

**Table S1: Characterisation of studies detecting disseminated AMR genes in sewage**

Country/ study site	Year of sample collection	Targeted/detected ARB	ARG associated with resistance to	AMR Molecular Detection		Phenotypic/ genotypic analysis	Reference
				Technology	Platform		
Bangladesh/ Mymensing	2018	<i>E. coli</i> and <i>Salmonella</i> Spp.	Tetracycline ( <i>tetA</i> ).	PCR	-	Both	[35]
China/ West China	2017-2018	<i>Acinetobacter cumulans</i> (novel species)	Carbapenems ( <i>blaNDM-1</i> , <i>blaOXA-58</i> and <i>blaOXA-23</i> ); other β-lactams ( <i>blaPER-1</i> ); aminoglycosides ( <i>aac(6')-Ib</i> , <i>aac(3)-IId</i> and <i>aph6</i> ); bleomycin ( <i>ble</i> ); macrolides ( <i>mphE</i> and <i>msrE</i> ); rifampin ( <i>arr-3</i> ); sulfonamides ( <i>sul1</i> and <i>sul2</i> ) and tetracycline ( <i>tet39</i> and <i>tetY</i> ).	WGS	Illumina HiSeq	Both	[36]
USA/ Larimer County, Colorado	2016-2017	<i>E. coli</i>	Carbapenems (e.g., <i>blaOXA</i> , <i>blaCTX-M</i> , and <i>blaTEM</i> ); aminoglycoside (e.g., <i>aac(6')</i> , <i>aac(3)</i> and <i>aph(6)</i> ); fluoroquinolone ( <i>qnr</i> and <i>mfd</i> ); phenicol ( <i>cat</i> ) and sulfonamid ( <i>sul</i> ).	PCR and WGS	Illumina MiSeq	Both	[37]
Japan/	2017	<i>K. pneumoniae</i>	Carbapenems ( <i>blaKPC-2</i> ).	WGS	Illumina N	Both	[38]

Tokyo Bay					extSeq		
Ireland *	2017	<i>Enterobacteriaceae</i> ( <i>E. coli</i> , <i>Enterobacter cloacae</i> complex, <i>K. pneumoniae</i> , <i>Citrobacter freundii</i> and <i>Klebsiella oxytoca</i> )	Carbapenems ( <i>blaNDM</i> , <i>blaOXA-48</i> , <i>blaIMP</i> and <i>blaVIM</i> ).	PCR and WGS	Illumina HiSeq	Both	[39]
Poland/ Olsztyn	2016	<i>Bacteroides fragilis</i> group ( <i>Parabacteroides distasonis</i> , <i>B. fragilis</i> , <i>B. thetaiotaomicron</i> , <i>B. ovatus</i> and <i>B. cacae</i> )	Fluoroquinolone ( <i>bexA</i> , <i>qnrB</i> and <i>qnrS</i> ); macrolide; lincosamide and streptogramin ( <i>ermF</i> , <i>linA</i> and <i>mefA</i> ); tetracycline ( <i>tetQ</i> , <i>tetA</i> , <i>tetB</i> , <i>tetL</i> and <i>tetM</i> ); chloramphenicol ( <i>catA1</i> ) and β-lactam ( <i>cepA</i> and <i>cfxA</i> ).	PCR	-	Both	[40]
Global collection from 60 countries (74 cities)	2016	ND <sup>#</sup>	β-lactam (e.g., variants of <i>blaOXA</i> ); aminoglycoside ( <i>aadA</i> ); tetracycline (e.g., <i>tetQ</i> and <i>tetW</i> ); sulfonamide (variants of <i>sul1</i> ); macrolide (e.g., <i>mphE</i> , <i>msrD</i> , <i>msrE</i> , <i>ermB</i> , <i>ermF</i> and <i>mefA</i> ) and streptomycin (e.g., <i>strA</i> and <i>strB</i> ).	WGS	Illumina HiSeq	Genotype	[41]

Sweden/ Örebro	2016	ND <sup>#</sup>	$\beta$ -lactam (e.g., <i>blaCTX-M-1</i> , <i>blaSHV-156G</i> , <i>blaSHV-238G240E</i> , <i>blaACT-1</i> , <i>blaACT-5/7</i> , <i>blaCMY-10</i> , <i>blaDHA</i> , <i>blaFOX</i> , <i>blaLAT</i> , <i>blaMIR</i> and <i>blaMOX</i> ); fluoroquinolone (variants of <i>qnrB-1</i> and <i>aacC2</i> ); macrolide ( <i>mefA</i> ); aminoglycoside ( <i>aadA1</i> ); tetracycline ( <i>tetA</i> and <i>tetB</i> ) and macrolide-lincosamide-streptogramin B ( <i>ermB</i> ).	qPCR	-	Genotype	[42]
Lebanon/ Al-Qaa refugee camp	2016	<i>E. coli</i>	$\beta$ -lactam ( <i>blaCTX-M-14</i> , <i>blaOXA-1</i> , <i>blaSHV-12</i> and <i>blaCMY-2</i> ); aminoglycoside ( <i>aac(6)-Ib</i> , <i>acc(3)-II</i> and <i>acc(3)-II</i> ); fluoroquinolone (variants of <i>gyrA</i> and <i>parC</i> ) and <i>int-I1</i> .	PCR and DNA sequencing	Sanger sequencing	Both	[43]
Multiple countries**	2015-2016	ND <sup>#</sup>	Aminoglycoside ( <i>aadA</i> and <i>strB</i> ); $\beta$ -lactam (e.g., <i>blaOXA</i> , <i>blaGES</i> and <i>blaVEB</i> ); macrolide-lincosamide-streptogramin B ( <i>ereA</i> , <i>ermF</i> , and <i>matA/mel</i> ); sulfonamides ( <i>sul1</i> ); tetracyclines ( <i>tetM</i> and <i>tetQ</i> ); amphenicol ( <i>cmxA</i> ) and quinolone ( <i>qnrSrtF11</i> ).	PCR	-	Genotype	[44]
Burkina Faso/ Ouagadougou	2015	ND <sup>#</sup>	Sulfonamide (including <i>sul1</i> and <i>sul2</i> ); aminoglycoside (e.g., <i>aadA13</i> , <i>aadA</i> and <i>aadA15</i> ); tetracycline ( <i>tetB</i> ); $\beta$ -lactam (e.g., <i>blaOXA-226</i> , <i>blaOXA-256</i> , <i>blaOXA-347</i> , <i>blaOXA-46</i> , <i>blaSHV-100</i> , <i>blaGES-21</i> and <i>blaAIM-1</i> ) and macrolide-lincosamide-streptogramin B (e.g., <i>ermB</i> and <i>ermF</i> ).	WGS	Illumina HiSeq	Genotype	[45]

United Kingdom/ East of England	2014-2015	<i>Enterococcus faecium</i>	Vancomycin ( <i>vanA</i> ); aminoglycoside ( <i>aac(6')</i> - <i>aph(2")</i> and <i>aacA</i> ); trimethoprim ( <i>dfrG</i> ); macrolide-lincosamide-streptogramin B ( <i>msrC</i> , <i>ermB</i> and <i>ermF</i> ); spectinomycin ( <i>spw</i> ); chloramphenicol ( <i>cat</i> ) and tetracycline ( <i>tetM</i> and <i>tetS</i> ).	DNA sequencing	Illumina HiSeq	Both	[46]
South Africa/ Alice	2014	<i>Enterococcus spp.</i>	Vancomycin ( <i>vanB</i> , <i>vanC1</i> and <i>vanC2/3</i> ) and macrolide ( <i>ermB</i> ).	PCR	-	Both	[47]
China/ Beijing	2013	ND <sup>#</sup>	Sulfonamide (e.g., <i>sul1</i> , <i>sul2</i> and <i>sul 3</i> ); aminoglycoside (e.g., <i>aac3</i> and <i>aac6</i> ); tetracycline (e.g., <i>tet41</i> and <i>tetC</i> ); β-lactam (e.g., <i>bla<sub>penA</sub></i> and <i>bla<sub>pepEC</sub></i> ); macrolide- lincosamide-streptogramin B (e.g., <i>msrE</i> and <i>ermF</i> ); fluoroquinolone ( <i>oqxBgb</i> ); vancomycin ( <i>vanR-F</i> and <i>van R-M</i> ) and trimethoprim ( <i>dfrK</i> ).	WGS	Illumina HiSeq	Genotype	[48]
China/ Northern China	2013	ND <sup>#</sup>	Tetracycline ( <i>tetO</i> , <i>tetT</i> , <i>tetQ</i> , <i>tetW</i> and <i>tetM</i> ); β-lactam ( <i>bla<sub>OXA-1</sub></i> ); sulphonamide ( <i>sul1</i> and <i>sul2</i> ) and macrolide ( <i>ermB</i> ).	PCR	-	Genotype	[49]

Brazil/ Curitiba	2012-2013	<i>E. coli</i> , <i>K. pneumoniae</i> and <i>K. oxytoca</i>	$\beta$ -lactam (e.g., <i>bla</i> <sub>CTX-M</sub> , <i>bla</i> <sub>SHV</sub> and <i>bla</i> <sub>GES-5</sub> ) and fluoroquinolone ( <i>oqxAB</i> , <i>aac-6'-Ib-cr</i> , <i>qnrS</i> , <i>qnrB</i> and variants of <i>gyrA</i> ).	PCR and DNA sequencing	Sanger sequencing	Both	[50]
Germany/ South-west Germany	2012-2013	ND <sup>#</sup>	Aminoglycoside ( <i>ant(3")</i> ); $\beta$ -lactam ( <i>bla</i> <sub>OXA</sub> ); macrolide ( <i>ermB</i> and <i>macB</i> ); multidrug ( <i>acrB</i> ); quinolone ( <i>parC</i> and <i>gyrA</i> ); rifamycin ( <i>rpoB</i> ) and tetracycline ( <i>tetW</i> , <i>tetO</i> and <i>tet32</i> ).	WGS	Illumina HiSeq	Genotype	[51]
South Africa/ Amathole District Municipality	2012-2013	<i>E. coli</i>	Aminoglycosides ( <i>strA</i> and <i>aadA</i> ); phenicols ( <i>catI</i> , <i>catII</i> and <i>cmlA1</i> ); $\beta$ -lactam ( <i>bla</i> <sub>TEM</sub> and <i>ampC</i> ) and tetracyclin ( <i>tetA</i> , <i>tetB</i> , <i>tetD</i> , <i>tetK</i> and <i>tetM</i> ).	PCR	-	Both	[52]
South Africa The kwini	2012	Not specify**	Sulfonamide ( <i>sul1</i> , <i>sul2</i> and <i>sul3</i> ) and tetracycline ( <i>tetM</i> ).	PCR and DNA sequencing	Sanger sequencing	Both	[53]
Brazil/ Rio de Janeiro	2011	ND <sup>#</sup>	$\beta$ -lactam ( <i>bla</i> <sub>CfxA</sub> , <i>bla</i> <sub>ACC</sub> , <i>bla</i> <sub>OXA-10</sub> , <i>bla</i> <sub>CEPA</sub> and <i>bla</i> <sub>Fox</sub> ).	WGS	Roche 454 sequencer	Genotype	[54]
Finland/ Helsinki Estonia/ Tartu and Tallinn	2010-2011	ND <sup>#</sup>	Tetracycline ( <i>tetC</i> and <i>tetM</i> ); $\beta$ -lactam ( <i>bla</i> <sub>oxa-58</sub> , <i>bla</i> <sub>shv-34</sub> and <i>bla</i> <sub>ctx-m-32</sub> ) and sulphonamide ( <i>sul1</i> and <i>sul2</i> ).	PCR	-	Genotype	[55]

India/ New Delhi	2010-2011	<i>V. cholerae</i> , <i>E. coli</i> , <i>Aeromonas caviae</i> , <i>Shigella boydii</i> , <i>Enterobacteria</i> and <i>Aeromonas</i>	β-lactam ( <i>bla</i> <sub>NDM-1</sub> ).	PCR	-	Both	[56]
China/Hong Kong, Shatin	2010	ND <sup>#</sup>	β-lactam ( <i>ampC</i> , <i>bla</i> <sub>VEB-3</sub> , <i>bla</i> <sub>VIM-2</sub> and <i>bla</i> <sub>IMP-1</sub> ); aminoglycoside ( <i>aacA4</i> , <i>aadA1</i> , <i>aadA2</i> , <i>aadA2b</i> and <i>aadA24</i> ); sulphonamides ( <i>sull</i> ); trimethoprim ( <i>dfrA1</i> ); tetracycline ( <i>tetA</i> , <i>tetC</i> , <i>tetG</i> , <i>tetM</i> and <i>tetB</i> ); macrolide ( <i>ermB</i> and <i>macB</i> ) and aminoglycoside ( <i>aadB</i> ).	PCR and WGS	Illumina HiSeq	Genotype	[57]
Brazil/ Rio de Janeiro	2008	<i>K. pneumoniae</i> , <i>Enterobacter cloacae</i> and <i>E. coli</i>	β-lactam ( <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>SHV</sub> and <i>bla</i> <sub>CTX-M</sub> ).	PCR and DNA sequencing	Sanger sequencing	Both	[58]
Germany/ Bielefeld-Heepen	2006	Not specified**	Aminoglycoside (e.g., <i>aacA</i> , <i>aac(6')</i> and <i>aph</i> ); β-lactam (e.g., <i>bla</i> <sub>TLA-2</sub> , <i>bla</i> <sub>CTXM-27</sub> , <i>bla</i> <sub>GES-3</sub> , <i>bla</i> <sub>IMP-13</sub> and <i>bla</i> <sub>OXA-58</sub> ); chloramphenicol (e.g., <i>cmxA</i> and <i>cat</i> ); fluoroquinolone (e.g., <i>qnr</i> and <i>qnrB4</i> ); erythromycin ( <i>ereA2</i> and <i>arr2</i> ); tetracycline ( <i>tetB(P)</i> , <i>tetL</i> , <i>tetM</i> , <i>tetS</i> and <i>tetX</i> ); sulfonamide ( <i>sul1</i> and <i>sul3</i> ); trimethoprim ( <i>dfrD</i> ) and multidrug ( <i>mexD</i> , <i>mexI</i> and <i>mexY</i> ).	PCR and DNA sequencing	Sanger sequencing	Both	[59]
China/ North China	2004-2005	Multiple bacteria***	Tetracycline ( <i>tetA</i> , <i>tetW</i> , <i>tetC</i> , <i>tetJ</i> , <i>tetL</i> , <i>tetD</i> , <i>tetY</i> and <i>tetK</i> ).	PCR and DNA sequencing	Sanger sequencing	Both	[60]

China/ Southern, Northern and Eastern regions	‡	ND <sup>#</sup>	Sulphonamide ( <i>sul1</i> and <i>sul2</i> ); tetracycline ( <i>tetX</i> , <i>tetW</i> and <i>tetQ</i> ) and class 1 integron ( <i>intI1</i> ).	PCR and WGS	Illumina MiSeq	Genotype	[61]
China/ Hong Kong	‡	ND <sup>#</sup>	aminoglycoside (e.g., <i>aadA</i> ); bacitracin ( <i>bacA</i> ); macrolide-lincosamide-streptogramin B (e.g., <i>macB</i> , <i>mefA</i> and <i>ermB</i> ); tetracycline ( <i>tet32</i> , <i>tetM</i> , <i>tetQ</i> and <i>tetO</i> ); sulfonamide ( <i>sul1,2</i> ); vancomycin ( <i>vanR</i> ); trimethoprim ( <i>dfrA</i> ); quinolone ( <i>qnrS</i> ); β-lactam (e.g., <i>blaOXA-1</i> , <i>blaTEM</i> , <i>blaCMY-2</i> and <i>blaCTX-M</i> ) and multiple drugs (e.g., <i>acrA</i> , <i>mdtH</i> , <i>mdtL</i> and <i>mdtO</i> ).	WGS	Illumina HiSeq	Genotype	[62]
Germany*	‡	<i>E. coli</i> , <i>K. pneumoniae</i> and <i>A. baumannii</i>	Colistin ( <i>mcr-1</i> ); erythromycin ( <i>ermB</i> ); tetracycline ( <i>tetM</i> ); β-lactam (e.g., <i>blaOXA1</i> , <i>blaTEM</i> , <i>blaCMY-2</i> and <i>blaCTX-M</i> ) and chloramphenicol ( <i>catA2</i> and <i>cmlA5</i> ).	PCR	-	Genotype	[63]

South Africa/ Fort Beaufort and Alice	‡	<i>Aeromonas</i> spp.	β-lactam ( <i>bla<sub>TEM</sub></i> ) and class 1 integron.	PCR	-	Both	[64]
South Africa/ Fort Beaufort and Alice	‡	<i>Vibrio</i> spp.	Dulfonamide ( <i>sul2</i> ), chloramphenicol ( <i>floR</i> ), trimetophrim ( <i>dfr18</i> and <i>dfrA1</i> ), tetracycline ( <i>tetA</i> ), streptomycin ( <i>strB</i> ) and β-lactam ( <i>bla<sub>TEM</sub></i> ).	PCR	-	Both	[65]
India/ Hyderabad Sweden / Skövde	‡	<i>E. coli</i>	Quinolone (variants of <i>GyrA</i> and <i>ParC</i> ).	PCR and WGS	the GS FLX+ system	Genotype	[66]
Swedish/ Uppsala	2003	<i>Enterococcus hirae</i> , <i>E. faecium</i> and <i>E. durans</i>	Vancomycin ( <i>vanA</i> and <i>vanB</i> ).	PCR	-	Both	[67]
Portugal/ Continental territory	ND	<i>E. faecium</i> , <i>E. gallinarum</i> and <i>E. casseliflavus</i>	Vancomycin ( <i>vanA</i> and <i>vanC1/2</i> ).	PCR	-	Both	[68]
USA****	2003-2006	<i>Clostridium perfringens</i>	Tetracycline ( <i>tetA</i> , <i>tetB</i> and <i>tetM</i> ) and macrolide ( <i>ermB</i> , <i>ermQ</i> and <i>mefA</i> ).	PCR and DNA sequencing	Sanger sequencing	Genotype	[69]

China/ Hong Kong and Shanghai	2007-2008	<i>Enterobacteriaceae</i>	Trimetophrim ( <i>dfr17</i> ) and spectinomycin/streptomycin ( <i>aadA5</i> ).	PCR and DNA sequencing	Sanger sequencing	Both	[70]
USA/Sao Paolo							
South Africa/ Eastern Cape	2007-2008	<i>Listeria ivanovii</i> and <i>L. innocua</i>	Sulphonamide ( <i>sulII</i> ).	PCR	-	Both	[71]
South Africa/ Eastern Cape	2010	<i>Vibrio</i> spp.	Trimethoprim ( <i>dfr18</i> and <i>dfrA1</i> ), chloramphenicol ( <i>floR</i> ), tetracycline ( <i>tetA</i> ), streptomycin ( <i>strB</i> ) and sulfamethoxazole ( <i>sulII</i> ).	PCR	-	Both	[72]
USA/ Northern Colorado	2008-2009	ND <sup>#</sup>	Tetracycline ( <i>tetB</i> , <i>tetC</i> , <i>tetO</i> and <i>tetW</i> ).	PCR	-	Both	[73]
France/ Seine River							
Belgium/ Scheldt River	2008-2009	Heterotrophic bacteria	ND	PCR and DNA sequencing	Sanger sequencing	Both	[74]
Spain/ Barcelona	‡	MRSA and <i>E. coli</i>	β-lactam ( <i>bla<sub>TEM</sub></i> and <i>bla<sub>CTX-M</sub></i> ) and penicillin- binding protein ( <i>mecA</i> ).	PCR and DNA sequencing	Sanger sequencing	Genotype	[75]

Brazil/ Rio Grande	2005 and 2007	<i>Pseudomonas aeruginosa</i>	ND	PCR	-	Both	[76]
United Kingdom	‡	ND <sup>#</sup>	Class 1 integron.	PCR	-	Genotype	[77]
China/ Jiangshou	2010	<i>E. coli</i> , <i>Klebsiella</i> spp. and <i>Aeromonas</i> spp.	Class 1 integronases ( <i>intI1</i> and <i>qacED1</i> ) and sulfonamid ( <i>sull</i> ).	PCR	-	Genotype	[78]
Poland	2008-2010	<i>E. coli</i>	Class 1 and 2 integron.	PCR	-	Genotype	[79]
Nicaragua/ Leon	20008-2009	<i>E. coli</i>	β-lactam ( <i>bla</i> <sub>CTX-M</sub> , <i>bla</i> <sub>SHV</sub> , <i>bla</i> <sub>OXA</sub> and <i>bla</i> <sub>TEM</sub> ).	PCR and DNA sequencing	Sanger sequencing	Both	[80]
Poland	2008-2010	<i>Enterobacteriaceae</i>	Class 1 and 2 integron.	PCR and DNA sequencing	Sanger sequencing	Both	[81]
Brazil	2007	<i>Pseudomonas</i> spp.	Class 1 integron.	PCR	-	Both	[82]
Australia/ Queensland	2011	<i>E. coli</i>	β-lactam, including <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>SHV</sub> , <i>bla</i> <sub>CTX-M</sub> and <i>bla</i> <sub>OXA</sub> .	PCR and DNA sequencing	Sanger sequencing	Both	[83]

Austria/ Graz and Styria	2009	<i>E. coli</i>	$\beta$ -lactam ( <i>bla</i> <sub>CTX-M-15</sub> , <i>bla</i> <sub>TEM</sub> and <i>bla</i> <sub>SHV</sub> ).	PCR and DNA sequencing	Sanger sequencing	Both	[84]
Spain/ Catalonia	2009-2010	<i>E. coli</i>	$\beta$ -lactam ( <i>bla</i> <sub>CTX-M-15</sub> , <i>bla</i> <sub>TEM</sub> and <i>bla</i> <sub>SHV</sub> ).	PCR	-	Both	[85]
Poland/ Gulf of Gdansk	2011	<i>Enterococcus</i> spp.	Tetracyclin ( <i>tetM</i> , <i>tetL</i> and <i>tetS</i> ); fluoroquinolon (variants of <i>gyrA</i> and <i>parC</i> ); streptomycin ( <i>ant(6')</i> - <i>Ia</i> ) and glycopeptide ( <i>aac(6')</i> - <i>Ie-aph(2")</i> ).	PCR and DNA sequencing	Sanger sequencing	Both	[86]
China/ Shanghai	2013-2014	ND <sup>#</sup>	Erythromycin ( <i>ereA</i> , <i>ereB</i> , <i>ermA</i> , <i>ermB</i> , <i>ermC</i> , <i>mefA/mefE</i> and <i>msrA/msrB</i> ).	PCR	-	Both	[87]
Japan/ Miyazaki	‡	<i>Enterococcus</i> spp.	Vancomycin ( <i>vanC1</i> and <i>vanC2/C3</i> ).	PCR and DNA sequencing	-	Both	[88]
China/Jiangsu	‡	Heterotrophic bacteria	$\beta$ -lactam ( <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>SHV</sub> and <i>bla</i> <sub>CTX_M</sub> ); sulfonamide ( <i>sul1</i> , <i>sul2</i> and <i>sul3</i> ); tetracycline ( <i>tetA</i> , <i>tetB</i> , <i>tetE</i> and <i>tetO</i> ) and streptomycin ( <i>strA</i> and <i>strB</i> ).	PCR	-	Both	[89]
France/ Limoges Luxembourg/ Esch-sur Alzette	2010-2011	<i>Acinetobacter johnsoni</i> , <i>Aeromonas allosacharophiles</i> and <i>Citrobacter</i> spp.	Class 3 integron with oxillinase gene cassette.	PCR	-	Both	[90]

United States/Florida	2014	<i>Enterococcus</i> spp.	Vancomycin ( <i>vanA</i> ).	PCR and WGS	Illumina	Both	[91]
South Korea/	‡	ND <sup>#</sup>	Tetracycline ( <i>tetM</i> , <i>tetQ</i> , <i>tetA</i> , <i>tetB</i> , <i>tetE</i> , <i>tetG</i> , <i>tetH</i> and <i>tetX</i> ); sulfonamides ( <i>sul1</i> and <i>sul2</i> ); macrolides ( <i>ermB</i> and <i>ermC</i> ); quinolones ( <i>qnrD</i> ) and β-lactam ( <i>bla<sub>TEM</sub></i> ).	PCR	-	Genotype	[92]
China/Xiangmen and LongYan	2014	ND <sup>#</sup>	Aminoglycosides, beta-lactams, bicyclomycin, chloramphenicol, fosfomycin, gentamicin and macrolides.	PCR and WGS	Illumina Miseq300	Genotype	[93]
Singapore	2014	<i>Pseudomonas</i> spp., <i>Klebsiella</i> spp., <i>Enterobacter</i> spp. and <i>Citrobacter</i> spp.	β-lactam ( <i>blaSHV</i> , <i>bla<sub>NDM-1</sub></i> , <i>bla<sub>CTX</sub></i> and <i>bla<sub>KPC</sub></i> ).	PCR	-	Both	[94]
Austria/Graz	2011-2012	Enterobacteriaceae, MRSA and vancomycin-resistant <i>Enterococci</i>	β-lactam ( <i>bla<sub>CTX-M</sub></i> ).	PCR and DNA sequencing	Sanger sequencing	Both	[95]
China	‡	ND <sup>#</sup>	<i>blaSHV/TEM</i> and <i>sul1</i> .	PCR	-	Genotype	[96]

Algeria/ Bejaia and Tizi Ouzou	2011- 2012	<i>E. coli</i> , <i>Klebsiella pneumoniae</i> , <i>Acinetobacter</i> spp., <i>Aeromonas</i> spp. and <i>Pseudomonas</i> spp.	$\beta$ -lactam ( <i>bla</i> <sub>CTX-M</sub> , <i>bla</i> <sub>TEM</sub> , <i>bla</i> <sub>SHV</sub> , <i>bla</i> <sub>OXA-48-like</sub> , <i>bla</i> <sub>OXA-23</sub> and <i>bla</i> <sub>OXA-51</sub> ); quinolone ( <i>qnrB</i> and <i>qnrS</i> ); tetracycline ( <i>tetA</i> , <i>tetB</i> and <i>tetC</i> ); trimethoprim ( <i>dfrA1</i> ); aminoglycoside ( <i>aac(3)-IIc</i> ( <i>aacC2</i> ) and <i>aac(6')-1b</i> ) and sulfonamid ( <i>sul1</i> and <i>sul2</i> ).	PCR	-	Both	[97]
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\* - study site is not mentioned.

# - identification of bacteria species was not performed.

\*\* - Portugal, Spain, Ireland, Cyprus, Germany, Finland and Norway.

\*\*\*\* - Arizona, California, Florida, Idaho, Indiana, Iowa, Georgia, Maryland, Massachusetts, Montana, North Carolina, Pennsylvania, Texas and Washington States.

‡ - year of collection was not mentioned.