## Northern Health Authority Annual Antibiogram 2025\*

Greater than or equal to 80% susceptibility = GOOD CHOICE Less than 80% susceptibility = POOR CHOICE Cefotaxime or Ceftriaxone Nitrofurantoin<sup>u</sup> Ciprofloxacin Moxifloxacin # of isolates Ceffazidine **Northern Health Authority Gram Negative Organisms** % Susceptible Escherichia coli 4362 88 62 87 92 100 93 94 83 74 97 Citrobacter freundii 79 R R R R R R R 100 92 96 89 87 91 279 R R R 99 94 94 88 88 34 Enterobacter cloacae complexa Klebsiella aerogenesb 65 R R R 100 92 95 97 91 162 88 R 91 59 92 88 94 99 94 95 94 95 96 Klebsiella oxytoca \_ R 97 Klebsiella pneumoniae 541 96 92 94 94 100 97 97 93 88 39 46 R R Morganella morganii R R 100 85 90 76 74 Proteus mirabilis 240 98 86 91 99 100 93 90 88 93 97 54 93 98 91 Serratia marcescens R R R R 93 76 R R Pseudomonas aeruginosa 324 R 89 R 93 90 99 R 87 R R R 92 Stenotrophomonas maltophilia 25 R R R Haemophilus influenzae 79 82 81 95 100 67 100

Notes

Percent susceptible for each organism/antimicrobial combination was generated by including the first isolate of that organism encountered on a given patient.

Organisms with less than 30 isolates have decreased statistical validity.

- Legend Drug not tested or indicated
  - Urinary tract infections only
  - R Intrinsic resistance
  - <sup>a</sup> Enterobacter cloacae complex includes Enterobacter asburiae, E. cloacae and E. hormaechei
  - Previously known as Enterobacter aerogenes



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Greater than or equal to 80%	Less than 80% susceptibility = POOR CHOICE																
Northern Health Authority Gram Negative Organisms % Susceptible	#ofisol	Amoxicillis	Ampicin:	Piperacillin,	Cefazoli.	Cefuros:	Cefixim	/ &	Ceftazia:	Merope	Gentam.	Tobram	TMP/SAA	Ciprofic	Moxiflox	Nitrofura	- σητοίηυ
Escherichia coli	4362	88	62	98	87	_	89	92	_	100	93	94	83	74	_	97	ı
Citrobacter freundii	79	R	R	R	R	R	R	R	_	100	92	96	89	87	-	91	
Enterobacter cloacae complex <sup>a</sup>	279	R	R	_	R	_	_	_	_	99	94	94	88	88	-	34	
Klebsiella aerogenes <sup>b</sup>	65	R	R	_	R	_	_	_	_	100	92	95	97	91	_	9	
Klebsiella oxytoca	162	88	R	91	59	-	92	88	94	99	94	95	94	95	_	96	
Klebsiella pneumoniae	541	96	R	97	92	-	93	94	94	100	97	97	93	88	-	39	
Morganella morganii	46	R	R	-	R	R	-	-	_	100	85	90	76	74	-	-	
Proteus mirabilis	240	98	86	100	91	-	97	99	-	100	93	90	88	93	-	-	
Serratia marcescens	54	R	R	_	R	R	_	_	_	93	93	76	98	91	_	R	
Pseudomonas aeruginosa	324	R	R	89	-	-	-	R	93	90	-	99	R	87	-	-	
Stenotrophomonas maltophilia	25	R	R	R	_	_	_	R	R	R	_	_	92	_	_	_	
Haemophilus influenzae	79	82	81	_	-	95	-	100	-	_	-	-	67	-	100	-	

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Northern Health Authority Gram Positive Organisms % Susceptible	# of isola.	Amoxic	Ampiciliis	Cloxacin	Penicillis	Cefazoli	/ 0	Erythrom	Clindam	Ciprofice	Moxiflox	Gent Su.	TMP-SM	Tetracuci.	Nitrofine	Vancom.	Linezolin	Di.
Staphylococcus aureus †	2398	_	_	76	_	76	_	74	86	_	_	-	92	96	-	100	-	
S. aureus (MRSA) ‡	571	R	R	R	R	R	R	R	88	-	_	-	89	92	-	100	100	
Staphylococcus epidermidis	303	-	_	53	_	54	_	47	75	_	_	_	77	89	-	100	-	
Staphylococcus lugdunensis	138	-	-	94	-	94	-	95	95	-	-	-	99	99	-	100	100	
Enterococcus faecalis	518	99 <sup>B</sup>	99	R	-	_	R	R	R	84	-	85	-	_	99	100	-	
Enterococcus faecium	95	R	R	R	-	_	R	R	R	12	-	88	-	-	_	50‡	93	
Group A Streptococcus	775	100°	100°	-	100	-	100	75	77	_	_	-	R	-	_	99	-	
Group B Streptococcus	265	100°	100°	-	100	-	100 <sup>\$</sup>	62	59	_	-	-	R	-	_	100	-	
Group C Streptococcus	111	100°	100 <sup>c</sup>	_	100	-	100 <sup>\$</sup>	76	74	_	_	-	R	-	_	100	-	
Group G Streptococcus	167	100°	100°	-	100	-	100 <sup>\$</sup>	77	75	_	_	_	R	-	_	100	-	
Streptococcus pneumoniae	76	87	-	-	100 <sup>2</sup> 72 <sup>3</sup>	-	100 <sup>2</sup> 87 <sup>3</sup>	64	-	_	100	-	66	_	_	100	-	

Notes Percent susceptible for each organism/antimicrobial combination was generated by including the first isolate of that organism encountered on a given patient.

Organisms with less than 30 isolates have decreased statistical validity.

#### Legend

- Drug not tested or indicated
- \$ AST method not validated for this organism/antibiotic combination
- Total Staphylococcus aureus isolates (includes MSSA and MRSA)
- # Represents clinical isolates excludes colonization
- <sup>u</sup> Urinary tract infections only
- R Intrinsic resistance
- Non-meningitis
- <sup>3</sup> Meningitis
- <sup>A</sup> Extrapolated from Cloxacillin
- <sup>B</sup> Extrapolated from Ampicillin
- c Extrapolated from Penicillin
- \* Prepared by Ally King, Regional Program Lead, Microbiology, based on 2024 NHA cumulative AST data from Jan. 1 2024 to Dec. 31 2024, As per CLSI document M39-A4 Analysis and presentation of cumulative antimicrobial susceptibility tests data approved guideline 4th edition

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Northern Health Authority Gram Positive Organisms % Susceptible	# of isola.	Amoxicin.	Ampicillis	Cloxacinii	Penicillis	Cefazolis	/ 0	Erythron	Clindam.	Ciprofic.	Moxiflox.	Gent Sur	TMP-SAN	Tetrac <sub>vo.</sub>	Nitrofur	Vancom	uycin Linezolia	7
Staphylococcus aureus †	2398	_	_	76	_	76	_	74	86	_	-	-	92	96	_	100	_	
S. aureus (MRSA) ‡	571	R	R	R	R	R	R	R	88	-	_	_	89	92	_	100	100	
Staphylococcus epidermidis	303	_	_	53	_	54	_	47	75	_	-	_	77	89	_	100	-	
Staphylococcus lugdunensis	138	_	_	94	_	94	-	95	95	-	_	-	99	99	-	100	100	
Enterococcus faecalis	518	99 <sup>B</sup>	99	R	-	_	R	R	R	84	-	85	-	_	99	100	-	
Enterococcus faecium	95	R	R	R	-	_	R	R	R	12	-	88	-	-	_	50‡	93	
Group A Streptococcus	775	100°	100°	-	100	_	100	75	77	_	-	-	R	-	_	99	_	
Group B Streptococcus	265	100 <sup>c</sup>	100 <sup>c</sup>	-	100	_	100 <sup>\$</sup>	62	59	_	_	-	R	_	_	100	_	
Group C Streptococcus	111	100°	100°	_	100	_	100 <sup>\$</sup>	76	74	_	_	_	R	_	_	100	-	
Group G Streptococcus	167	100°	100°	-	100	-	100 <sup>\$</sup>	77	75	-	-	-	R	-	-	100	-	
Streptococcus pneumoniae	76	87	-	-	100 <sup>2</sup> 72 <sup>3</sup>	-	100² 87³	64	-	-	100	-	66	-	-	100	-	

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<sup>\*</sup> Prepared by Ally King, Regional Program Lead, Microbiology, based on 2024 NHA cumulative AST data from Jan. 1 2024 to Dec. 31 2024, As per CLSI document M39-A4 Analysis and presentation of cumulative antimicrobial susceptibility tests data – approved guideline – 4th edition