

TABLE 1.Taxonomy of bacteria expressing chromosomally determined AmpC β -lactamases

Phylum, class, and order	Genus and species	GenBank protein accession no.	Reference(s)
<i>Actinobacteria</i>	<i>Mycobacterium smegmatis</i>	YP_888266	92
<i>Proteobacteria</i>			
<i>Alphaproteobacteria</i>			
	<i>Ochrobactrum anthropi</i>	CAC04522	127, 226
	<i>Rhodobacter sphaeroides</i>	YP_355256	24
	<i>Chromobacterium violaceum</i>	NP_900980	87
<i>Betaproteobacteria</i>			
<i>Neisseriales</i>			
	<i>Laribacter hongkongensis</i>	AAT46346	167
<i>Gammaproteobacteria</i>			
<i>Aeromonadales</i>			
	<i>Aeromonas caviae</i>	AAM46773	95
	<i>Aeromonas hydrophila</i>	YP_857635	11, 334
	<i>Aeromonas jandae</i> ^a	AAA83416	272
	<i>Aeromonas salmonicida</i>	ABO89301	120
	<i>Aeromonas veronii</i> bv. <i>sobria</i>	CAA56561	333, 334
<i>Enterobacteriales</i>			
	<i>Buttiauxella agrestis</i>	AAN17791	90
	<i>Citrobacter braakii</i>	AAM11668	223
	<i>Citrobacter freundii</i>	AAM93471	178
	<i>Citrobacter murlinae</i>	AAM11664	12, 223
	<i>Citrobacter youngae</i>	CAD32304	12
	<i>Citrobacter werkmanii</i>	AAM11670	223
	<i>Edwardsiella tarda</i>	ABO48510	312
	<i>Enterobacter aerogenes</i>	AAO16528	266
	<i>Enterobacter asburiae</i>	CAC85157	279
	<i>Enterobacter cancerogenus</i>	AAM11666	223
	<i>Enterobacter cloacae</i>	P05364	101
	<i>Enterobacter dissolvens</i>	CAC85359	279
	<i>Enterobacter hormaechei</i>	CAC85357	279
	<i>Enterobacter intermedius</i> ^b	CAC85358	279
	<i>Erwinia rhapontici</i>	AAP40275	225
	<i>Escherichia albertii</i>	EDS93081	310
	<i>Escherichia fergusonii</i>	AAM11671	223
	<i>Escherichia coli</i>	NP_418574	144
	<i>Hafnia alvei</i>	AAF86691	107, 320
	<i>Morganella morganii</i>	AAC68582	260, 264
	<i>Providencia stuartii</i>	CAA76739	68

^aOriginally named *Aeromonas sobria*.^bAlternate name, *Kluyvera intermedia*.^c*Shigella* strains with enhanced virulence and a 190-kb chromosomal deletion that includes *bla*_{AmpC} have been described ([208](#)).