

Clonal Distribution and Its Association With the Carbapenem Resistance Mechanisms of Carbapenem-Non-Susceptible *Pseudomonas aeruginosa* Isolates From Korean Hospitals

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SUPPLEMENTAL DATA

Supplemental Data Table S1. Characteristics of 155 carbapenem-non-susceptible *P. aeruginosa* isolates collected in three Korean hospitals between 2011 and 2019 (provided in Excel format)

Supplemental Data Table S2. Primers used in this study

Primer	Sequence (5' to 3')		Use	Reference
GES-F	ATGCGCTTCATTCACGCAC			
GES-R	CTATTTGTCCGTGCTCAGG	<i>bla_{GES}</i>	Detection	[24]
KPC-F	ATCGCCGTCTAGTTCTGCTG			
KPC-R	CGTTGACGCCCAATCC	<i>bla_{KPC}</i>	Detection	[24]
blaIMP-F	GAATAGRRTGGCTTAAYTCTC			
blaIMP-R	CCAAACYACTASGTTATC	<i>bla_{IMP}</i>	Detection	[25]
blaNDM-F	TTGGCCTTGCTGTCCTTG			
blaNDM-R	ACACCAGTGACAATATCACCG	<i>bla_{NDM}</i>	Detection	[26]
blaSPM-F	CTGCTTGGATTCATGGGCGC			
blaSPM-R	CCTTTTCCGCGACCTTGATC	<i>bla_{SPM}</i>	Detection	[15]
blaVIM-F	GTTTGGTCGCATATCGCAAC			
blaVIM-R	AATGCGCAGCACCAGGATAG	<i>bla_{VIM}</i>	Detection	[15]
oxa-48-F	TTGGTGGCATCGATTATCGG			
oxa-48-R	GAGCACTTCTTTTGTGATGGC	<i>bla_{oxa-48}</i>	Detection	[15]
oxa-58-F	CGATCAGAATGTTCAAGCGC			
oxa58-R	ACGATTCTCCCCTCTGCGC	<i>bla_{oxa-58}</i>	Detection	[15]
armA-F	ATTCTGCCTATCCTAATTGG			
armA-R	ACCTATACTTTATCGTCGTC	<i>armA</i>	Detection	[27]
rmtB-F	GCTTTCTGCGGGCGATGTAA			
rmtB-R	ATGCAATGCCGCGCTCGTAT	<i>rmtB</i>	Detection	[27]
oprD-F	CCAAGAAACACTGCGTGCTA			
oprD-R	GGGGTGTTTTTCGCAGAGTAA	<i>oprD</i>	Amplification	[24]
oprD-seq-F	CCTGCGTAGGTGGCATAGAG			
oprD-seq-R	TCCAAGACCATGCTGAAGTG	<i>oprD</i>	Sequencing	[24]
blaIMP-Full-F	<u>GTTTTCCAGTCACGACGTTATGAGC</u>	<i>bla_{IMP}</i>	Amplification	[28]

	AAGTTATCTGTATTCTT		and	
blaIMP-Full-R	<u>CAGGAAACAGCTATGACCATGATCGT</u> TTAACCCCTTTAACCGCCTG		sequencing	
blaNDM-Full-F	<u>GTTTTCCCAGTCACGACGTTATGGAA</u> TTGCCCAATATTATGCACCCGGTCG		Amplification	
blaNDM-Full-R	<u>CAGGAAACAGCTATGACCATGATCAG</u> CGCAGCTTGTCGGCCATG	<i>bla_{NDM-1}</i>	and sequencing	[28]
blaVIM-Full-F	<u>GTTTTCCCAGTCACGACGTTATGTTC</u> AAACTTTTGAGTAAG		Amplification	
blaVIM-Full-R	<u>CAGGAAACAGCTATGACCATGACTAC</u> TCAACGACTGAGCGAT	<i>bla_{VIM}</i>	and sequencing	[28, 29]
ampC-F	GGGCTGGCCTCGAAAGAGGAC		Gene	
ampC-R	GCACCGAGTCGGGGAAGTCA	<i>ampC</i>	expression	[30]
rpsL-F	GCTGCAAACTGCCCGCAACG		Gene	
rpsL-R	ACCCGAGGTGTCCAGCGAACC	<i>rpsL</i>	expression	[15]

Underlined sequences indicate regions of the universal primer that are not complementary to the template.