

## Erratum to: Prevalence and molecular epidemiology of acquired AmpC $\beta$ -lactamases and carbapenemases in *Enterobacteriaceae* isolates from 35 hospitals in Spain

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### Erratum to: Eur J Clin Microbiol Infect Dis DOI 10.1007/s10096-012-1737-0

Several errors

1.- Page 2, First column, First paragraph, line14-16:  
It should be eight new enzymes and *bla*<sub>CMY-96</sub> should be deleted.

Accordingly, it should be: "Eight new bla genes were described: *bla*<sub>CMY-54</sub>, *bla*<sub>CMY-55</sub>, *bla*<sub>CMY-56</sub>, *bla*<sub>CMY-57</sub>,

*bla*<sub>DHA-6</sub>, *bla*<sub>DHA-7</sub>, *bla*<sub>FOX-8</sub> and *bla*<sub>IMP-28</sub>" instead of "Nine new bla genes were described: *bla*<sub>CMY-54</sub>, *bla*<sub>CMY-55</sub>, *bla*<sub>CMY-56</sub>, *bla*<sub>CMY-57</sub>, *bla*<sub>CMY-96</sub>, *bla*<sub>DHA-6</sub>, *bla*<sub>DHA-7</sub>, *bla*<sub>FOX-8</sub> and *bla*<sub>IMP-28</sub>."

2.- Page 3, second column, third paragraph, line10-13:  
It should be seven new pAmpC and *bla*<sub>CMY-96</sub> should be deleted.

Accordingly, it should be: "Nucleotide sequence analysis of all bla genes identified seven new pAmpC (*bla*<sub>CMY-54</sub>,

The online version of the original article can be found at <http://dx.doi.org/10.1007/s10096-012-1737-0>.

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**Table 2** Prevalence of pAmpC and carbapenemases in the 674 Enterobacteriaceae strains studied

Micro-organism ( <i>n</i> )	pAmpC	No. of strains (%)	Carbapenemases	No. of strains (%)	pAmpC + carbapenemases	No. of strains (%)
<i>E. coli</i> (466) <sup>a</sup>	<b>CMY-2</b>	<b>406 (87.1)</b>	VIM-1	1 (0.2)		
	<b>DHA-1</b>	<b>37 (7.9)</b>				
	ACC-1	3 (0.6)				
	ACC-1+FOX	1 (0.2)				
	CMY-2t <sup>a</sup>	3 (0.6)				
	CMY-4	3 (0.6)				
	CMY-7	1 (0.2)				
	CMY-27	2 (0.4)				
	CMY-43	1 (0.2)				
	CMY-48	1 (0.2)				
	CMY-54	1 (0.2)				
	CMY-55	1 (0.2)				
	CMY-57	1 (0.2)				
	DHA-6	1 (0.2)				
	FOX-3	1 (0.2)				
	FOX-8	2 (0.4)				
<i>K. pneumoniae</i> (111)	<b>DHA-1</b>	<b>68 (61.3)</b>	VIM-1	9 (8.1)	DHA-1 + VIM-1	1 (0.9)
	CMY-2	18 (16)	IMP-22	9 (8.1)		
	ACC-1	3 (2.7)				
	CMY-56	1 (0.9)				
	CMY-4	1 (0.9)				
	CMY-2+DHA-1	1 (0.9)				
<i>P. mirabilis</i> (63)	<b>CMY-2</b>	<b>56 (88.8)</b>			CMY-2 + VIM-1	1 (1.6)
	ACC-1	3 (4.7)				
	DHA-1	3 (4.7)				
<i>K. oxytoca</i> (18)	DHA-1	5 (27.7)	<b>VIM-1</b>	<b>10 (55.5)</b>		
	CMY-2	2 (11.1)	IMP-28	1 (5.5)		
<i>E. cloacae</i> (11)			<b>VIM-1</b>	<b>9 (81.8)</b>	DHA-7 + VIM-1	2 (18.2)
<i>C. koseri</i> (3)	CMY-2	1				
	DHA-1	2				
<i>S. enterica</i> (1)	DHA-1	1				
<i>P. penneri</i> (1)	CMY-2	1				

<sup>a</sup>CMY-2t: CMY-2 related enzyme. Three isolates yielded positive PCR results for the internal fragment of CIT enzymes and sequences of the fragments were identical to CMY-2, despite several attempts to amplify the full gene consistently failing

*bla*<sub>CMY-55</sub>, *bla*<sub>CMY-56</sub>, *bla*<sub>CMY-57</sub>, *bla*<sub>DHA-6</sub>, *bla*<sub>DHA-7</sub> and *bla*<sub>FOX-8</sub>" instead of " Nucleotide sequence analysis of all *bla* genes identified eight new pAmpC (*bla*<sub>CMY-54</sub>, *bla*<sub>CMY-55</sub>, *bla*<sub>CMY-56</sub>, *bla*<sub>CMY-57</sub>, *bla*<sub>CMY-96</sub>, *bla*<sub>DHA-6</sub>, *bla*<sub>DHA-7</sub> and *bla*<sub>FOX-8</sub>)".

3.- Page 3, first column, second paragraph (Nucleotide sequence accession numbers), line 4:

The accession number HQ267531 (*bla*<sub>CMY-96</sub>), should be deleted.

4.- Regarding table 2, please see corrected file :