bla</i> _{CTX-M-1-like}	688	60	11
CTX-M_R	CGATATCGTTGGTGGTRCCAT				
SHV_schles_F	TTTATCGGCCYTCACTCAAGG	<i>bla</i> _{SHV}	776	50	12
SHV_schles_R	AGATAAATCACCACAATGCGC				
<i>mgrB</i> _can_f	AAGGCGTTCATTCTACCAC	<i>mgrB</i>	253	56	9
<i>mgrB</i> _can_R	TTAAGAAGGCCGTGCTATCC				
<i>mgrB</i> _new_1F	TTTGTATGATCCCTGGCGTGA	<i>mgrB</i>	269	56	This study
<i>mgrB</i> _new_1R	AATAGTGCAAATGCCGCTGA				
<i>mgrB</i> _new_2F	ACTGATAGTCATCATAGCAG	<i>mgrB</i>	344	56	This study
<i>mgrB</i> _new_2R	AGCAGCGCTTGCAGCAGCTC				
<i>pmrA</i> _F	CTCCACATGTTTCTGGTTGT	<i>pmrA</i>	960	58	This study
<i>pmrA</i> _R	TCGCAGGATAATCTGTTCTC				
<i>pmrB</i> _F	TTCTCTTATCGTCCTGCTTG	<i>pmrB</i>	1411	58	This study
<i>pmrB</i> _R	CAATACCCTGGAAGTGCATA				
<i>phoP</i> _F	GCAGGTGTCTGACAGGGATT	<i>phoP</i>	1467	58	This study
<i>phoP</i> _R	AGGCTGAATACCCACAGGAC				
<i>phoQ</i> _F	ATGACAGCGGGAAGATATGC	<i>phoQ</i>	672	58	This study
<i>phoQ</i> _R	GAGCGTCAGACTACTATCGA				

Supplementary Table S2. Presence of antibiotic resistance genes in clinical *Klebsiella pneumoniae* isolates as determined by ResFinder (<https://cge.food.dtu.dk/services/ResFinder/>). The selected threshold for identification was 98% with a minimum length of 80%. All isolates harboured *oqx*A, *oqx*B and *fos*A.

	Veterinary isolates						Human isolates											
Antibiotic class	IHIT27665	IHIT27662	IHIT27663	IHIT27664	IHIT32358	IHIT33535	KP_03	KP_53	KP_54	KP_1153	KP_1200	KP_1377	KP_1585	KP_1710	KP_1954	KP_2442	KP_3405	KP_3996
Beta-lactam <i>(bla)</i> ^b	TEM-1B, SHV-28, CTX-M-1	SHV-27-like	SHV-1	SHV-36	SHV-28	TEM-1B, SHV-28, CTX-M-15	SHV-1, OXA-1	TEM-1B, SHV-1, OXA-1, CTX-M-15	SHV-11, OXA-1, DHA-1	TEM-1B, SHV-11, CTX-M-15	TEM-1B, SHV-27, OXA-1, CTX-M-15	TEM-1A, SHV-12, OXA-9, KPC-2	TEM-1B, SHV-11, OXA-1, CTX-M-15	TEM-1B, SHV-11, OXA-1, CTX-M-15	TEM-1A, OXA-9, KPC-2	TEM-1B, SHV-28, OXA-1, CTX-M-15	TEM-1A, SHV-28, OXA-1, CTX-M-15	SHV-1, KPC-3
Aminoglycoside	<i>aph(4)-la</i> , <i>aac(3)-IV</i>	-	-	-	<i>strA</i> , <i>strB</i> , <i>aadA2</i>	<i>aac(3)-IIa</i> , <i>strA</i> , <i>strB</i>	<i>aac(6')/lb-cr</i> , <i>aac(3)-IIa</i> , <i>strA</i> , <i>aadA2</i>	<i>aadA2</i> , <i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i> , <i>b-cr</i> , <i>aph(3')-la</i>	<i>aac(6')-lb</i> , <i>aph(6)-Id</i> , <i>aph(3')-lb</i>	<i>aac(3)-IIa</i> , <i>aac(6')/lb-cr</i> , <i>aph(3')-la</i> , <i>aadA2</i>	<i>aph(3')-la</i> , <i>aac(6')-lb</i> , <i>aadA2</i>	<i>aadA2</i> , <i>aac(6')/lb-cr</i> , <i>aac(3)-IId</i> , <i>aph(3')-la</i>	<i>aac(6')/lb-cr</i> , <i>aadA2</i>	<i>aph(3')-la</i> , <i>strA</i> , <i>aadA2</i> , <i>aac(6')-lb</i>	<i>aac(6')/lb-cr</i> , <i>aac(3)-IIa</i> , <i>aph(3')-la</i> , <i>aadA1</i> , <i>aadA2</i>	<i>aac(3)-IId</i> , <i>aac(6')/lb-cr</i>	<i>armA</i>
Fluoroquinolone	-	-	-	-	-	<i>qnrB1</i>	<i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i> , <i>qnrB4</i>	<i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i> , <i>qnrB4</i>	<i>aac(6')/lb-cr</i> , <i>qnrB4</i>	<i>aac(6')/lb-cr</i>	<i>aac(6')/lb-cr</i>	-
Trimethoprim	-	-	-	-	<i>dfrA12</i>	<i>dfrA14</i>	<i>dfrA12</i>	<i>dfrA12</i>	-	<i>dfrA12</i>	<i>dfrA12</i>	<i>dfrA12</i>	<i>dfrA30</i>	<i>dfrA1</i>	<i>dfrA14</i> , <i>dfrA12</i>	<i>dfrA12</i>	<i>dfrA14</i> , <i>dfrA17</i>	-
Sulphonamide	-	-	-	-	-	<i>sul2</i>	<i>sul1</i> , <i>sul2</i>	<i>sul1</i>	<i>sul1</i>	<i>sul2</i>	<i>sul1</i>	<i>sul1</i>	<i>sul1</i>	<i>sul1</i>	<i>sul1</i> , <i>sul2</i>	<i>sul1</i> , <i>sul3</i>	-	-
Tetracycline	-	-	-	-	<i>tet(A)</i>	<i>tet(A)</i>	<i>tet(A)</i>	<i>tet(A)</i>	-	<i>tet(A)</i>	<i>tet(A)</i>	-	<i>tet(D)</i>	<i>tet(A)</i>	-	<i>tet(A)</i>	-	-
MLS^a	<i>mph(A)</i> , <i>mph(B)</i>	-	-	-	-	-	<i>mph(A)</i>	<i>mph(A)</i>	<i>mph(A)</i>	<i>mph(A)</i>	<i>mph(A)</i>	-	<i>mph(A)</i>	<i>mph(A)</i>	<i>mph(A)</i>	<i>mph(A)</i>	-	<i>msr(E)</i> , <i>mph(E)</i>
Phenicol	<i>floR</i>	-	-	-	<i>catA1</i>	-	<i>catA1</i> , <i>catB4</i>	<i>catB4</i>	<i>catB3</i>	<i>catA2</i>	<i>catB4</i>	<i>catA1</i>	<i>catA1</i> , <i>catB4</i>	<i>catA1</i> , <i>catB4</i>	<i>catA1</i>	<i>catA1</i> , <i>catA4</i> , <i>floR</i> , <i>cmIA1</i>	<i>catB4</i>	-

^aMLS, Macrolide-lincosamide-streptogramin B; ^b ESBLs and carbapenemases are indicated in bold.

Supplementary Table S3. Minimum inhibitory concentrations (MICs) for clinical *Klebsiella pneumoniae* isolates. All MICs were determined by antibiotic gradient tests unless otherwise indicated.

	Veterinary isolates						Human isolates											
Antibiotic agent	IHIT27662	IHIT27663	IHIT27664	IHIT27665	IHIT32358	IHIT33535	KP_03	KP_53	KP_54	KP_1153	KP_1200	KP_1377	KP_1585	KP_1710	KP_1954	KP_2442	KP_3405	KP_3996
Ampicillin	128	64	64	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256
Ampicillin/sulbactam ^a	2	2	2	>256	>256	16	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256
Piperacillin	2	8	8	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256
Cefuroxime	4	4	2	>256	4	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256	>256
Cefepime ^b	0.064	0.032	0.032	16	0.125	2	32	>256	8	>256	>64	>64	>256	48	>256	>64	>256	>256
Ertapenem	0.016	0.016	0.016	8	0.032	0.064	0.125	>32	>32	>32	12	32	>32	0.125	>32	0.25	>32	>32
Gentamicin	0.5	0.5	0.5	8	1	32	32	2	1	2	>256	2	>256	1	2	16	16	>256
Levofloxacin ^b	≤0.125	≤0.125	≤0.125	4	>8	>8	1	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32
Tigecycline	0.5	0.5	0.5	4	2	1	0.5	2	0.5	1	8	1	2	2	4	2	2	2
Colistin ^c	8	8	8	8	4	2	8	8	8	4	4	4	4	8	16	2	2	2
Colistin ^d	24	12	24	24	12	32	24	48	256	256	6	12	16	256	256	6	8	12
Colistin ^e	>16	>16	>16	>16	>16	>16	8	>16	>16	>16	8	>16	>16	>16	>16	8	>16	8

^aSulbactam was used at a fixed concentration of 4 mg/L as recommended by EUCAST.

^bMICs were determined by either antibiotic gradient test or VITEK[®] 2 analysis.

^cMICs were determined using ETEST[®] and Mueller-Hinton agar (Oxoid).

^dMICs were determined using ETEST[®] and Mueller-Hinton E agar (bioMérieux).

^eMICs were determined by VITEK[®] 2 analysis.

PmrH

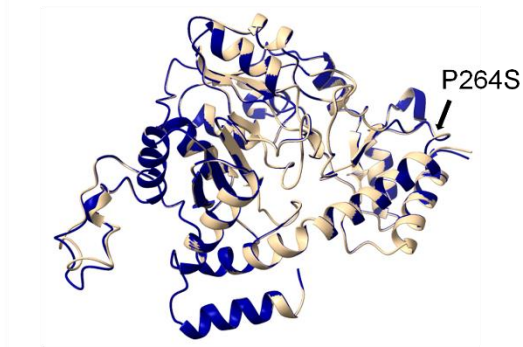


Figure S1 Superposition of PmrH of *K. pneumoniae* MGH78578 in blue (accession number WP_004889389.1) and PmrH of KP_03 in beige. Amino acid substitution P264S is marked by an arrow.

PmrB

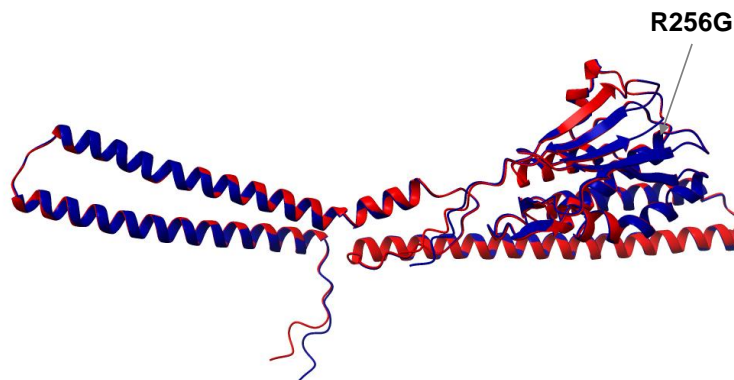


Figure S2 Superposition of PmrB of *K. pneumoniae* MGH78578 in blue and PmrB (R256G) of IHIT27665 in red. The arrow marks amino acid substitution R256G.

PmrB

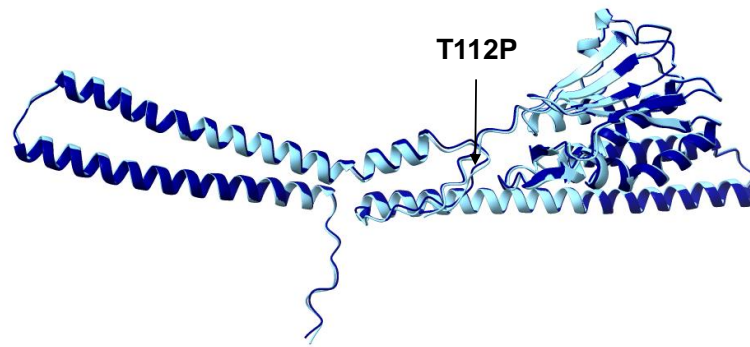


Figure S3 Superposition of PmrB of *K. pneumoniae* MGH78578 in dark blue and PmrB (T112P) of KP_54 (light blue). The arrow marks amino acid substitution T112P.