Sullivan Nicolaides PATHOLOGY
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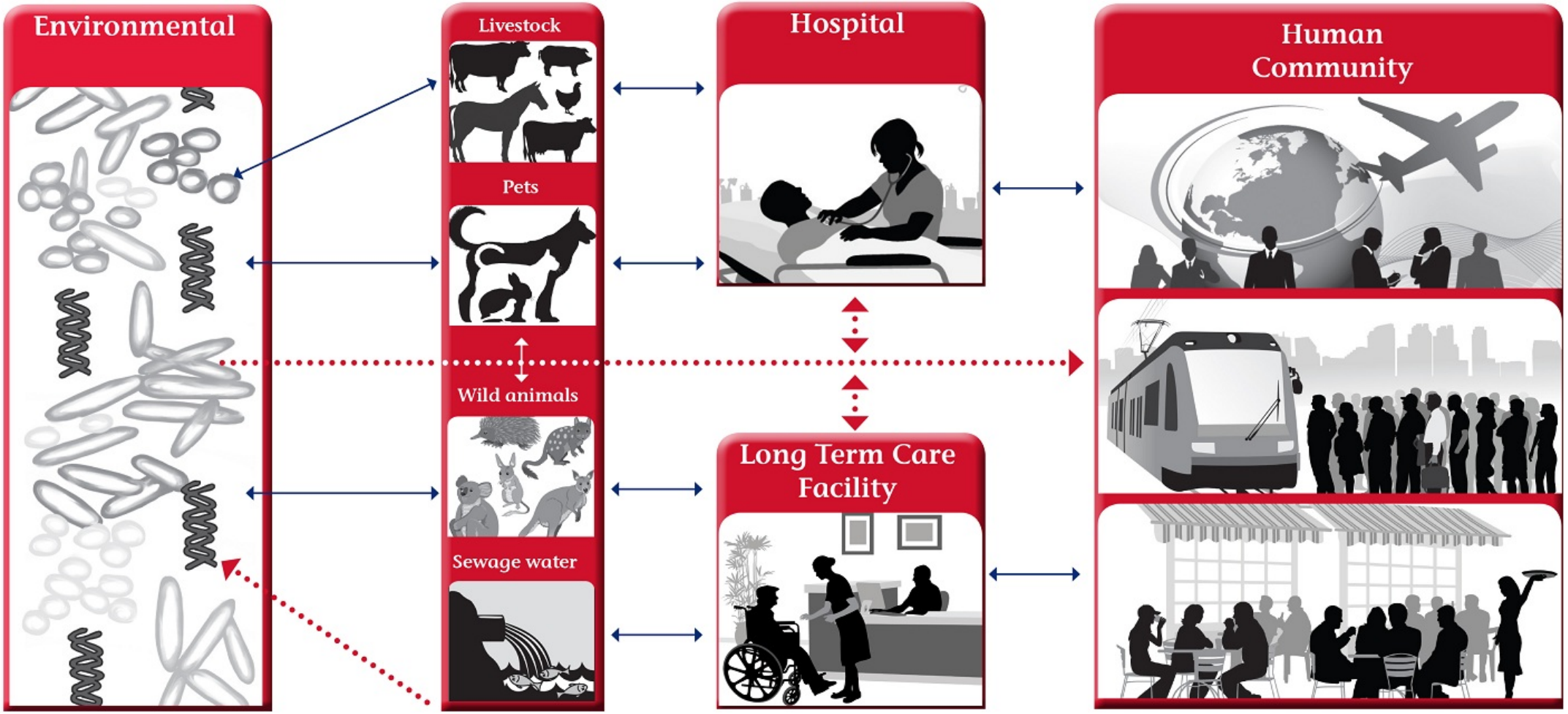
Community Antibiogram Jan - Dec 2017



Community Antibiogram Jan - Dec 2017

Cumulative antibiograms from community isolates can be a useful resource to inform empirical therapy recommendations as per Therapeutic Guidelines.

This document provides a cumulative antibiogram of isolates collected from patients in the community. Sullivan Nicolaides pathology services patients from all over Queensland as well as Northern New South Wales.

The data is divided into two parts

1. Urinary Antibiogram
   Note: only a subset of *S.saprophyticus* isolates have had routine susceptibility performed.
2. Non Urinary Antibiogram

Organisms are listed in descending order of frequency.
Organisms are colour coded according to whether they are Gram Positive or Gram Negative.
Only the first isolate of a given species per patient per year per subtype (e.g urine, non urine) is included.
Since 2012, susceptibility testing to produce these antibiograms is performed using EUCAST microbroth dilution and disc diffusion.[1]
Expert EUCAST rules in Antimicrobial Susceptibility testing have been applied.[2]
Where the total number of isolates tested is < 30, results are considered statistically invalid in accordance with CLSI M39-A4.[3]
Where only a subset (< 95%) of isolates from a particular organism group have been tested, reported susceptibilities are usually not indicative of the true susceptibility because of the selective nature of testing only more resistant isolates. These occasions are marked with an * and susceptibility results should be interpreted with caution.

Signal resistances:
Some important antibiotic resistances have their origin or circulate in the community. At the end of each antibiogram, signal resistances are summarised

These organisms include

- *Enterobacteriaceae* resistant to third or fourth generation cephalosporins due to the presence of Extended Spectrum Beta Lactamases (ESBLs)
- *Enterobacteriaceae* resistant to third or fourth generation cephalosporins due to the presence of Plasmid Mediated AMPC production (PAMPs)
- *Enterobacteriaceae* resistant to carbapenems due to the presence of a plasmid mediated carbapenamase (CPE)
- Non *Enterobacteriaceae* (e.g *Acinetobacter* spp; *Pseudomonas aeruginosa*) resistant to carbapenems due to the presence of a plasmid mediated carbapenamase (CPNE)
- Vancomycin resistant Enterococci (VRE)
- Methicillin resistant *Staphylococcus aureus* (MRSA)
- Vancomycin heteroresistant, intermediate and resistant *Staphylococcus aureus* (hVISA, VISA, VRSA)
- Penicillin intermediate and resistant *Streptococcus pneumoniae* noting that breakpoints differ according to clinical condition (meningitis, pneumonia, other) and mode of administration
- Penicillin intermediate and resistant viridans *Streptococci*

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Colour coding for all antibiograms tables is shown below.

	Gram Positive Organism		≥90% of isolates susceptible		≥90% of isolates susceptible (where sample size <95% of total isolates tested)
	Gram Negative Organism		70-89% of isolates susceptible		70-89% of isolates susceptible (where sample size <95% of total isolates tested)
	Antibiotic Not recommended to be used in children without specialist advice		<70% of isolates susceptible		<70% of isolates susceptible (where sample size <95% of total isolates tested)
	Restricted or 2nd Line Antibiotics	**R**	Intrinsic Resistance is present with this organism-antibiotic combination	*	Sample size <95% of the total isolates tested
	Restricted or 2nd Line Antibiotics and Antibiotic Not recommended to be used in children without specialist advice	**%**	Percentage of isolates sensitive to this particular antibiotic	**n**	Number of isolates tested with this antibiotic

References:

1. http://www.eucast.org/clinical_breakpoints/
2. http://www.eucast.org/expert_rules_and_intrinsic_resistance/
3. Clinical and Laboratory Standards Institute (CLSI). Guideline for Analysis and Presentation of Cumulative Antimicrobial Susceptibility (M39-A4). 3rd ed. Pennsylvania: CLSI, 2014

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Urine Antibiogram

All SNP Community

Organism Group	No. Organisms	%Total		Amoxicillin	Amoxicillin-clavulanate	Ticarcillin-clavulanate	Pipericillin-tazobactam	Cefalexin	Ceftriaxone	Ceftazidime	Meropenem	Gentamicin	Amikacin	Trimethoprim	Nitrofurantoin	Norfloxacin	Ciprofloxacin	Fusidic Acid	Rifampicin	Gentamicin (High Level)	Fosfomycin	Quinupristin-dalfopristin	Vancomycin
All isolates	**95487**	**100**																					
Escherichia spp	62864	65.8	%	58	88	64	95	93	92		100	95	99	77	99	92					99		
			n	62853	*53285	*30573	*31771	62690	*31931		*30726	*31990	*30775	62851	62813	62863					*2391		
Enterococcus spp	9212	9.6	%	99				10						R	99	96			20	65		45	87
			n	9209				*10							9189	9197			*176	*181		*179	*210
Klebsiella spp	7449	7.8	%	R	96	86	91	95	95		100	98	100	90	85	96							
			n		*6163	*3070	*3444	7420	*3459		*3110	*3464	*3117	7447	7442	7449							
Staphylococcus saprophyticus	2438	2.6	%	21	97									93	100	100							
			n	*460	*471									2422	2432	2430							
Proteus mirabilis	2326	2.4	%	89	96	99	100	98	99		100	98	99	86	R	98							
			n	2324	*1964	*1104	*1193	2326	*1199		*1111	*1199	*1114	2326		2326							
Pseudomonas aeruginosa	2136	2.2	%	R	R	44	94	R	R	96	96	96	95	R	R	95	92						
			n			*1790	2132			2136	*1812	2136	*1820			2062	2128						
Enterobacter cloacae complex	1648	1.7	%	R	R	67	80	R	80		99	97	99	86	86	95							
			n			*1192	*1387		*1433		*1222	*1434	*1229	1648	1646	1648							
Citrobacter koseri, amalonaticus	1364	1.4	%	R	97	94	96	94	98		100	99	100	98	96	99							
			n		*1125	*626	*690	1362	*691		*633	*694	*633	1364	1360	1364							
ß haemolytic Streptococci Group B	1300	1.4	%	100					100					91	100	75							100
			n	1300					*14					1300	1294	*1008							*15
Enterobacter aerogenes	1001	1.0	%	R	R	82	83	R	86		100	99	100	97	72	97							
			n			*733	*788		*790		*742	*791	*743	1001	1001	1001							

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Urine Antibiogram

All SNP Community

Organism Group	No. Organisms	%Total		Amoxicillin	Amoxicillin-clavulanate	Ticarcillin-clavulanate	Pipericillin-tazobactam	Cefalexin	Ceftriaxone	Ceftazidime	Meropenem	Gentamicin	Amikacin	Trimethoprim	Nitrofurantoin	Norfloxacin	Ciprofloxacin	Fusidic Acid	Rifampicin	Gentamicin (High Level)	Fosfomycin	Quinupristin-dalfopristin	Vancomycin
All isolates	**95487**	**100**																					
Coagulase negative Staphylococci	770	0.8	%	22	73									61	100	89		70	90				100
			n	761	769									766	765	761		*89	*89				*90
Staphylococcus aureus (ALL)	655	0.7	%	24	88									93	100	94		95	100				100
			n	655	654									646	644	636		*102	*102				*103
Morganella spp	540	0.6	%	R	R	85	99	R	92		99	96	99	89	R	94							
			n			*492	537		537		*497	537	*498	540		540							
Citrobacter freundii complex	424	0.4	%	R	R	67	76	R	71		100	96	99	89	97	96							
			n			*293	*331		*333		*302	*333	*303	424	424	424							
Serratia spp	353	0.4	%	R	R	93	97	R	97		99	97	98	92	R	93							
			n			*287	346		347		*290	347	*290	353		353							
viridans Streptococci	190	0.2	%	97										65	99	59							
			n	189										190	190	188							
Proteus spp other	120	0.1	%	R	52	100	100	R	84		99	97	100	78	R	99							
			n		*104	*86	*107		*107		*87	*107	*87	120		120							
Providencia spp	92	0.1	%	R	R	98	99	R	95		100	84	99	89	R	99							
			n			*86	92		92		*86	92	*86	92		92							
Acinetobacter spp	84	0.1	%	R	R	91	95	R	R		100	100	99	R	2	82							
			n			*67	*77				*71	84	*71		84	*73							
Miscellaneous GPC†	77	0.1	%	97										21	99	91							
			n	76										*72	77	77							

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Organism Group	No. Organisms	%Total		Amoxicillin	Amoxicillin-clavulanate	Ticarcillin-clavulanate	Pipericillin-tazobactam	Cefalexin	Ceftriaxone	Ceftazidime	Meropenem	Gentamicin	Amikacin	Trimethoprim	Nitrofurantoin	Norfloxacin	Ciprofloxacin	Fusidic Acid	Rifampicin	Gentamicin (High Level)	Fosfomycin	Quinupristin-dalfopristin	Vancomycin
All isolates	**95487**	**100**																					
ß haemolytic Streptococci Group A	71	0.1	%	100										100	100	89							
			n	71										*65	*65	*64							
Raoultella spp	59	0.1	%	R	93	96	94	93	94		100	97	96	93	97	98							
			n		*41	*24	*35	59	*35		*25	*35	*25	59	59	59							
Miscellaneous Enterobacteriaceae†	53	0.1	%	32	80	85	80	72	95		100	100	100	92	91	96							
			n	53	*49	*34	*40	53	*42		*36	*42	*36	53	53	53							
Salmonella spp	53	0.1	%	94	98				92					98	100	98							
			n	53	*47				*12					53	53	53							
ß haemolytic Streptococci Other (nonA nonB)	44	0.0	%	100										100	100	86							
			n	44										*41	*41	*36							
Pseudomonas spp	30	0.0	%	R	0	0	77	0			57	100	95	3	3	89							
			n		*25	*20	30	30			*21	30	*21	30	30	*19							

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Signal Resistances: Where the tables below contain no data no multiresistant organisms have been detected.

Urine: Extended spectrum beta lactamase producing Enterobacteriaceae (ESBL)

Organism Group	Organism Name	No. Positive	% of Strain
Citrobacter freundii complex n = 424	Citrobacter braakii	1	0.2
	Citrobacter freundii	8	1.9
Citrobacter koseri, amalonaticus n = 1364	Citrobacter koseri	3	0.2
Enterobacter aerogenes n = 1001	Enterobacter aerogenes	4	0.4
Enterobacter cloacae complex n = 1648	Enterobacter cloacae	30	1.8
Escherichia spp n = 62868	Escherichia coli	2155	3.4
Klebsiella spp n = 7450	Klebsiella oxytoca	5	0.1
	Klebsiella pneumoniae	125	1.7
Morganella spp n = 540	Morganella morganii	1	0.2
Proteus mirabilis n = 2327	Proteus mirabilis	4	0.2
Proteus spp other n = 120	Proteus penneri	2	1.7
Raoultella spp n = 59	Raoultella ornithinolytica	1	1.7
Serratia spp n = 353	Serratia marcescens	2	0.6

Urine: Plasmid mediated AMPC producing Enterobacteriaceae (PAMP)

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Organism Group	Organism Name	No. Positive	% of Strain
Citrobacter koseri, amalonaticus n = 1364	Citrobacter koseri	7	0.5
Enterobacter aerogenes n = 1001	Enterobacter aerogenes	1	0.1
Enterobacter cloacae complex n = 1648	Enterobacter cloacae	1	0.1
Escherichia spp n = 62868	Escherichia coli	1090	1.7
Klebsiella spp n = 7450	Klebsiella oxytoca	7	0.1
	Klebsiella pneumoniae	31	0.4
Proteus mirabilis n = 2327	Proteus mirabilis	8	0.3
Proteus spp other n = 120	Proteus penneri	4	3.3
	Proteus vulgaris	1	0.8
Raoultella spp n = 59	Raoultella ornithinolytica	1	1.7

Urine: Plasmid mediated Carbapenemase producing Enterobacteriaceae (CPE)

Organism Group	Organism Name	No. Positive	% of Strain
Citrobacter freundii complex n = 424	Citrobacter freundii	1	0.2
Enterobacter cloacae complex n = 1648	Enterobacter cloacae	2	0.1
Escherichia spp n = 62868	Escherichia coli	1	0.0
Klebsiella spp n = 7450	Klebsiella pneumoniae	2	0.0

Urine: Plasmid mediated Carbapenemase producing Non Enterobacteriaceae (CPNE)

Organism Group	Organism Name	No. Positive	% of Strain

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Urine: Vancomycin Resistant Enterococci (VRE)			
Organism Group	**Organism Name**	**No. Positive**	**% of Strain**
Enterococcus spp n = 9213	Enterococcus faecalis(VRE) - VAN-A	1	0.0
	Enterococcus faecium (VRE) - VAN-A	9	0.1
	Enterococcus faecium (VRE) - VAN-B	12	0.1

Urine: Methicillin Resistant Staphylococcus aureus (MRSA)			
Organism Group	**Organism Name**	**No. Positive**	**% of Strain**
Staphylococcus aureus (ALL) n = 655	S.aureus (non-multiresistant MRSA)	54	8.2
	Staphylococcus aureus (MRSA)	1	0.2
	Staphylococcus aureus (UK EMRSA-15)	23	3.5

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Non-urine Antibiogram

All SNP Community

Organism Group	No. Organisms	%Total		Penicillin	Amoxicillin	Flucloxacillin	Amoxicillin-clavulanate	Ticarcillin-clavulanate	Pipericillin-tazobactam	Cefalotin	Cefazolin	Ceftriaxone	Ceftazidime	Cefepime	Meropenem	Gentamicin	Amikacin	Sulpha-trimethoprim	Norfloxacin	Ciprofloxacin	Fusidic Acid	Rifampicin	Gentamicin (High Level)	Erythromycin/Clarithromycin	Azithromycin	Clindamycin	Tetracycline	Mupirocin	Quinupristin-dalfopristin	Vancomycin
All isolates	**73173**	**100**																											
Staphylococcus aureus (ALL)	**41197**	**56.3**	%	**18**		**89**	**89**			**89**								**98**		**97**	**92**	**100**		**83**		**84**	**97**	97	90	**100**
			n	41179		41190	41158			41193								41192		*3801 8	*3803 9	*3803 9		41195		41171	41192	*3409 2	*10	*3802 4
ß haemolytic Streptococci Group A	**7897**	**10.8**	%	**100**						**100**		99						**99**						**93**		**95**	**85**			100
			n	7893						7835		*5002						7894						7894		7889	7893			*5175
Pseudomonas aeruginosa	**7876**	**10.8**	%		R		R	52	**97**		R	R	**98**	**98**	**98**	**96**	95	R		**96**										
			n					*6182	7868				7870	7625	7872	7876	*6230			7870										
Haemophilus influenzae	**4512**	**6.2**	%		68		**86**					80						**71**		93							**99**			
			n		4511		4510					*15						4512		*15							4511			
ß haemolytic Streptococci Other (nonA nonB)	**1786**	**2.4**	%	**100**						**100**		97						**100**						**81**		**83**	**72**			100
			n	1786						1769		*864						1786						1784		1781	1785			*896
Campylobacter spp	**1506**	**2.1**	%																	**74**				**94**	**94**		**81**			
			n																	1506				1506	1505		1506			
Moraxella catarrhalis	**1244**	**1.7**	%		1		**100**											**92**						**100**			**100**			
			n		1244		1244											1244						1243			1244			
Salmonella spp	**1214**	**1.7**	%		**91**		**98**					**99**			**100**			**98**	**94**	**96**					85		50			
			n		1212		1190					1183			1180			1214	1214	1183					*13		*10			
Streptococcus pneumoniae	**1125**	**1.5**	%	**62**						81		84						**72**						**73**		**77**	**76**			100
			n	1121						*86		*676						1123						1122		*1030	1124			*683

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Non-urine Antibiogram

All SNP Community

Organism Group	No. Organisms	%Total		Penicillin	Amoxicillin	Flucloxacillin	Amoxicillin-clavulanate	Ticarcillin-clavulanate	Pipericillin-tazobactam	Cefalotin	Cefazolin	Ceftriaxone	Ceftazidime	Cefepime	Meropenem	Gentamicin	Amikacin	Sulpha-trimethoprim	Norfloxacin	Ciprofloxacin	Fusidic Acid	Rifampicin	Gentamicin (High Level)	Erythromycin/Clarithromycin	Azithromycin	Clindamycin	Tetracycline	Mupirocin	Quinupristin-dalfopristin	Vancomycin
All isolates	**73173**	**100**																											
Aeromonas spp	**1108**	**1.5**	%		R		62	86	92		0	99		100	78	100	100	97	97	100							92			
			n				1066	*973	*89		*496	*1045		*1029	*1023	1070	*310	1108	*1034	*1050							*13			
Coagulase negative Staphylococci	**517**	**0.7**	%	41		80	80			80								92		94	92	100		79		87	84			100
			n	515		514	515			516								516		*483	*482	*482		516		516	517			*484
viridans Streptococci	**403**	**0.6**	%	97						100		88						92					100	82		90	69			99
			n	402						*270		*103						*293					*17	401		400	403			*115
Escherichia spp	**402**	**0.5**	%		53		78	63	94		82	92		94	100	94	98	80		89										
			n		402		402	*373	400		402	401		401	401	402	*377	402		401										
Neisseria gonorrhoeae	**238**	**0.3**	%	5								100								76					95					
			n	238								238								237					232					
Klebsiella spp	**206**	**0.3**	%		R		98	95	96		84	99		99	100	99	100	96		99										
			n				206	*183	205		206	206		205	203	206	*184	206		203										
Corynebacterium spp	**184**	**0.3**	%	79						92								66						70		61	99			100
			n	184						*71								*133						*135		184	184			*126
Fastidious GNB†	**170**	**0.2**	%	91	100		100		100		100	100		100	100	85		99		100							99			
			n	*74	166		168		*10		*38	*19		*34	*16	*34		169		*24							*159			
Enterobacter cloacae complex	**169**	**0.2**	%		R		R	85	94		R	93		99	100	99	100	95		99										
			n					*143	167			169		167	169	169	*143	169		169										

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Non-urine Antibiogram

All SNP Community

Organism Group	No. Organisms	%Total		Penicillin	Amoxicillin	Flucloxacillin	Amoxicillin-clavulanate	Ticarcillin-clavulanate	Pipericillin-tazobactam	Cefalotin	Cefazolin	Ceftriaxone	Ceftazidime	Cefepime	Meropenem	Gentamicin	Amikacin	Sulpha-trimethoprim	Norfloxacin	Ciprofloxacin	Fusidic Acid	Rifampicin	Gentamicin (High Level)	Erythromycin/Clarithromycin	Azithromycin	Clindamycin	Tetracycline	Mupirocin	Quinupristin-dalfopristin	Vancomycin
All isolates	**73173**	**100**																											
Stenotrophomonas maltophilia	**165**	**0.2**	%															**97**												
			n															165												
Serratia spp	**164**	**0.2**	%		R		R	**98**	**100**		R	**100**		**100**	**100**	**100**	**99**	**99**		**98**										
			n					*149	164			163		161	162	164	*149	164		162										
Non fermenting GNB†	**162**	**0.2**	%		**8**		79	94	98		3	57		40	95	33	18	**95**		**57**							100			
			n		160		*131	*62	*127		*134	*99		*119	*126	*134	*74	159		160							*52			
Vibrio spp	**95**	**0.1**	%		**22**		97		100		52	100		100	100	100		**98**		**100**							**99**			
			n		92		*32		*24		*50	*78		*32	*24	*28		95		92							95			
Yersinia spp	**93**	**0.1**	%		**1**		6	93	100		0	99		99	100	100	100	**99**		100										
			n		93		*80	*80	*38		*77	*81		*81	*81	*82	*81	93		*81										
ß haemolytic Streptococci Group B	**89**	**0.1**	%	**99**						100		100						**99**						**74**		**75**	**23**			100
			n	88						*69		*58						87						86		87	87			*62
Proteus mirabilis	**80**	**0.1**	%		**90**		**98**	**100**	**100**		**86**	**100**		**100**	**100**	**100**	**99**	**89**		**100**										
			n		80		80	76	80		79	80		79	79	80	76	80		79										
Enterococcus spp	**75**	**0.1**	%	90						R								R				25	70	R		R	**26**		34	95
			n	*67																		*16	*23				*68		*29	*66
Pseudomonas spp	**73**	**0.1**	%		R		**8**	13	**79**		R	100		**96**	**75**	**97**	98	**14**		**100**										
			n				73	*48	73			*10		70	73	73	*48	73		73										

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Organism Group	No. Organisms	%Total		Penicillin	Amoxicillin	Flucloxacillin	Amoxicillin-clavulanate	Ticarcillin-clavulanate	Pipericillin-tazobactam	Cefalotin	Cefazolin	Ceftriaxone	Ceftazidime	Cefepime	Meropenem	Gentamicin	Amikacin	Sulpha-trimethoprim	Norfloxacin	Ciprofloxacin	Fusidic Acid	Rifampicin	Gentamicin (High Level)	Erythromycin/Clarithromycin	Azithromycin	Clindamycin	Tetracycline	Mupirocin	Quinupristin-dalfopristin	Vancomycin
All isolates	**73173**	**100**																											
Acinetobacter spp	**61**	**0.1**	%		R		R	76	87		R	R	45	55	100	98	96	95		100										
			n					*51	*45				60	60	61	61	*52	61		61										
Shigella spp	**44**	**0.1**	%		53		72	51	97		47	92		94	100	69	69	30	68	86					27					
			n		43		*36	*35	*32		*34	*36		*36	*36	*35	*36	44	44	*36					*11					
Enterobacter aerogenes	**40**	**0.1**	%		R		R	92	93		R	95		100	100	100	100	100		98										
			n					*36	40			40		40	40	40	*36	40		40										
Neisseria meningitidis	**38**	**0.1**	%	29								100								97						100				
			n	*35								38								38						*20				
Nocardia spp	**35**	**0.0**	%				22					56					100	97		21			33							
			n				*32					34					34	*33		34			*21							
Morganella spp	**33**	**0.0**	%		R		R	90	100		R	91		97	100	97	100	94		94										
			n					*31	33			32		33	33	33	*31	33		33										

Community Antibiogram Jan - Dec 2017

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Signal Resistances: Where the tables below contain no data no multiresistant organisms have been detected.

NonUrine: Extended spectrum beta lactamase producing Enterobacteriaceae (ESBL)

Organism Group	Organism Name	No. Positive	% of Strain
Enterobacter cloacae complex n = 171	Enterobacter cloacae	2	1.2
Escherichia spp n = 490	Escherichia coli	25	5.1
Klebsiella spp n = 215	Klebsiella pneumoniae	3	1.4
Salmonella spp n = 1215	Salmonella species	2	0.2

NonUrine: Plasmid mediated AMPC producing Enterobacteriaceae (PAMP)

Organism Group	Organism Name	No. Positive	% of Strain
Escherichia spp n = 490	Escherichia coli	7	1.4
Klebsiella spp n = 215	Klebsiella pneumoniae	1	0.5

NonUrine: Plasmid mediated Carbapenemase producing Enterobacteriaceae (CPE)

Organism Group	Organism Name	No. Positive	% of Strain

NonUrine: Plasmid mediated Carbapenemase producing Non Enterobacteriaceae (CPNE)

Organism Group	Organism Name	No. Positive	% of Strain

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NonUrine: Vancomycin Resistant Enterococci (VRE)			
Organism Group	**Organism Name**	**No. Positive**	**% of Strain**
Enterococcus spp n = 80	Enterococcus faecium (VRE) - VAN-B	3	3.8

NonUrine: Methicillin Resistant Staphylococcus aureus (MRSA)			
Organism Group	**Organism Name**	**No. Positive**	**% of Strain**
Staphylococcus aureus (ALL) n = 41201	S.aureus (non-multiresistant MRSA)	3978	9.7
	Staphylococcus aureus (MRSA)	104	0.3
	Staphylococcus aureus (UK EMRSA-15)	336	0.8

NonUrine: Streptococcus pneumoniae Penicillin Susceptibility ** (Oral penicillin V breakpoints)			
Organism Group	**MIC category**	**No.**	**% of Strain**
Streptococcus pneumoniae n = 1123	Sensitive ≤ 0.06 mg/L	697	62.1
	Intermediate 0.12 - 1 mg/L	405	36.1
	Resistant ≥ 2 mg/L	21	1.9

NonUrine: Streptococcus alpha haem (viridans Streptococci)			
Organism Group	**MIC category**	**No.**	**% of Strain**
viridans Streptococci n = 398	Sensitive ≤ 0.06 mg/L	387	97.2
	Intermediate 0.12 - 1 mg/L	11	2.8

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Blood Culture Antibiogram															All SNP Community
Organism Group	**No. Organisms**	**%Total**		Penicillin	Flucloxacillin	Amoxicillin-clavulanate	Cefalotin	Sulpha-trimethoprim	Ciprofloxacin	Fusidic Acid	Rifampicin	Erythromycin/Clarithromycin	Clindamycin	Tetracycline	Vancomycin
All isolates	**195**	**100.0**													
Coagulase negative Staphylococci	82	42.1	%	21	64	64	64	79	89	91	100	65	78	80	99
			n	82	81	81	81	81	79	78	78	81	82	82	82

Signal Resistances: Where the tables below contain no data no multiresistant organisms have been detected.

Blood Culture: Extended spectrum beta lactamase producing Enterobacteriaceae (ESBL)

Organism Group	Organism Name	No. Positive	% of Strain
Escherichia spp n = 28	Escherichia coli	1	3.6

Blood Culture: Plasmid mediated AMPC producing Enterobacteriaceae (PAMP)

Organism Group	Organism Name	No. Positive	% of Strain

Blood Culture: Plasmid mediated Carbapenemase producing Enterobacteriaceae (CPE)

Organism Group	Organism Name	No. Positive	% of Strain

Blood Culture: Plasmid mediated Carbapenemase producing Non Enterobacteriaceae (CPNE)

Community Antibiogram Jan - Dec 2017

Organism Group	Organism Name	No. Positive	% of Strain
Blood Culture: Vancomycin Resistant Enterococci (VRE)			
Organism Group	**Organism Name**	**No. Positive**	**% of Strain**
Blood Culture: Methicillin Resistant Staphylococcus aureus (MRSA)			
Organism Group	**Organism Name**	**No. Positive**	**% of Strain**
Staphylococcus aureus (ALL) n = 14	S.aureus (non-multiresistant MRSA)	1	7.1

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Blood Culture: Streptococcus pneumoniae Penicillin Susceptibility ** (Oral penicillin V breakpoints)			
Organism Group	**MIC category**	**No.**	**% of Strain**
Streptococcus pneumoniae n = 4	Sensitive ≤ 0.06 mg/L	3	75.0
	Intermediate 0.12 - 1 mg/L	1	25.0

Blood Culture: Streptococcus alpha haem (viridans Streptococci)			
Organism Group	**MIC category**	**No.**	**% of Strain**
viridans Streptococci n = 89	Sensitive ≤ 0.06 mg/L	77	86.5
	Intermediate 0.12 - 1 mg/L	9	10.1
	Resistant ≥ 2 mg/L	3	3.4

† Miscellaneous organism groupings detail report.