



Supplemental Figure S1.

Radial Dendrogram produced by MLST on the basis of seven housekeeping genes (*gapA*, *infB*, *mdh*, *pgi*, *phoE*, *rpoB*, *tonB*). Allele sequences were concatenated to generate a 3,003 bp concatemer for each strain. A phylogenetic tree was constructed in MEGA ver. 6 using UPGMA. Distances were computed using Maximum Composite Likelihood and the reliability of the tree was assessed by bootstrap analysis with 1,000 replicates. Sequences of 113 isolates present in the PubMLST database were included in the analysis. The isolates analysed are displayed in bold.

Table S1. Allelic profiles of *Klebsiella* sp. isolates and reference strains analysed by MLST

	<i>gapA</i>	<i>infB</i>	<i>mdh</i>	<i>pgi</i>	<i>phoE</i>	<i>rpoB</i>	<i>tonB</i>	cluster
<i>K. oxytoca</i> PHS-890	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-891	3	5	21	13	20 (⁶² C→T)	6	12	B1

<i>K. oxytoca</i> PHS-892	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-893	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-894	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-895	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-896	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-897	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-898	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> PHS-899	3	5	21	13	20 (⁶² C→T)	6	12	B1
<i>K. oxytoca</i> CB4063	5	6	19	10	46	5	9 (⁷⁵ A→G ⁷⁸ G→A)	B2
<i>K. oxytoca</i> CB4074	1	2	2	1	4	1	2	A
<i>K. oxytoca</i> CB4072	3	4	15 (³³⁰ A →T)	8	25	3	4	B1
<i>K. oxytoca</i> Oman 61	1	3	13	24	2 (³²¹ G →A)	10	10	A
<i>K. oxytoca</i> ATCC 13182	2	2	2	3	19	2	2	A
<i>K. oxytoca</i> CCUG 15788	16 (16 diff.)	25 (20 diff.)	39 (49 diff.)	23 (20 diff.)	15 (35 diff.)	24 (17 diff.)	34 (55 diff.)	out