

Supplementary Table 1: Resistance mechanisms and carbapenem MICs of *Enterobacteriaceae* yielding false, doubtful or non-interpretable results with one of the four evaluated assays.

Species	Carbapenemase production	Other resistance mechanisms	MIC (µg/ml) Imipenem	MIC (µg/ml) Meropenem	MIC (µg/ml) Ertapenem	BYG	BCT	RCNP	NRCK
<i>K. pneumoniae</i>	None	DHA	0,25	0,047	0,38	P	N	N	N
<i>E. cloacae</i>	None	AmpC overexpression and phenotypically assessed impermeability	0,75	0,25	2	N	D	N	N
<i>K. oxytoca</i>	None	K-OXY overexpression	0,19	0,064	0,032	N	D	N	N
<i>K. pneumoniae</i>	KPC-3		> 32	> 32	> 32	P	D	P	P
<i>K. pneumoniae</i>	OXA-48	DHA, CTX-M9	0,75	0,5	2	P	D	P	P
<i>E. coli</i>	OXA-48		3	0,5	0,5	P	D	P	P
<i>E. cloacae</i>	None	AmpC overexpression and phenotypically assessed impermeability	1	0,19	0,5	N	N	D	N
<i>K. pneumoniae</i>	None	DHA	0,38	0,047	0,5	N	N	D	N
<i>K. pneumoniae</i>	None	ESBL and phenotypically assessed impermeability	0,5	1,5	6	N	N	D	N
<i>K. pneumoniae</i>	None	CTX-M1 and phenotypically assessed impermeability	0,75	3	24	N	N	D	N
<i>K. pneumoniae</i>	None	CTX-M1 and phenotypically assessed impermeability	1,5	6	>32	N	N	D	N
<i>E. coli</i>	None	CTX-M1	0,25	0,032	0,006	N	N	D	N
<i>E. aerogenes</i>	None	AmpC overexpression	0,5	0,38	1,5	N	N	D	N

<i>K. pneumoniae</i>	None	DHA	0,25	0,094	0,38	N	N	D	N
<i>K. pneumoniae</i>	None	ESBL and phenotypically assessed impermeability	0,25	1	3	N	N	D	N
<i>E. cloacae</i>	None	AmpC overexpression and phenotypically assessed impermeability	2	1	2	N	N	D	N
<i>K. pneumoniae</i>	None	Phenotypically assessed impermeability	1	2	24	N	N	D	N
<i>E. coli</i>	None	CMY-2, ESBL (CTX-M1)	0,38	0,047	0,25	N	N	D	N
<i>E. cloacae</i>	None	AmpC overexpression, ESBL and phenotypically assessed impermeability	2 (0,5)	0,25	1	N	N	D	N
<i>E. cloacae</i>	OXA-48	CTX-M1	1,5	0,75	3	P	P	D	P
<i>K. pneumoniae</i>	OXA-48	CTX-M1	3	1	1,5	P	P	D	P
<i>K. pneumoniae</i>	OXA-48	CTX-M1	0,38	0,38	0,75	P	P	D	P
<i>E. coli</i>	OXA-48	CTX-M1	1,5	0,25	0,38	P	P	D	D
<i>E. coli</i>	OXA-48		3	0,5	1,5	P	P	D	D
<i>K. pneumoniae</i>	OXA-48		1,5	0,38	0,38	P	P	D	D
<i>K. pneumoniae</i>	OXA-48	ESBL	2	1	0,75	P	P	D	D
<i>E. cloacae</i>	OXA-48	CTX-M9	2	2	6	P	P	D	D
<i>E. cloacae</i>	OXA-48	CTX-M9, SHV	24	32	> 32	P	P	D	D
<i>K. pneumoniae</i>	None	ESBL	0,25	0,38	1	N	N	N	D

<i>E. coli</i>	None	CTX-M1 and phenotypically assessed impermeability	1	3	16	N	N	N	D
<i>E. aerogenes</i>	None	ESBL and phenotypically assessed impermeability	16	4	> 32	N	N	N	D
<i>S. marcescens</i>	None	AmpC overexpression and phenotypically assessed impermeability	6	3	> 32	N	N	N	D
<i>K. pneumoniae</i>	OXA-48	CTX-M1	0,75	0,5	1,5	P	P	P	D
<i>K. pneumoniae</i>	OXA-48	CTX-M1	0,75	0,5	1	P	P	P	D
<i>K. pneumoniae</i>	OXA-48		1,5	1,5	1	P	P	P	D
<i>K. pneumoniae</i>	None	ESBL	0,125	0,016	0,023	N	N	D	NI
<i>K. pneumoniae</i>	None	DHA and phenotypically assessed impermeability	6	2	> 32	N	N	N	NI
<i>K. pneumoniae</i>	None	DHA	0,75	0,19	0,38	N	N	N	NI
<i>K. pneumoniae</i>	OXA-48	CTX-M1	1	0,5	1,5	P	P	P	NI
<i>E. cloacae</i>	None	AmpC overexpression	0,5	0,38	0,38	N	N	N	NI
<i>E. cloacae</i>	None	AmpC overexpression and phenotypically assessed impermeability	0,5	0,25	1,5	N	N	N	NI
<i>E. coli</i>	OXA-48	CTX-M1	1,5	0,19	0,38	P	P	P	N
<i>K. pneumoniae</i>	OXA-48	CTX-M1	1	1,5	1,5	P	P	P	N

P = positive ; N = negative; D = doubtful ; NI = non-interpretable