

Antibiogram of Carbapenemase Producing Organisms - February, 2022

Table 1: Aggregate antibiogram of carbapenemase producing Enterobacterales, Acinetobacter and Pseudomonas isolates by number and percentage susceptible to each antimicrobial, received by MDU PHL 01/01/2020 - 31/12/2021

Beta-lactamase class	CPO Gene	Organism	N	Count of susceptible isolates and susceptibility proportion (%)													
				Aminoglycosides (see note 1)			Monobactams	Cephalosporins					Fluoroquinolones	Carbapenems	Tetracyclines	Miscellaneous agents	
				Amikacin	Gentamicin	Tobramycin	Aztreonam	Ceftazidime/Avibactam	Ceftazidime	Ceftolozane/Tazobactam	Ceftriaxone	Cefepime	Ciprofloxacin	Meropenem	Tigecycline	Colistin	Fosfomycin
Class A	KPC-2	<i>Klebsiella pneumoniae</i>	6	-	3/4 (75.0%)	1/4 (25.0%)	0/4 (0.0%)	5/6 (83.3%)	0/4 (0.0%)	0/6 (0.0%)	0/4 (0.0%)	0/4 (0.0%)	-	0/6 (0.0%)	-	4/6 (66.7%)	6/6 (100.0%)
Class B	NDM-1	<i>Citrobacter freundii</i> complex	10	10/10 (100.0%)	8/10 (80.0%)	8/10 (80.0%)	6/10 (60.0%)	*	0/10 (0.0%)	*	0/8 (0.0%)	0/8 (0.0%)	8/10 (80.0%)	3/10 (30.0%)	7/7 (100.0%)	9/10 (90.0%)	10/10 (100.0%)
		<i>Enterobacter cloacae</i>	8	5/8 (62.5%)	0/8 (0.0%)	0/8 (0.0%)	1/8 (12.5%)	*	0/8 (0.0%)	*	0/5 (0.0%)	0/5 (0.0%)	1/7 (14.3%)	4/8 (50.0%)	8/8 (100.0%)	8/8 (100.0%)	7/8 (87.5%)
		<i>Escherichia coli</i>	21	19/20 (95.0%)	18/20 (90.0%)	16/20 (80.0%)	16/20 (80.0%)	*	0/20 (0.0%)	*	0/16 (0.0%)	0/16 (0.0%)	7/18 (38.9%)	3/20 (15.0%)	19/19 (100.0%)	20/20 (100.0%)	20/21 (95.2%)
		<i>Klebsiella oxytoca</i>	9	8/8 (100.0%)	6/8 (75.0%)	6/8 (75.0%)	8/8 (100.0%)	*	0/8 (0.0%)	*	0/5 (0.0%)	0/5 (0.0%)	4/7 (57.1%)	2/9 (22.2%)	5/5 (100.0%)	9/9 (100.0%)	9/9 (100.0%)
		<i>Klebsiella pneumoniae</i>	10	7/10 (70.0%)	8/10 (80.0%)	6/10 (60.0%)	6/10 (60.0%)	*	0/10 (0.0%)	*	0/10 (0.0%)	0/10 (0.0%)	0/5 (0.0%)	6/10 (60.0%)	8/8 (100.0%)	10/10 (100.0%)	9/10 (90.0%)
		<i>Pseudomonas aeruginosa</i>	5	0/5 (0.0%)	0/5 (0.0%)	0/5 (0.0%)	3/5 (60.0%)	*	0/5 (0.0%)	*	*	0/5 (0.0%)	1/5 (20.0%)	0/5 (0.0%)	*	4/5 (80.0%)	^
	IMP-4	<i>Citrobacter freundii</i> complex	11	10/11 (90.9%)	0/11 (0.0%)	0/11 (0.0%)	4/11 (36.4%)	*	0/11 (0.0%)	*	0/9 (0.0%)	4/9 (44.4%)	6/11 (54.5%)	8/11 (72.7%)	11/11 (100.0%)	11/11 (100.0%)	11/11 (100.0%)
		<i>Enterobacter cloacae</i>	62	61/62 (98.4%)	3/62 (4.8%)	1/62 (1.6%)	15/62 (24.2%)	*	1/62 (1.6%)	*	1/58 (1.7%)	17/58 (29.3%)	18/50 (36.0%)	58/62 (93.5%)	42/43 (97.7%)	59/61 (96.7%)	55/62 (88.7%)
		<i>Escherichia coli</i>	7	7/7 (100.0%)	1/7 (14.3%)	0/7 (0.0%)	7/7 (100.0%)	*	1/7 (14.3%)	*	1/7 (14.3%)	6/7 (85.7%)	4/7 (57.1%)	6/7 (85.7%)	7/7 (100.0%)	7/7 (100.0%)	7/7 (100.0%)
		<i>Klebsiella oxytoca</i>	8	8/8 (100.0%)	2/8 (25.0%)	0/8 (0.0%)	6/8 (75.0%)	*	0/8 (0.0%)	*	0/6 (0.0%)	4/6 (66.7%)	6/8 (75.0%)	8/8 (100.0%)	7/7 (100.0%)	8/8 (100.0%)	8/8 (100.0%)
		<i>Klebsiella pneumoniae</i>	12	12/12 (100.0%)	1/12 (8.3%)	1/12 (8.3%)	11/12 (91.7%)	*	1/12 (8.3%)	*	1/11 (9.1%)	7/11 (63.6%)	3/12 (25.0%)	10/12 (83.3%)	10/10 (100.0%)	12/12 (100.0%)	12/12 (100.0%)
		<i>Serratia marcescens</i>	11	10/10 (100.0%)	4/9 (44.4%)	0/10 (0.0%)	10/10 (100.0%)	*	0/10 (0.0%)	*	0/8 (0.0%)	0/8 (0.0%)	1/8 (12.5%)	6/11 (54.5%)	2/8 (25.0%)	*	7/11 (63.6%)
	NDM-5	<i>Escherichia coli</i>	36	30/34 (88.2%)	22/35 (62.9%)	16/35 (45.7%)	6/35 (17.1%)	*	0/35 (0.0%)	*	0/29 (0.0%)	0/29 (0.0%)	0/27 (0.0%)	8/36 (22.2%)	35/35 (100.0%)	36/36 (100.0%)	35/36 (97.2%)
	NDM-7	<i>Klebsiella pneumoniae</i>	6	6/6 (100.0%)	6/6 (100.0%)	6/6 (100.0%)	6/6 (100.0%)	*	0/6 (0.0%)	*	0/6 (0.0%)	0/6 (0.0%)	6/6 (100.0%)	1/6 (16.7%)	-	6/6 (100.0%)	5/5 (100.0%)
	VIM-2	<i>Pseudomonas aeruginosa</i>	6	0/5 (0.0%)	0/5 (0.0%)	0/5 (0.0%)	5/5 (100.0%)	*	0/5 (0.0%)	*	*	0/4 (0.0%)	0/5 (0.0%)	1/6 (16.7%)	*	2/6 (33.3%)	^
Class D	OXA-181	<i>Escherichia coli</i>	8	7/8 (87.5%)	7/8 (87.5%)	6/8 (75.0%)	2/8 (25.0%)	8/8 (100.0%)	2/8 (25.0%)	1/8 (12.5%)	2/6 (33.3%)	3/6 (50.0%)	1/6 (16.7%)	8/8 (100.0%)	8/8 (100.0%)	8/8 (100.0%)	8/8 (100.0%)
		<i>Klebsiella pneumoniae</i>	5	-	4/4 (100.0%)	3/4 (75.0%)	3/4 (75.0%)	5/5 (100.0%)	1/4 (25.0%)	2/5 (40.0%)	-	-	-	5/5 (100.0%)	4/4 (100.0%)	5/5 (100.0%)	5/5 (100.0%)
	OXA-23	<i>Acinetobacter baumannii</i>	10	0/9 (0.0%)	0/9 (0.0%)	0/9 (0.0%)	^	*	^	^	^	^	0/9 (0.0%)	0/10 (0.0%)	^	9/10 (90.0%)	*
		<i>Proteus mirabilis</i>	5	4/5 (80.0%)	4/5 (80.0%)	4/5 (80.0%)	5/5 (100.0%)	5/5 (100.0%)	5/5 (100.0%)	5/5 (100.0%)	5/5 (100.0%)	-	5/5 (100.0%)	5/5 (100.0%)	*	*	5/5 (100.0%)
	OXA-48	<i>Escherichia coli</i>	7	7/7 (100.0%)	5/7 (71.4%)	5/6 (83.3%)	1/7 (14.3%)	6/7 (85.7%)	3/7 (42.9%)	5/7 (71.4%)	1/5 (20.0%)	3/5 (60.0%)	3/6 (50.0%)	7/7 (100.0%)	7/7 (100.0%)	7/7 (100.0%)	7/7 (100.0%)
Total			263	211/245 (86.1%)	102/253 (40.3%)	79/253 (31.2%)	121/245 (49.4%)	37/262 (14.1%)	14/245 (5.7%)	17/252 (6.7%)	11/215 (5.1%)	40/190 (21.0%)	74/206 (35.9%)	149/262 (56.9%)	180/196 (91.8%)	234/261 (89.7%)	226/251 (90.0%)

Legend	
*	Intrinsic Resistance Reported
-	Less than 4 isolates tested
^	No interpretive criteria

Percentage susceptible (%)				
0-20%	21-40%	41-60%	61-80%	81-100%

** No isolates tested reported susceptibility to Ampicillin or Ampicillin/Sulbactam

Note 1 Aminoglycoside susceptible/resistance differentiates between organisms with and without acquired resistance mechanisms only. For systemic infections aminoglycosides must be used in combination with other active therapy.

Carbapenemase producing Enterobacterales isolates identified or submitted to the Microbiological Diagnostic Unit Public Health Laboratory (MDU PHL) between 01/01/2020 and 31/12/2021 with available susceptibility data are included in the analysis above. Antimicrobial susceptibility testing was performed by broth microdilution, except for fosfomycin (agar dilution). Aggregate antibiograms exclude antimicrobials for an organism and carbapenemase gene combinations where fewer than five valid results were available. Aggregate antibiograms are categorised by carbapenemase gene sub-type(s) and isolates may contain other antimicrobial resistance (AMR) mechanisms not indicated. Carbapenemase gene subtypes have been determined by whole genome sequence analysis. EUCAST 2021 clinical breakpoints have been used for all susceptibility interpretations (1). Intermediate susceptibility included as susceptible in aggregate antibiograms.

(1) The European Committee on Antimicrobial Susceptibility Testing. Breakpoint Tables for Interpretation of MICs and Zone Diameters. Version 11.0, 2021. <http://www.eucast.org>.