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Rapid Phenotypic Detection of AmpC β-Lactamases, Extended-Spectrum- β-Lactamases and Metallo-β-Lactamases in Enterobacteriaceae Using A Resazurin Microtitre Assay with Inhibitor-Based Methods --Manuscript Draft--

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Abstract:	Dissemination of antibiotic resistance in Enterobacteriaceae mediated by AmpC, ESBL and MBL β -lactamases is clinically significant. A simple, rapid method for the detection of these resistance phenotypes would greatly improve chemotherapeutic recommendation. This technology would provide valuable input in our surveillance of resistance on a global stage, particularly if the methodology could be applicable to resource poor settings. A resazurin microtitre plate (RMP) assay incorporating cloxacillin (CX), clavulanic acid (CA), and EDTA for the rapid phenotypic identification of AmpC, ESBL, MBL and the co-existence of β -lactamases has been developed. A total of 44 molecularly characterised Enterobacteriaceae clinical isolates producing AmpCs, ESBLs, co-producers of ESBL and AmpC, MBLs, and co-producers of ESBL and MBL were phenotypically examined using the RMP assay. The ceftazidime (CAZ)-based and cefotaxime (CTX)-based RMP assay successfully detected all 15 AmpC, 12 ESBL, 9 MBL producers, 6 ESBL-AmpC co-producers, and 2 ESBL-MBL co-producers of ESBL and AmpC alone, while the CTX-based assay performed better in identifying co-producers of ESBL and AmpC. There was no difference in detection of ESBL and MBL producers. The findings of the present study suggest that use of the RMP assay with particular β -lactamase inhibitors explicitly detects three different β -lactamases, as well as co-existence of β -lactamases within 6 h. This assay is applicable to carry out in any laboratory, is cost-effective and easy to interpret. It could be implemented in screening patients, controlling infection and for surveillance purposes.					

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24	

25 Abstract

Dissemination of antibiotic resistance in Enterobacteriaceae mediated by AmpC, ESBL and 26 MBL β -lactamases is clinically significant. A simple, rapid method for the detection of these 27 resistance phenotypes would greatly improve chemotherapeutic recommendation. This 28 technology would provide valuable input in our surveillance of resistance on a global stage, 29 particularly if the methodology could be applicable to resource poor settings. A resazurin 30 31 microtitre plate (RMP) assay incorporating cloxacillin (CX), clavulanic acid (CA), and EDTA for the rapid phenotypic identification of AmpC, ESBL, MBL and the co-existence of 32 β-lactamases has been developed. A total of 44 molecularly characterised Enterobacteriaceae 33 clinical isolates producing AmpCs, ESBLs, co-producers of ESBL and AmpC, MBLs, and 34 co-producers of ESBL and MBL were phenotypically examined using the RMP assay. The 35 ceftazidime (CAZ)-based and cefotaxime (CTX)-based RMP assay successfully detected all 36 15 AmpC, 12 ESBL, 9 MBL producers, 6 ESBL-AmpC co-producers, and 2 ESBL-MBL 37 co-producers without false positive results. The CAZ-based assay was more reliable in 38 detecting AmpC alone, while the CTX-based assay performed better in identifying co-39 producers of ESBL and AmpC. There was no difference in detection of ESBL and MBL 40 producers. The findings of the present study suggest that use of the RMP assay with 41 42 particular β -lactamase inhibitors explicitly detects three different β -lactamases, as well as coexistence of β -lactamases within 6 h. This assay is applicable to carry out in any laboratory, is 43 cost-effective and easy to interpret. It could be implemented in screening patients, controlling 44 infection and for surveillance purposes. 45

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49 Introduction

Infections caused by multidrug-resistant Enterobacteriaceae are among one of the most 50 serious threats to human health (Tangden & Giske 2015). The production of β -lactamases 51 52 mediated by both chromosomal and plasmid genes are crucial mechanisms of antibiotic resistance. AmpC β-lactamases were the first bacterial enzymes found to inactivate penicillin, 53 first identified in Escherichia coli in 1940 prior to penicillins being medically introduced 54 55 (Jacoby, 2009). Plasmid-mediated AmpC β-lactamase (pAmpC)-producing Enterobacteriaceae can confer resistance to multiple antibiotics, such as penicillins, 56 oxyimino-cephalosporins (e.g. cefotaxime, ceftazidime and ceftriaxone) 57 and 7- α methoxycephalosporins (e.g. cefoxitin and cefotetan) (Gupta et al., 2012). CMY-type β -58 lactamases, especially CMY-2 are the highest prevalence of pAmpC in E. coli that have been 59 60 commonly identified in many regions of the world (Doi et al., 2010; Liebana et al., 2013). These enzymes are inhibited by boronic acid (BA), and its derivatives (Beesley et al., 1983) 61 and cloxacillin (CX) (Jacoby, 2009). AmpC β-lactamases are poorly inhibited by clavulanic 62 63 acid (CA) (Bush et al., 1995).

64 Enterobacteriaceae can also produce powerful enzymes called extended-spectrum βlactamases (ESBLs), which characteristically confer resistance to penicillins, cephalosporins, 65 66 and monobactams, but remain susceptible to β -lactamase inhibitors (CA, tazobactam, sulbactam) cephamycins and carbapenems (Bradford, 2001). The majority of ESBL-67 producing clinical isolates of Enterobacteriaceae harbour TEM-, SHV-, and CTX-M-type β-68 lactamases. These enzymes have gained medical significance in both hospital and community 69 settings, across all ages and demographics (Lukac et al., 2015). Furthermore, carbapenem-70 inactivating β -lactamase, namely metallo- β -lactamase (MBL) has been identified with 71 72 increasing prevalence over the past few years (Patrice et al., 2011). MBL can hydrolyse all classes of bicyclic β-lactams (penicillins, cephalosporins and carbapenems) and all 73

74 commercially available β-lactamase inhibitors (such as CA), but lack hydrolytic activity toward monocyclic β -lactams (monobactams) (Bebrone, 2007). The activity of this enzyme is 75 facilitated by zinc-ions to degrade β-lactams and is inhibited by metal-chelating 76 ethylenediaminetetraacetic acid (EDTA) and dipicolinic acid (Walsh et al., 2005). 77 Imipenemase (IMP), Verona integron-encoded metallo-β-lactamase (VIM), and New Delhi 78 metallo-β-lactamase (NDM) are common widespread MBLs. These metalloenzymes were 79 frequently identified in *Pseudomonas aeruginosa* and *Acinetobacter* spp., but currently an 80 increasing emergence of these have been documented in Enterobacteriaceae worldwide (Patel 81 82 & Bonomo,2013).

The early detection of the presence of these resistant strains at the phenotypic level would aid 83 infection-control practitioners to minimise the dissemination of these bacteria and also help to 84 85 select appropriate antimicrobial treatment (Pitout & Laupland, 2008). The development of a rapid method for the identification of these β-lactamase producing strains in clinical practice 86 is a challenging issue. The resazurin (Alamar blue) reduction assay is a simple and 87 88 inexpensive colorimetric method that has extensively been used as an indicator for cell growth, cell viability, toxicity and indirect antimicrobial susceptibility testing. This dye is 89 non-toxic to cells and stable in culture media. A blue coloured resazurin is irreversibly 90 converted to a pink coloured resorufin by active cells (O'Brien et al., 2000; Palomino et 91 al.,2002). The present study uses the resazurin dye in conjunction with three distinct β -92 93 lactamase inhibitors, CX, CA, and EDTA, for the phenotypic detection of AmpC, ESBL, MBL and co-expression of β -lactamases. 94

95 Methods

96 Bacterial Isolates, Chemicals and Antibiotics

97 A total of 44 Enterobacteriaceae clinical isolates were used in this study. The bacterial strains included E. coli (n=13), Klebsiella pneumoniae (n=16), Enterobacter aerogenes (n=2), 98 Enterobacter cloacae (n=8), Morganella morganii (n=1), Citrobacter freundii (n=3), and 99 100 *Klebsiella ozaenae* (n=1); 15 were AmpC producers, 12 were ESBL producers, 6 were ESBL and AmpC co-producers, 9 were MBL producers, and 2 were ESBL and MBL co-producers 101 (Table 1). The control strains were E. coli ATCC 25922 (non-β-lactamase producer), E. 102 cloacae ATCC BAA-1143 (blaACT-32), E. coli NCTC 13352 (blaTEM-10) and E. coli NCTC 103 13353 (bla_{CTX-M-15}). All organisms had been identified by biochemical, phenotypic and 104 105 molecular tests including combined disc, PCR and sequencing methods as previously described (Pérez-Pérez & Hanson, 2002; Woodford, 2010; Roschanski et al., 2014). All 106 antibiotics and chemicals employed were purchased from Sigma Aldrich, UK. Mueller-107 108 Hinton Broth (MHB) powder was obtained from Scientific Laboratory Supplies, UK. Antibiotics and β -lactamase inhibitors were prepared in accordance with the recommendation 109 of manufacturers. 110

111 Resazurin Microtitre Plate Assay (RMP)

The production of ESBL, AmpC, MBL and co-β-lactamases was phenotypically detected 112 using the RMP assay based upon the broth microdilution method. An overnight culture of the 113 appropriate test bacterium was pelleted for 10 min and washed twice with normal saline at 114 4,000 rpm for 5 min before adjustment to a 0.5 McFarland suspension according to the 115 recommendations of the Clinical Laboratory Standards Institute (CLSI) in document M07-A9 116 (Clinical Laboratory Standards Institute, 2012). An aliquot (20 µL) of a 5 x 10⁶ CFU/mL of 117 118 organism were added to the wells containing 180 µL of MHB, amended with resazurin at a concentration of 20 µg/mL, and serial twofold dilution of cefotaxime (CTX) or ceftazidime 119 (CAZ) with a fixed concentration of β -lactamase inhibitors; CX at 200 µg/mL, EDTA at 0.5 120 121 mM, and CA at 4 µg/mL. Following incubation at 37 °C, the lowest concentration showing no colorimetric change from blue to pink within 6 h was considered as the MIC value.
Interpretation criteria of the result obtained is explained in Table 2. The experiment was
carried out in three independent replicates.

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126 **Results and Discussion**

127 β -Lactam antibiotics are most frequently prescribed for the treatment of infections caused by Gram negative bacteria, in particular Enterobacteriaceae. Resistance to β-lactams is mediated 128 129 by both plasmid- and chromosomally-encoded β -lactamases. The rapid global spread of these resistant bacteria is facilitated by inter- and intra-species gene transmission, poor sanitation 130 and hygiene, increasing global trade, international travel and tourism, and antibiotic misuse 131 (Rogers et al., 2011; Aung et al., 2012; Laxminarayan et al., 2013). The consequence is 132 increased treatment failure and health risks. The development of rapid and accurate methods 133 for the detection and characterisation of different types of β-lactamase remains an urgent 134 necessity. The early detection is crucial in guiding appropriate antimicrobial therapy. The 135 common conventional detection methods for the production of β -lactamase are still time-136 consuming (approximately 24 to 48 h) (Nordmann et al., 2012). Here, we developed the RMP 137 assay with inhibitor-based methods for the early phenotypic detection of ESBL, AmpC, 138 MBL, and co-production of β -lactamases. 139

The findings from the RMP assay were very simple and clear to interpret as illustrated in Figure 1. The results showed that the RMP CAZ-based assay was more reliable than the CTX-based assay for early AmpC detection. An eightfold or higher MIC ratio was observed when comparing the well without CX to other wells only for AmpC producers. Other MIC ratios revealed values lower than 8, which indicated negative results for ESBL and MBL producers. The CAZ-based assay successfully detected 15 of all 15 AmpC-producing 146 isolates, while 14 of 15 strains were detected by the CTX-based RMP assay. The AmpC control strain clearly showed hyperproduction of AmpC β-lactamase indicated by a high MIC 147 value in the well without an AmpC inhibitor (Table 1). The production of AmpC β -lactamase 148 149 in Gram negative pathogens, especially Enterobacteriaceae is clinically important. Accurate and rapid detection methods for these enzymes should be urgently developed. This would 150 ensure effective antimicrobial therapy and improve clinical management as a consequence of 151 clinical failure in the treatment of AmpC producers with broad-spectrum cephalosporins 152 (Polsfuss *et al.*,2011). Cloxacillin is well-described as an AmpC β -lactamase inhibitor, but it 153 154 cannot distinguish between plasmid-mediated and chromosome-borne AmpC (Jacoby, 2009). This study found that CAZ in conjunction with CX and EDTA was more sensitive for early 155 detection of AmpC-producing Enterobacteriaceae compared with the CTX-based assay. This 156 157 result is consistent with a previous report where CAZ showed the best performance compared with CTX, cefmetazole, or moxalactam in combination with 3-aminophenylboronic acid 158 (APB) in detection of AmpC β -lactamases (Yagi *et al.*,2005). Another previous research 159 160 study reported no difference in sensitivity and specificity between CTX- and CAZ-based detection in the presence of CA with and without BA in detection of pAmpC (Jeong et 161 al., 2009). In this study, some AmpC producers such as, K. pneumoniae 2001 (DHA-1), E. 162 coli 2002 (DHA-1), and *M. morganii* 2010 were partially inhibited by CX as indicated by the 163 high MIC values after exposure to CX. In those isolates we suggest that the resistance was 164 165 not only mediated by AmpC β-lactamase, but might also be conferred by alteration in the permeability of the outer membrane. 166

167 The MIC ratio for wells without CA compared to wells with CA gave a value ≥8 in all tested 168 ESBL-producing Enterobacteriaceae employed in this study. The findings unambiguously 169 indicated that both CAZ-based and CTX-based assays gave reproducible results in detecting 170 all ESBL isolates. Even though CLSI has recommended a standardised confirmatory test for

ESBL-producing Enterobacteriaceae, previous reports found that using CTX and CAZ alone 171 and in combination with CA demonstrated a high percentage of false-negative results in 172 detection of ESBL producers due to co-production of β -lactamase enzymes (Jeong et 173 al.,2009). In the current study inhibitor-based methods using three different β -lactamase 174 inhibitors were applied to improve the sensitivity and specificity for phenotypic detection of 175 ESBL in clinical isolates of Enterobacteriaceae. This modification allows the successful 176 identification of all ESBLs in both CAZ-and CTX-based assays. Several techniques for 177 detecting the production of ESBL have been proposed such as, double-disc synergy test, Etest 178 179 and automated methods. The sensitivities and specificities for the double-disc test and Etest were 80% to 95%, whilst automated susceptibility methods were 80% to 99% (Drieux et 180 al.,2008; Gazin et al.,2012). In addition, the ESBL NDP (Nordmann/Dortet/Poirel) test has 181 182 been developed for rapid detection with an excellent sensitivity and specificity of 92.6% and 100%, respectively. However, this test was limited to detecting ESBL without presence of 183 AmpC or other β -lactamases (Nordmann et al., 2012). 184

185 All MBL producers were detected by the RMP assay, including 9 MBL-producing isolates and 2 co-producers of ESBL and MBL. The CTX-based assay gave similar results with the 186 CAZ-based assay for the rapid phenotypic detection of MBL producers. The MIC ratio of 187 non-\beta-lactamase-producing E. coli ATCC 25922 for both CTX- and CAZ-based assays was 188 1, suggesting no ESBL, AmpC, and MBL production in this bacterium. A raised global 189 dissemination of MBL-producing Enterobacteriaceae has drawn researchers to develop 190 reliable techniques for detection of MBLs (Hattori et al., 2013). Yan and colleagues 191 conducted double-disc, combined disc, and Etest techniques for the detection of Gram 192 negative bacilli producing MBLs. They found that the 2-mercaptopropionic acid (2-MPA) 193 double-disc test using CAZ and cefepime, with and without CA, exhibited highest sensitivity 194 (100%) compared with the combined disc test (86.7%) and Etest (36.7%). Even the double-195

196 disc test showed high sensitivity, but there were some disadvantages, including subjective interpretation of results in some samples, and also volatile and odorous properties of 2-MPA 197 (Yan et al., 2004). More recently, double-disc synergy tests using sodium mercaptoacetic 198 199 acid (SMA) and EDTA as MBL inhibitors, and CAZ, imipenem and meropenem as indicator antibiotics have emerged. The results showed that CAZ in combination with SMA can detect 200 only MBL-producing strains, whereas a combination of meropenem and SMA revealed the 201 best performance (sensitivity, 79.1%; specificity, 100%) in detection of co-production of β -202 lactamases (Hattori et al., 2013). Yagi and colleagues also suggested the broth microdilution 203 204 method incorporating CAZ or CTX with SMA for successful detection of IMP-type MBL (Yagi et al., 2005). Our findings revealed a good performance in the RMP assay for either 205 206 MBL alone or in combination with ESBL.

ESBL and AmpC co-producers were positively detected as indicated by a \geq 8 MIC ratio for 207 208 wells containing either CA or CX compared to wells containing both CA and CX. The CTXbased RMP assay was more reliable in phenotypic characterisation of the co-existence of 209 210 ESBL and AmpC β-lactamase compared with the CAZ-based RMP assay (Table 1). Co-211 production of different types of β -lactamase in the same strain are one of the most common causes of phenotypic detection failure. The most prevalent isolates co-producing pAmpCs 212 and ESBLs are K. pneumoniae and E. coli (Goossens & Grabein, 2005). These co-β-213 lactamases may result in false-negative tests for the detection of ESBLs or AmpCs 214 (Coudron, 2005). A method that can distinguish between ESBLs and pAmpCs is important for 215 both treatment and epidemiology (Song et al., 2007). Furthermore, co-expression of MBL 216 together with ESBLs (commonly CTX-M-type) and AmpC β-lactamases (mostly CMY-217 enzymes) appears to be difficult to detect. Consequently, these may be a major cause of false-218 negative tests for detection of co-MBL, AmpC and ESBL in Gram negative pathogens 219 (Hattori et al., 2013; Tangden & Giske, 2015). 220

221 Although simple phenotypic tests for ESBL, AmpC and MBL have been proposed by several researchers, and indeed conducted in clinical laboratory practice (Yagi et al., 2005; Jeong et 222 al.,2009), there is no standard guideline for phenotypic detection of AmpC that has been 223 224 recommended by the CLSI (Clinical Laboratory Standards Institute, 2012). Broth microdilution methods have become more popular in clinical diagnostic laboratories since 225 semi-automated antimicrobial susceptibility test systems have become commercially 226 227 available. This test generally involves overnight incubation in the determination of antibiotic susceptibility profiling (Yagi et al., 2005). The present study using the RMP assay allows for 228 229 the detection of ESBLs, AmpCs, MBLs and co-producers within 6 h. RMP assays could allow clinicians to detect these enzymes in rural clinical laboratories in resource poor 230 settings. 231

Phenotypic tests for β-lactamase production are carried out in most clinical diagnostic 232 233 laboratories because they are easy to implement and are cost effective. This detection is important in terms of epidemiological purposes and to restrict the dissemination of β -234 235 lactamase-mediated resistance mechanisms (Poulou et al., 2014). Phenotypic tests alone cannot distinguish between the specific bla genes of β-lactamases (TEM, SHV, CTX for 236 ESBLs, CMY, MOX for AmpCs, and IMP, NDM, VIM for MBLs) (Pitout & Laupland, 2008; 237 Pitout et al., 2010), therefore, several reference laboratories use molecular methods for 238 identification of those *B*-lactamase-encoded genes. Molecular approaches are restricted 239 largely to reference laboratories for epidemiological studies because of the complexity and 240 diversity of distinct point mutations of β -lactamases (Sundsfjord *et al.*, 2004). 241

Although we used small samples of clinical isolates, a previous study found that use of CTX or CAZ combined with CA for detecting ESBL and CTX or CAZ plus BA for identifying pAmpC demonstrated 100% in both sensitivity and specificity (Jeong et al.,2009). In our study, the RMP assay detected all ESBL, AmpC, and ESBL-AmpC co-producers. Therefore, 246 we expect that our RMP assay could reach 100% in sensitivity and specificity similar to previous reports when a larger number of clinical isolates are tested. This assay can 247 successfully distinguish co-expression of ESBL and AmpC or ESBL and MBL in the same 248 strain. In future work, isolates with other resistance mechanisms, including alteration of 249 permeability of bacterial membrane and modification of membrane proteins should be 250 included in total isolates screened with RMP assay. The early detection of β -lactamase 251 enzymes may reduce a delay in the initiation of appropriate antimicrobial therapy and could 252 produce effective treatment. RMP assay is a useful method that can detect and characterise 253 254 ESBL, AmpC, MBL and co-production of ESBL and AmpC or ESBL and MBL within 6 h. It could be implemented in microbiological diagnostic laboratories in resource poor areas to 255 help clinicians screen patients, control infection, and provide epidemiological surveillance. 256

257

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266 Transparency declaration

267 The authors have no conflict of interest to declare.

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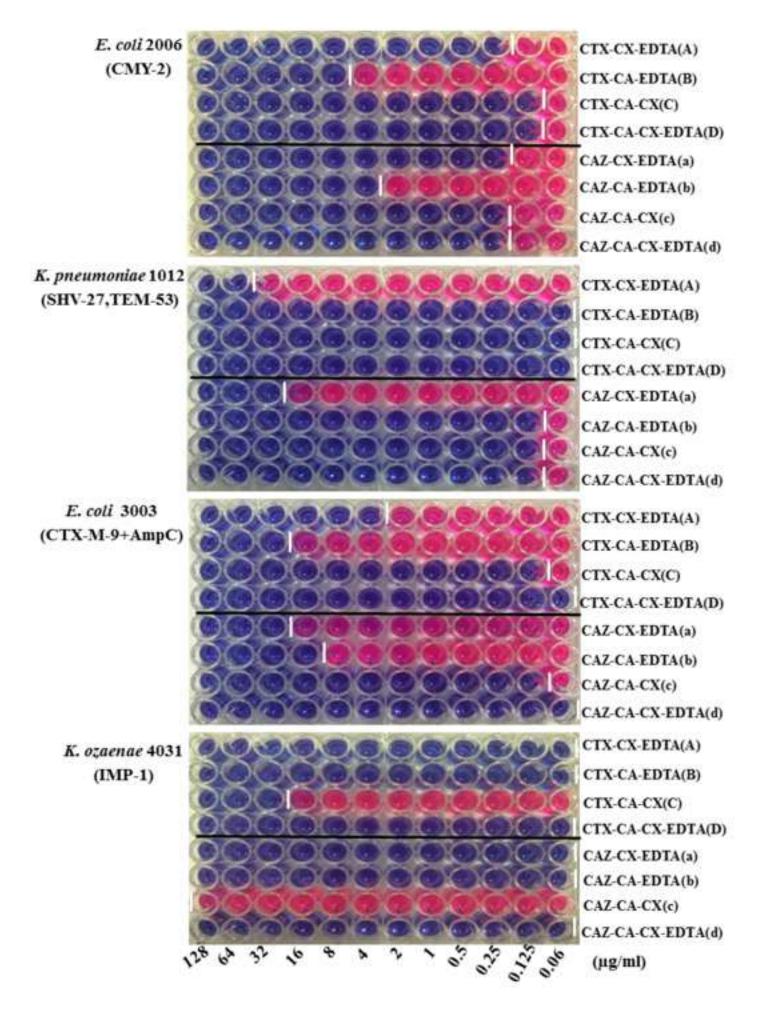
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390	Figure 1 Schematic representation of results from resazurin microtitre plate assay at 6

h. CTX = cefotaxime; CAZ=ceftazidime; CA=clavulanic acid (4 μ g/ml); CX = cloxacillin

392 (200 μ g/ml); EDTA = ethylenediaminetetraacetic acid (0.5 mM).



Strains	Resistance	MICs of CTX -based (µg/ml)				MICs of CAZ -based (µg/ml)				Phenotypic
	Mechanism	А	В	С	D	a	b	с	d	results
*E. coli ATCC 25922	Negative control	≤0.06	≤0.06	≤0.06	≤0.06	≤0.06	≤0.06	≤0.06	≤0.06	None
K. pneumoniae 2001	DHA-1	4	≥256	8	4	8	≥256	16	8	AmpC
E. coli 2002	DHA-1	4	32	4	4	16	≥256	16	16	AmpC
E. coli 2003	AmpC	≤0.06	0.25	≤0.06	≤0.06	0.125	1	0.125	0.125	AmpC
E. coli 2004	CMY-2	≤0.06	32	≤0.06	≤0.06	≤0.06	64	≤0.06	5 ≤0.06	AmpC
E. coli 2005	AmpC	≤0.06	2	≤0.06	≤0.06	0.125	4	0.125	0.125	AmpC
E.coli 2006	CMY-2	0.25	8	0.125	0.125	0.25	4	0.25	0.25	AmpC
E. aerogenes 2007	ACT-31	≤0.06	32	≤0.06	≤0.06	≤0.06	4	≤0.06	5 ≤0.06	AmpC
E. cloacae 2008	FOX	≤0.06	16	≤0.06	≤0.06	0.125	32	0.125	0.125	AmpC
E. cloacae 2009	ACR-32	0.125	64	0.125	0.125	0.125	32	0.125	0.125	AmpC
M. morganii 2010	AmpC	2	64	2	2	8	≥256	8	8	AmpC
E. cloacae 2011	AmpC	≤0.06	64	≤0.06	≤0.06	≤0.06	32	≤0.06	≤0.06	AmpC
C. freundii 2026	CMY-112	≤0.06	32	≤0.06	≤0.06	≤0.06	32	≤0.06	5≤0.06	AmpC
K. pneumoniae 2020	FOX	≤0.06	2		≤0.06	0.5	16	0.5	0.5	AmpC
<i>E. coli</i> 2021	FOX-3	≤0.06	0.5	≤0.06	≤0.06	≤0.06	8	0.125	0.125	AmpC
E. cloacae 2022	ACT-32	4	128	4	4	2	≥256	2	2	AmpC
*E. cloacae BAA-1143	ACT-32	4	128	4	4	2	128	4	4	AmpC
<i>E. coli</i> 1004	TEM-214	64	0.125	0.125	0.125	16	0.25	0.25	0.125	ESBL
E. coli 1005	CTX-M-3	1	≤0.06		≤0.06	0.5	≤0.06		5≤0.06	ESBL
E. coli 1006	CTX-M-3	64	0.125	 ≤0.06		1	0.125	0.125	0.125	ESBL
E. coli 1008	CTX-M-3	≥256	0.25	0.125	0.125	64	1	0.5	0.5	ESBL
E. coli 1009	TEM	32	0.125	≤0.06		4	0.125	0.125	0.125	ESBL
K. pneumoniae 1010	CTX-M-15,SHV-27	64	≤0.06		_0.06	16	≤0.06	≤0.06	≤0.06	ESBL
K. pneumoniae 1012	SHV-27,TEM-53	64	_0.06		_0.06	32	0.125	0.125	0.125	ESBL
K. pneumoniae 1012	SHV-27	32	_0.06	_0.06	_0.06	32	0.5	0.5	0.5	ESBL
K. pneumoniae 1015	SHV-27,TEM-71	16	< 0.06		_0.06	8	0.125	0.125	0.125	ESBL
* <i>E. coli</i> 13352	TEM-10	1	_0.06		_0.06	128	0.5	0.5	0.5	ESBL
*E. coli 13353	CTX-M-15	128	_0.06		_0.06	32	≤0.06	≤0.06	≤0.06	ESBL
K. pneumoniae 1021	SHV-18	128	_0.06		_0.06	32	_0.06	_	_0.06	ESBL
K. pneumoniae 1021	SHV-27,TEM-115	32	_0.06		_0.06	16	_0.06 ≤		5 <u>≤</u> 0.06	ESBL
K. pneumoniae 1022 K. pneumoniae 1023	SHV-27, TEM-115	52 64	<u>_</u> 0.00		<u>_</u> 0.00 ≤0.06	16	<u>_</u> 0.00 ≤0.06		i ≤0.06	ESBL
E. cloacae 3007	CTX-M-9, MIR-1 ACT-1	16	<u>_0.00</u> 64		<u>_</u> 0.06	10	32			AmpC+ESBL
E. cloacae 3009	TEM-214,SHV-12+ ACT	16	1		<u>_</u> 0.00 ≤0.06	4	0.25			AmpC+ESBL
E. aerogenes 3022	SHV-12,CTX-M-9+ ACT-32		32		<u>_</u> 0.00 ≤0.06	0.5	2	0.25	0.25	
<i>E. coli</i> 3003	CTX-M-9+AmpC	4	32		<u>_</u> 0.00 ≤0.06	32	16	0.125		AmpC+ESBL
<i>C. freundii</i> 3004	CTX-M-3+AmpC	2	32		<u>≤</u> 0.00	0.25	4	≤0.06		AmpC+ESBL AmpC+ESBL
C. freundii 3005	CTX-M-3+ ACT-1	4	32		<u>≤</u> 0.00	32	4	<u> </u>	0.25	AmpC+ESBL AmpC+ESBL
K. ozaenae 4031		≠ ≤0.06	0.125			0.5				MBL
K. pneumoniae 4031	IMP-1			32	≤0.06		0.5	128	≤0.06	
K. pneumoniae 4033 K. pneumoniae 4034	VIM-1+SHV-102 VIM-1+SHV-12	4	≤0.06	32	≤0.06	128	≤0.06 ≤0.06	≥256	≤0.06	
		8	≤0.06	8 >256	≤0.06	128	≤0.06	128	≤0.06	
K. pneumoniae 4036	NDM-1	≤0.06 0.125	≤0.06 0.125	≥256	≤0.06 0.125	≤0.06	≤0.06	≥256	≤0.06	MBL
K. pneumoniae 4025	VIM-type	0.125	0.125	≥256	0.125	≤0.06	≤0.06	≥256	≤0.06	MBL
E. cloacae 4004	NDM-1	≤0.06	≤0.06	≥256	≤0.06	≤0.06	≤0.06	≥256	≤0.06	MBL
K. pneumoniae 4017	VIM-type	≤0.06	≤0.06	≥256	≤0.06	≤0.06	≤0.06	≥256	≤0.06	MBL
K. pneumoniae 4023	NDM-1	0.125	0.125	1	0.125	≤0.06	≤0.06	2	≤0.06	MBL
K. pneumoniae 4024	NDM-1	≤0.06	≤0.06	≥256	≤0.06	≤0.06	≤0.06	≥256	≤0.06	MBL
E. coli 4011	NDM-1	≤0.06	≤0.06	≥256	≤0.06	≤0.06	≤0.06	≥256	≤0.06	MBL
E. cloacae 4005	NDM	8	8	≥256	4	2	2	≥256	1	MBL

Table 1. Bacterial strains used in this investigation and results of phenotypic β -lactamase detection at 6 h using CTX -based^{ABCD} and CAZ-based^{abcd} assays.

CTX= cefotaxime; CAZ= ceftazidime; CA= clavulanic acid (4 μ g/ml); CX=cloxacillin (200 μ g/ml); EDTA = ethylenediaminetetraacetic acid (0.5mM); A (a) = CTX (CAZ)+ CX+EDTA; B (b) = CTX(CAZ)+CA+EDTA; C (c) = CTX(CAZ)+CA+CX; D (d) = CTX (CAZ)+CA+CX+EDTA. * control strain

MIC ratio ≥8	ESBL	AmpC	MBL
A/D or a/d	+	-	-
B/D or b/d	-	+	-
C/D or c/d	-	-	+
A/D or a/d and B/D or b/d	+	+	-
A/D or a/d and C/D or c/d	+	-	+
B/D or b/d and C/D or c/d	-	+	+
A/D or a/d and B/D or b/d and C/D or c/d	+	+	+

Table 2 Interpretation criteria of ceftazidime (CAZ)-based and cefotaxime (CTX)-based

 detection using resazurin microtitre plate assay.

A (a) = CTX (CAZ)+ CX+EDTA B (b) = CTX(CAZ)+CA+EDTA C (c) = CTX(CAZ)+CA+CX D (d) = CTX (CAZ)+CA+CX+EDTA