## 2016 Alaska State Antibiogram

The following tables show the proportion of isolates of various bacterial species that tested susceptible to various antibiotics during 2016. These data were aggregated from the antibiograms produced by Alaska hospitals in order to create aggregate regional resistance pattern summaries. These antibiograms can be helpful for health care providers in selecting appropriate "presumptive" antimicrobial therapy for their patients until specific individual laboratory test results are available. They can also be helpful for determining antibiotic stewardship priorities within hospitals and emerging resistance patterns in a broader service area.

- **Methodology:** Individual hospitals prepared their own facility antibiograms, which were shared with the Alaska Section of Epidemiology. Aggregated susceptibility percentages were calculated as the proportion of all tested isolates for the region that were susceptible. Values are only reported when more than one facility provided data for the given species-antibiotic combination. Intrinsic resistance is indicated with an "R", following the guidance of CLSI document M100-S24.
- Multi-Drug Resistant Organisms of Note:
  - o Vancomycin-resistant Staphylococcus aureus (VRSA): no cases of VRSA have ever been reported in Alaska. VRSA is reportable to the Alaska Section of Epidemiology.
  - o Carbapenem-resistant Enterobacteriaceae (CRE): there were 11 cases of CRE reported in Alaska in 2016.
- Legend:
  - o The top value in each square is the percent of isolates of that species that tested susceptible to that antibiotic.
  - o The lower value in each square indicates the number of tested isolates for that bacteria-antibiotic combination.
  - o "R" indicates intrinsic resistance to that antibiotic, while "S" indicates definitional susceptibility.
  - o "NED" indicates that there was Not Enough Data to report the value: either only one facility reported data for that drug-bug combination or <30 isolates were tested.
- **Limitations:** Individual facilities often use different methods to test for antimicrobial susceptibility, different methods to build their antibiograms, and different antibiotics in their pharmacies. These factors limit interpretation of these data. Additionally, antimicrobial susceptibility testing done in the laboratory does not always predict how effective that drug will be when used to treat a patient. Data are not stratified by infection site, which influences antibiotic choice and effectiveness.
- **Contributing Facilities:** Thanks to all the hospitals in Alaska for participating in this project to the extent of their ability. These statewide data include all the hospitals used in the Regional Antibiograms, plus Fairbanks Memorial Hospital.

For more information and the methods used for the analyses, please see the "Regional Antibiogram Project — Alaska, 2014–2015" Epidemiology *Bulletin*.

MSSA   13%   3%   S   98%   99%   100%   100%   85%   85%   94%   98%   99%   99%   99%   99%   99%   98%   99%   99%   99%   98%   99%	Statewide data  Species	Penicillin	Ampicillin	Oxacillin	Ampicillin-sulbactam	Amoxicillin	Cefazolin	Ceftriaxone	Cefotaxime	Ciprofloxacin	Levofloxacin	Moxifloxacin	Daptomycin	Clindamycin	Erythromycin	Vancomycin	Gentamicin	Gent Syn	Trimethoprim-sulfamethoxazole	Linezolid	Tetracycline	Nitrofurantoin	Quinupristin-dalfopristin	Rifampin
MSSA   13%   3%   S   98%   99%   100%   100%   85%   85%   94%   98%   88%   68%   99%   99%   99%   99%   99%   99%   99%   99%   99%   99%   98%   145%	Total Staphylococcus aureus																							99%
MRSA	ΛΛΟΟΛ																							
MRSA	IVISSA			3																				
Staphylococcus lugdunensis   NED   93%   (201)   (201)   (370)   (484)   (957)   (1276)   (207)   (231)   (2391)   (1248)   (248)   (1062)   (2316)   (1687)   (2307)   (2002)   (50)   (951	MRSA			R																				99%
Staphylococcus lugdunensis   NED   93%   (60)   (		(978)			(201)	(201)	(370)	(484)		(957)			(231)	(2391)		(2468)	(1062)		(2316)	(1687)		(2002)	(50)	(951)
Coag-negative Staphylococcus   15%   0%   51%   51%   51%   48%   49%   73%   72%   82%   98%   61%   63%   99%   99%   99%   69%   99%   89%   99%   98%   98%   98%   61%   63%   99%   99%   89%   99%   99%   99%   99%   99%   99%   99%   99%   88%   99%   99%   88%   98%   99%   99%   88%   99%   99%   88%   98%   99%   99%   88%   98%   99%   99%   88%   99%   99%   88%   99%   99%   88%   98%   99%   88%   98%   99%   88%   98%   99%   88%   98%   99%   88%   98%   99%   88%   99%   88%   99%   88%   98%   99%   88%	Staphylococcus lugdunensis	NED		93%						97%	93%			90%	88%	100%	100%		NED	NED	98%	100%		
Composition				(60)						(60)	(60)			(60)	(60)	(60)	(60)				(60)	(60)		
R R R R R R R R R R R R R R R R R R R	Coag-negative Staphylococcus	15%	0%	51%	51%	51%	48%	49%		73%	72%	82%		61%	63%	99%	90%		69%	99%	89%	99%	98%	98%
Composition				(1310)	(239)	(239)						(82)											-	(516)
R   R   R   R   R   R   R   R   R   R	Enterococcus faecalis						R	R	R					R			R		R				R	
Composition	5.4		,				-	-					(233)	-	(295)	· ·			-					(121)
Enterococcus spp. 98% 99% (417) (431) 88% 90% (137) (362) 99% NED 100% 99% (294) (376) 98% (137) (362) 100% (431) 100% (294) (376) 100% (129) 100% 100% 100% 100% 100% 100% 100% 100	Enterococcus Jaecium	NED					К	ĸ	К	NED	NED			К			к		К		NED	NED		
Comp B Streptococcus   100%   S   Comp B Streptococcus   100%   Comp B Streptococcus	Enterococcus spp.	98%								88%	90%											99%		
Streptococcus pneumoniae (all)   92%   103   103   104   105   1																		1125						
Streptococcus pneumoniae (all)         92%         96%         96%         98%         98%         100%         93%	Group B Streptococcus									, ,	,			51%	NED	` ′								
(103)     (84)     (89)     (141)     (191)     (134)     (242)     (82)     (88)       S. pneumoniae - oral     64%     (193)     (193)     97%     88%     88%     88%     88%		(129)												(128)		(129)								
S. pneumoniae - oral 64% (193) S. pneumoniae - non-CSF 74% 97% 88%	Streptococcus pneumoniae (all)	92%						96%	96%		98%			92%	74%	100%			93%		93%			
(193)         97%         88%		(103)						(84)	(89)		(141)			(191)	(134)	(242)			(82)		(88)			
S. pneumoniae - non-CSF 74% 97% 88%	S. pneumoniae - oral																							
(38/)	S. pneumoniae - non-CSF																							.
	S nneumonige - meningitic																							
S pneumoniae - meningitis   65%         95%   86%	5 pheamomae - meimigitis																							
Viridans-group Streptococcus NED 97% 100%	Viridans-group Streptococcus							·	(307)							100%								
(31)	O																							.

Statewide data  Species	Amoxicillin+ clavulanic acid	Ampicillin	Ampicillin+Sulbactam	Piperacillin+Tazobactam	Cefazolin	Cefuroxime	Ceftriaxone	Ceftazidime	Cefepime	Cefotaxime	Cefotetan	Cefoxitin	Aztreonam	Gentamicin	Tobramycin	Amikacin	Ertapenem	Imipenem	Meropenem	Ciprofloxacin	Levofloxacin	Trimeth+Sulfa	Tetracycline	Nitrofurantoin
Citrobacter freundii	R	R	R	98%	R	R	96%	95%	84%	NED	R	R	NED	100%	100%	100%	NED	100%	100%	97%	97%	92%	NED	95%
				(124)			(124)	(123)	(96)					(124)	(118)	(103)		(34)	(77)	(124)	(124)	(124)		(110)
Enterobacter aerogenes	R	R	R	89%	R	R	86%	89%	100%	77%	R	R	83%	100%	98%	100%	NED	NED	100%	99%	99%	96%	98%	24%
	_		_	(127)	_		(127)	(76)	(89)	(40)	_	_	(83)	(94)	(107)	(101)			(101)	(127)	(127)	(127)	(58)	(118)
Enterobacter cloacae	R	R	R	88%	R	R	85%	86%	98%	78%	R	R	84%	98%	97%	100%	97%	99%	98%	98%	96%	92%	91%	38%
Frank reighting on li	070/	F.C.0/	F00/	(377)	040/	060/	(377)	(288)	(307)	(166)	000/	0.40/	(223)	(377)	(344)	(258)	(32)	(67)	(312)	(377)	(377)	(377)	(193)	(351)
Escherichia coli	87% (7022)	56%	58%	97%	<b>91%</b> (12125)	86%	<b>97%</b> (12125)	<b>97%</b> (7608)	92%	73%	98%	94%	91%	93%	95%	99%	99%	99%	<b>100%</b> (8001)	<b>85%</b> (12125)	85% (12125)	<b>77%</b> (12125)	81%	98%
ESBL E. coli	(7832) NED	(11582) <b>0%</b>	(10551) <b>83%</b>	(11493) <b>93%</b>	0%	(7827) NED	0%	0%	(8467)	(4258)	(543)	(3110)	(3/16)	(12124) <b>76%</b>	(10835)	(5990)	(3601)	(3879) NED	100%	41%	(12125) <b>41%</b>	38%	(5558) <b>48%</b>	(11813) <b>89%</b>
ESBL E. COII	INED	(58)	(30)	(58)	(58)	NED	(58)	(30)						(58)				INED	(31)	(58)	(58)	(58)	(31)	(54)
Klebsiella oxytoca	92%	0%	58%	92%	61%	87%	95%	99%	98%	NED		99%	86%	99%	98%	100%		100%	100%	96%	96%	93%	NED	85%
	(85)	(36)	(205)	(205)	(154)	(204)	(205)	(204)	(170)			(85)	(169)	(205)	(205)	(185)		(36)	(150)	(205)	(205)	(205)		(179)
Klebsiella pneumoniae	98%	R	89%	97%	94%	93%	98%	67%	97%	98%	98%	95%	99%	99%	98%	99%	99%	100%	99%	96%	97%	95%	88%	50%
·	(837)		(1365)	(1365)	(1358)	(1091)	(1365)	(1130)	(1103)	(440)	(63)	(192)	(739)	(1365)	(1293)	(817)	(213)	(282)	(993)	(1365)	(1365)	(1355)	(512)	(1292)
Proteus mirabilis	97%	86%	91%	99%	89%	95%	98%	99%	98%	99%	NED	93%	99%	84%	94%	99%	60%	55%	99%	89%	88%	90%	R	R
	(329)	(557)	(523)	(500)	(594)	(421)	(597)	(508)	(462)	(181)		(243)	(303)	(588)	(541)	(309)	(249)	(161)	(407)	(583)	(597)	(597)		ı
Pseudomonas aeruginosa	R	R	R	94%	R	R	R	91%	87%	R	R	R	60%	91%	98%	94%	R	75%	93%	87%	82%	R	R	R
				(858)				(835)	(837)				(162)	(870)	(874)	(526)		(263)	(569)	(895)	(895)			
Serratia marcesens	R	R	R	60%	R	R	99%	97%	95%		R	R	98%	100%	90%	98%	NED	NED	98%	99%	99%	100%	5%	R
				(30)			(73)	(61)	(59)				(46)	(73)	(59)	(48)			(57)	(73)	(73)	(73)	(31)	
Haemophilus influenzae		57%					100%														100%	68%		
		(78)					(78)														(87)	(78)		

# 2016 Alaska State Antibiogram: Anchorage-Mat-Su Region

The following tables show the proportion of isolates of various bacterial species that tested susceptible to various antibiotics during 2014 and 2015. These data were aggregated from the antibiograms produced by Alaska hospitals in order to create aggregate regional resistance pattern summaries. These antibiograms can be helpful for health care providers in selecting appropriate "presumptive" antimicrobial therapy for their patients until specific individual laboratory test results are available. They can also be helpful for determining antibiotic stewardship priorities within hospitals and emerging resistance patterns in a broader service area.

- **Methodology:** Individual hospitals prepared their own facility antibiograms, which were shared with the Alaska Section of Epidemiology. Aggregated susceptibility percentages were calculated as the proportion of all tested isolates for the region that were susceptible. Values are only reported when more than one facility provided data for the given species-antibiotic combination. Intrinsic resistance is indicated with an "R", following the guidance of CLSI document M100-S24.
- Multi-Drug Resistant Organisms of Note:
  - o Vancomycin-resistant Staphylococcus aureus (VRSA): no cases of VRSA have ever been reported in Alaska. VRSA is reportable to the Alaska Section of Epidemiology.
  - o Carbapenem-resistant Enterobacteriaceae (CRE): there were 7 cases of CRE in Anchorage/Mat-Su residents in 2016.
- Legend:
  - o The top value in each square is the percent of isolates of that species that tested susceptible to that antibiotic.
  - o The lower value in each square indicates the number of tested isolates for that bacteria-antibiotic combination.
  - o "R" indicates intrinsic resistance to that antibiotic, while "S" indicates definitional susceptibility.
  - o "NED" indicates that there was Not Enough Data to report the value: either only one facility reported data for that drug-bug combination or <30 isolates were tested.
- **Limitations:** Individual facilities often use different methods to test for antimicrobial susceptibility, different methods to build their antibiograms, and different antibiotics in their pharmacies. These factors limit interpretation of these data. Additionally, antimicrobial susceptibility testing done in the laboratory does not always predict how effective that drug will be when used to treat a patient. Data are not stratified by infection site, which influences antibiotic choice and effectiveness.
- Contributing Facilities: Thanks to the following facilities for providing data in support of this project:
  - Alaska Native Medical Center
  - Alaska Regional Hospital
  - o Providence Alaska Medical Center
  - Mat-Su Regional Medical Center

Anchorage+ Mat-Su Region data	Penicillin	Ampicillin	Oxacillin	Cefazolin	Ceftriaxone	Cefotaxime	Ciprofloxacin	Levofloxacin	Clindamycin	Erythromycin	Vancomycin	Gentamicin	Gent Syn	Trimethoprim-sulfamethoxazole	Linezolid	Tetracycline	Nitrofurantoin
Total Staphylococcus aureus	NED	NED	59%	56%			63%	63%	85%	39%	99%	98%		98%	100%	96%	99%
			(3555)	(2128)			(788)	(788)	(3555)	(788)	(3555)	(1797)		(3555)	(2767)	(3137)	(2215)
MSSA	NED	NED	S	100%			89%	90%	88%	63%	100%	98%		98%	100%	96%	100%
				(1205)			(433)	(433)	(2104)	(433)	(2104)	(1108)		(2104)	(1671)	(1671)	(1205)
MRSA	NED	NED	R	NED			31%	31%	81%	9%	100%	99%		98%	100%	96%	98%
							(355)	(355)	(1462)	(355)	(1462)	(689)		(1462)	(1107)	(1268)	(1462)
Coag-negative Staphylococcus	NED	NED	46%	37%			55%	55%	53%	35%	100%	83%		58%	100%	92%	100%
			(264)	(74)			(74)	(74)	(264)	(74)	(264)	(220)		(264)	(190)	(224)	(127)
Enterococcus faecalis	99%	100%		R	R	R	91%	95%	R	NED	100%	R	83%	R	99%	NED	99%
	(134)	(505)					(371)	(371)			(505)		(437)		(319)		(439)
Enterococcus faecium	NED	29%		R	R	R	NED	NED	R		53%	R	80%	R	98%	NED	NED
		(55)									(55)		(55)		(55)		
Streptococcus pneumoniae (all)								98%	NED	64%	100%						
								(44)		(44)	(145)						
S. pneumoniae - non-CSF	82%				97%	99%											
	(310)				(288)	(310)											
S pneumoniae - meningitis	70%				95%	97%											
	(310)				(288)	(310)											

Anchorage+ Mat-Su Region data  Species	Amoxicillin+ clavulanic acid	Ampicillin	Ampicillin+Sulbactam	Piperacillin+Tazobactam	Cefazolin	Cefuroxime	Ceftriaxone	Ceftazidime	Cefepime	Cefotaxime	Aztreonam	Gentamicin	Tobramycin	Amikacin	Meropenem	Ciprofloxacin	Levofloxacin	Trimeth+Sulfa	Tetracycline	Nitrofurantoin
Citrobacter freundii	R	R	R	98%	R	R	98%	98%	82%	NED	NED	100%	100%	NED	NED	99%	99%	92%	NED	97%
				(85)			(85)	(85)	(85)			(85)	(85)			(85)	(85)	(85)		(77)
Enterobacter aerogenes	R	R	R	87%	R	R	87%	NED	100%	NED	85%	NED	100%	100%	100%	100%	100%	100%	NED	22%
				(78)			(78)		(78)		(78)		(78)	(78)	(78)	(78)	(78)	(78)		(78)
Enterobacter cloacae	R	R	R	87%	R	R	85%	86%	98%	78%	84%	99%	98%	100%	97%	97%	97%	94%	91%	38%
				(277)			(277)	(215)	(277)	(152)	(215)	(277)	(277)	(215)	(277)	(277)	(277)	(277)	(152)	(277)
Escherichia coli	87%	55%	53%	97%	91%	83%	96%	96%	90%	68%	91%	93%	94%	100%	100%	84%	84%	82%	82%	98%
	(3349)	(5462)	(6005)	(6005)	(6005)	(6005)	(6005)	(2656)	(6005)	(3349)	(5163)	(6005)	(6005)	(4620)	(5462)	(6005)	(6005)	(6005)	(3349)	(5958)
Klebsiella oxytoca	NED		54%	91%	61%	84%	95%	98%	98%	NED	86%	98%	97%	NED	100%	96%	96%	92%	NED	88%
			(169)	(169)	(119)	(169)	(169)	(169)	(169)		(169)	(169)	(169)		(150)	(169)	(169)	(169)		(155)
Klebsiella pneumoniae	98%	R	89%	97%	93%	91%	98%	49%	97%	98%	99%	99%	98%	99%	99%	96%	96%	94%	87%	49%
	(349)		(823)	(823)	(823)	(823)	(823)	(698)	(803)	(349)	(699)	(823)	(823)	(603)	(717)	(823)	(823)	(823)	(349)	(791)
Proteus mirabilis	98%	82%	89%	99%	91%	94%	98%	98%	98%	99%	99%	95%	95%	100%	100%	89%	89%	89%	R	R
	(133)	(271)	(311)	(214)	(311)	(311)	(311)	(275)	(311)	(133)	(275)	(311)	(311)	(235)	(271)	(311)	(311)	(311)		
Pseudomonas aeruginosa	R	R	R	94%	R	R	R	91%	85%	R	NED	91%	98%	93%	93%	89%	83%	R	R	R
				(566)				(506)	(566)			(566)	(566)	(433)	(493)	(566)	(566)			

## 2016 Alaska State Antibiogram: Gulf Coast Region

The following tables show the proportion of isolates of various bacterial species that tested susceptible to various antibiotics during 2014 and 2015. These data were aggregated from the antibiograms produced by Alaska hospitals in order to create aggregate regional resistance pattern summaries. These antibiograms can be helpful for health care providers in selecting appropriate "presumptive" antimicrobial therapy for their patients until specific individual laboratory test results are available. They can also be helpful for determining antibiotic stewardship priorities within hospitals and emerging resistance patterns in a broader service area.

- **Methodology:** Individual hospitals prepared their own facility antibiograms, which were shared with the Alaska Section of Epidemiology. Aggregated susceptibility percentages were calculated as the proportion of all tested isolates for the region that were susceptible. Values are only reported when more than one facility provided data for the given species-antibiotic combination. Intrinsic resistance is indicated with an "R", following the guidance of CLSI document M100-S24.
- Multi-Drug Resistant Organisms of Note:
  - o Vancomycin-resistant Staphylococcus aureus (VRSA): no cases of VRSA have ever been reported in Alaska. VRSA is reportable to the Alaska Section of Epidemiology.
  - o Carbapenem-resistant Enterobacteriaceae (CRE): there was 1 case of CRE in a Gulf Coast resident in 2016.
- Legend:
  - o The top value in each square is the percent of isolates of that species that tested susceptible to that antibiotic.
  - o The lower value in each square indicates the number of tested isolates for that bacteria-antibiotic combination.
  - o "R" indicates intrinsic resistance to that antibiotic, while "S" indicates definitional susceptibility.
  - o "NED" indicates that there was Not Enough Data to report the value: either only one facility reported data for that drug-bug combination or <30 isolates were tested.
- **Limitations:** Individual facilities often use different methods to test for antimicrobial susceptibility, different methods to build their antibiograms, and different antibiotics in their pharmacies. These factors limit interpretation of these data. Additionally, antimicrobial susceptibility testing done in the laboratory does not always predict how effective that drug will be when used to treat a patient. Data are not stratified by infection site, which influences antibiotic choice and effectiveness.
- Contributing Facilities: Thanks to the following facilities for providing data in support of this project:
  - o Central Peninsula Hospital
  - South Peninsula Hospital
  - Providence Valdez Medical Center

Gulf Coast Region data Species	Penicillin	Ampicillin	Oxacillin	Ampicillin-sulbactam	Amoxicillin-clavanulate	Ceftriaxone	Ciprofloxacin	Levofloxacin	Daptomycin	Clindamycin	Erythromycin	Vancomycin	Gentamicin	Trimethoprim-sulfamethoxazole	Linezolid	Tetracycline	Nitrofurantoin	Rifampin
Total Staphylococcus	<b>10%</b>	6%	63%	60%	62%	<b>65%</b>	53%	54%	99%	81%	<b>39%</b>	99%	99%	<b>96%</b>	99%	95% (5.47)	99%	99%
aureus	(547) <b>14%</b>	(182) NED	(547) <b>S</b>	(182) NED	(182) NED	(386) NED	(547) <b>75</b> %	(547) <b>77</b> %	(386) NED	(474) <b>83</b> %	(495) <b>59</b> %	(547) <b>98%</b>	(182) NED	(547) <b>96</b> %	(547) <b>99</b> %	(547) <b>95</b> %	(142) NED	(526) <b>99</b> %
MSSA	(322)					.,,	(322)	(322)	.,,	(292)	(292)	(322)	.,,	(322)	(322)	(322)	.,,	(322)
MRSA	<b>0</b> % (217)	<b>0</b> % (82)	R	<b>4%</b> (82)	<b>1%</b> (82)	<b>3%</b> (148)	<b>15%</b> (217)	<b>17</b> % (217)	<b>99%</b> (148)	<b>78%</b> (195)	<b>5%</b> (195)	<b>99%</b> (217)	<b>98%</b> (82)	<b>96%</b> (217)	<b>100%</b> (217)	<b>94%</b> (217)	<b>100%</b> (104)	<b>100%</b> (204)
Staphylocccus epidermidis	<b>12%</b> (168)		<b>47%</b> (168)	<b>52%</b> (33)	<b>47%</b> (33)	<b>47%</b> (138)	<b>64%</b> (168)	<b>75%</b> (138)	<b>99%</b> (138)	<b>55%</b> (122)	<b>31%</b> (122)	<b>98%</b> (168)	<b>85%</b> (33)	<b>58%</b> (168)	<b>99%</b> (168)	<b>93%</b> (168)	<b>99%</b> (79)	<b>99%</b> (165)
Enterococcus faecalis	<b>99%</b> (234)	<b>99%</b> (234)				R	<b>78%</b> (234)	<b>84%</b> (234)	NED	R	NED	<b>99%</b> (234)	R	R	<b>94%</b> (234)	<b>23%</b> (234)	<b>100%</b> (207)	NED
Group B Streptococcus	<b>100%</b> (29)	S						NED		<b>62%</b> (29)	<b>52%</b> (29)	<b>100%</b> (29)			NED	NED		
Streptococcus pneumoniae	<b>90%</b> (31)							NED		NED	NED	NED		NED		NED		

Gulf Coast Region data	Amoxicillin+ clavanulanic acid	Ampicillin	Ampicillin+Sulbactam	Piperacillin+Tazobactam	Cefazolin	Cefuroxime	Ceftriaxone	Ceftazidime	Cefepime	Gentamicin	Tobramycin	Amikacin	Imipenem	Ciprofloxacin	Levofloxacin	Trimeth+Sulfa	Tetracycline	Nitrofurantoin	
Escherichia coli	90%	60%	64%	98%	92%	96%	99%	99%	100%	92%	94%	99%	99%	85%	85%	77%	81%	97%	
	(1051)	(1051)	(1051)	(1051)	(1051)	(982)	(1051)	(1051)	(303)	(1051)	(1051)	(817)	(817)	(1051)	(1051)	(1051)	(303)	(972)	
Klebsiella pneumoniae	99%	R	88%	98%	97%	99%	100%	100%	100%	99%	99%	100%	100%	98%	99%		92%	46%	
Kiebsiena priedmoniae	(221)		(221)	(221)	(221)	(217)	(221)	(221)	(51)	(221)	(221)	(174)	(174)	(221)	(221)		(51)	(201)	
Proteus mirabilis	93%	84%	86%	100%	89%	98%	97%	100%	NED	89%	89%	98%	93%	80%	86%	90%	R	R	
Froteus Iriii ubilis	(70)	(70)	(70)	(70)	(70)	(66)	(70)	(70)		(70)	(70)	(46)	(46)	(70)	(70)	(70)			
Degudomonas agruainosa	R	R	R	92%	R	R	R	86%	80%	86%	100%	99%	99%	75%	74%	R	R	R	
Pseudomonas aeruginosa   				(88)				(88)	(88)	(88)	(88)	(73)	(73)	(88)	(88)			l	

# 2016 Alaska State Antibiogram: Northern Region

The following tables show the proportion of isolates of various bacterial species that tested susceptible to various antibiotics during 2014 and 2015. These data were aggregated from the antibiograms produced by Alaska hospitals in order to create aggregate regional resistance pattern summaries. These antibiograms can be helpful for health care providers in selecting appropriate "presumptive" antimicrobial therapy for their patients until specific individual laboratory test results are available. They can also be helpful for determining antibiotic stewardship priorities within hospitals and emerging resistance patterns in a broader service area.

- **Methodology:** Individual hospitals prepared their own facility antibiograms, which were shared with the Alaska Section of Epidemiology. Aggregated susceptibility percentages were calculated as the proportion of all tested isolates for the region that were susceptible. Values are only reported when more than one facility provided data for the given species-antibiotic combination. Intrinsic resistance is indicated with an "R", following the guidance of CLSI document M100-S24.
- Multi-Drug Resistant Organisms of Note:
  - o Vancomycin-resistant Staphylococcus aureus (VRSA): no cases of VRSA have ever been reported in Alaska. VRSA is reportable to the Alaska Section of Epidemiology.
  - o Carbapenem-resistant Enterobacteriaceae (CRE): there were no cases of CRE reported in the Northern Region in 2016
- Legend:
  - o The top value in each square is the percent of isolates of that species that tested susceptible to that antibiotic.
  - o The lower value in each square indicates the number of tested isolates for that bacteria-antibiotic combination.
  - o "R" indicates intrinsic resistance to that antibiotic, while "S" indicates definitional susceptibility.
  - o "NED" indicates that there was Not Enough Data to report the value: either only one facility reported data for that drug-bug combination or <30 isolates were tested.
- **Limitations:** Individual facilities often use different methods to test for antimicrobial susceptibility, different methods to build their antibiograms, and different antibiotics in their pharmacies. These factors limit interpretation of these data. Additionally, antimicrobial susceptibility testing done in the laboratory does not always predict how effective that drug will be when used to treat a patient. Data are not stratified by infection site, which influences antibiotic choice and effectiveness.
- Contributing Facilities: Thanks to the following facilities for providing data in support of this project:
  - o Maniilaq Health Center
  - Norton Sound Health Center
  - o Samuel Simmonds Memorial Hospital

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Northern Region data Species	Amoxicillin-clavulanic acid	Ampicillin	Ampicillin-sulbactam	Oxacillin	Ciprofloxacin	Levofloxacin	Clindamycin	Erythromycin	Vancomycin	Gent Syn	Trimethoprim-sulfamethoxazole	Linezolid	Tetracycline	Nitrofurantoin	Rifampin
Total Staphylococcus aureus			NED	69%	69%	68%	79%	54%	95%		99%	99%	97%	100%	99%
				(495)	(495)	(495)	(491)	(216)	(495)		(495)	(495)	(495)	(381)	(393)
MSSA	99%		99%	s	87%	86%	77%	70%	87%		97%	NED	95%	100%	NED
IVISSA	(152)		(152)		(152)	(152)	(150)	(150)	(152)		(152)		(152)	(69)	
MRSA	NED	NED	NED	R		NED	61%	NED	NED		100%	100%	99%	100%	
IVINSA							(67)				(69)	(69)	(69)	(38)	
Coag negative Staphylococcus		NED	NED	45%	84%	84%	60%	NED	97%		83%	98%	93%	95%	98%
				(95)	(95)	(95)	(59)		(95)		(95)	(95)	(95)	(65)	(95)
Enterococcus faecalis		99%			96%	96%	R		100%	78%	R	99%	27%	100%	NED
		(97)			(97)	(97)			(97)	(50)		(97)	(97)	(94)	

Northern Region data Species	Amoxicillin+ clavulanic acid	Ampicillin	Ampicillin+Sulbactam	Piperacillin+Tazobactam	Ticaricillin-clavulanic acid	Cefazolin	Cefuroxime	Ceftriaxone	Ceftazidime	Cefepime	Cefotaxime	Cefotatan	Cefoxitin	Gentamicin	Tobramycin	Ertapenem	Imipenem	Ciprofloxacin	Levofloxacin	Trimeth+Sulfa	Tetracycline	Nitrofurantoin
Escherichia coli	85%	54%	61%	98%	96%	91%	100%	97%	97%	97%	86%	100%	98%	93%	95%	92%	100%	85%	85%	76%	84%	98%
	(982)	(982)	(982)	(982)	(543)	(982)	(804)	(982)	(982)	(982)	(543)	(543)	(543)	(981)	(982)	(543)	(804)	(982)	(982)	(982)	(543)	(973)
Klebsiella pneumoniae	97%	R	89%	100%	NED	98%	100%	100%	99%	100%	98%	100%	98%	99%	98%	NED	100%	97%	96%	99%	NED	62%
	(103)		(103)	(103)		(103)	(67)	(103)	(67)	(103)	(63)	(63)	(63)	(103)	(103)		(67)	(103)	(103)	(103)		(98)
Proteus mirabilis	100%	91%	96%	100%	NED	98%	94%	100%	100%	100%	NED	89%	89%	89%	89%	NED	100%	90%	93%	89%		R
	(45)	(45)	(45)	(45)		(45)	(35)	(45)	(45)	(45)		(45)	(45)	(45)	(45)		(35)	(45)	(45)	(45)		
Pseudomonas aeruginosa	R	R	R	98%			90%	R	94%	91%	NED	NED	94%	NED	94%		NED	87%	85%	R		R
				(47)			(40)		(47)	(47)			(47)		(47)			(47)	(47)			

# 2016 Alaska State Antibiogram: Southeast Region

The following tables show the proportion of isolates of various bacterial species that tested susceptible to various antibiotics during 2014 and 2015. These data were aggregated from the antibiograms produced by Alaska hospitals in order to create aggregate regional resistance pattern summaries. These antibiograms can be helpful for health care providers in selecting appropriate "presumptive" antimicrobial therapy for their patients until specific individual laboratory test results are available. They can also be helpful for determining antibiotic stewardship priorities within hospitals and emerging resistance patterns in a broader service area.

• **Methodology:** Individual hospitals prepared their own facility antibiograms, which were shared with the Alaska Section of Epidemiology. Aggregated susceptibility percentages were calculated as the proportion of all tested isolates for the region that were susceptible. Values are only reported when more than one facility provided data for the given species-antibiotic combination. Intrinsic resistance is indicated with an "R", following the guidance of CLSI document M100-S24.

#### • Multi-Drug Resistant Organisms of Note:

- o Vancomycin-resistant Staphylococcus aureus (VRSA): no cases of VRSA have ever been reported in Alaska. VRSA is reportable to the Alaska Section of Epidemiology.
- o Carbapenem-resistant Enterobacteriaceae (CRE): there was 1 case of CRE reported in a Southeast resident in 2016.

#### Legend:

- o The top value in each square is the percent of isolates of that species that tested susceptible to that antibiotic.
- o The lower value in each square indicates the number of tested isolates for that bacteria-antibiotic combination.
- o "R" indicates intrinsic resistance to that antibiotic, while "S" indicates definitional susceptibility.
- o "NED" indicates that there was Not Enough Data to report the value: either only one facility reported data for that drug-bug combination or <30 isolates were tested.
- **Limitations:** Individual facilities often use different methods to test for antimicrobial susceptibility, different methods to build their antibiograms, and different antibiotics in their pharmacies. These factors limit interpretation of these data. Additionally, antimicrobial susceptibility testing done in the laboratory does not always predict how effective that drug will be when used to treat a patient. Data are not stratified by infection site, which influences antibiotic choice and effectiveness.
- Contributing Facilities: Thanks to the following facilities for providing data in support of this project:
  - o Bartlett Regional Hospital
  - o Peacehealth Ketchikan Medical Center

o SEARHC

Petersburg Medical Center Sitka Community Hospital Wrangell Medical Center

outheast egion data Species	Penicillin	Ampicillin	Oxacillin	Ampicillin-sulbactam	Amoxicillin-clavanulate	Ceftriaxone	Ciprofloxacin	Levofloxacin	Moxifloxacin	Daptomycin	Clindamycin	Erythromycin	Vancomycin	Gentamicin	Gent Syn	Trimethoprim-sulfamethoxazole	Linezolid	Tetracycline	Nitrofurantoin	Quinupristin-dalfopristin	Rifampin	Strep syn
Total Staphylococcus	10%	0%	57%	50%	52%	51%	63%	63%	70%	100%	85%	47%	100%	99%		98%	99%	97%	99%	99%	99%	
aureus	(606)	(101)	(908)	(104)	(104)	(103)	(648)	(648)	(365)	(103)	(906)	(906)	(866)	(386)		(866)	(363)	(826)	(811)	(80)	(545)	
MSSA	<b>11%</b> (231)	<b>0%</b> (51)	S	<b>96%</b> (54)	<b>100%</b> (54)	<b>100%</b> (53)	<b>87%</b> (218)	<b>85%</b> (218)		<b>100%</b> (53)	<b>90%</b> (395)	<b>72%</b> (395)	<b>100%</b> (377)	<b>100%</b> (83)		<b>98%</b> (377)	<b>99%</b> (231)	<b>71%</b> (347)	<b>99%</b> (335)	<b>100%</b> (53)	<b>99%</b> (231)	
MRSA	<b>0</b> % (50)	<b>0</b> % (50)	R	<b>0</b> % (50)	<b>0</b> % (50)	<b>0</b> % (50)	<b>28%</b> (187)	<b>29%</b> (187)		<b>100%</b> (50)	<b>72%</b> (186)	<b>13%</b> (186)	<b>100%</b> (269)	<b>98%</b> (60)		<b>99%</b> (246)	<b>100%</b> (132)	<b>98%</b> (269)	<b>98%</b> (233)	<b>98%</b> (50)	<b>100</b> % (132)	
Coag-negative	19%		61%				72%	72%	73%		64%	39%	99%	96%		68%	100%	80%	100%		98%	
Staphylocccus	(63)		(98)				(74)	(74)	(52)		(81)	(82)	(98)	(53)		(91)	(58)	(98)	(67)		(64)	
Enterococcus faecalis	<b>100%</b> (127)	<b>100</b> % (141)					<b>92%</b> (137)	<b>95%</b> (123)		<b>100%</b> (40)		<b>6%</b> (114)	<b>100</b> % (141)		<b>88%</b> (118)		<b>100%</b> (113)	<b>21%</b> (127)	<b>100</b> % (127)		<b>70%</b> (40)	<b>93%</b> (105)

Southeast Region data	Amoxicillin+ clav	Ampicillin	Ampicillin-sulbactam	Piperacillin-tazobactam	Cefazolin	Cefuroxime	Ceftriaxone	Ceftazidime	Cefepime	Gentamicin	Tobramycin	Amikacin	lmipenem	Ciprofloxacin	Levofloxacin	Trimeth + sulfa	Tetracycline	Nitrofurantoin
Escherichia coli	90%	60%	64%	98%	92%	96%	99%	99%	100%	92%	94%	99%	99%	85%	85%	77%	81%	97%
LSCHEHCHIA COII	(1051)	(1051)	(1051)	(1051)	(1051)	(982)	(1051)	(1051)	(303)	(1051)	(1051)	(817)	(817)	(1051)	(1051)	(1051)	(303)	(972)
Klebsiella pneumoniae	99%	R	88%	98%	97%	99%	100%	100%	100%	99%	99%	100%	100%	98%	99%	94%	92%	46%
	(221)		(221)	(221)	(221)	(217)	(221)	(221)	(51)	(221)	(221)	(174)	(174)	(221)	(221)	(221)	(51)	(201)
Proteus mirabilis	93%	84%	86%	100%	89%	98%	97%	100%	NED	89%	89%	98%	93%	80%	86%	90%	R	R
FIOLEUS IIII UDIIIS	(70)	(70)	(70)	(70)	(70)	(66)	(70)	(70)		(70)	(70)	(46)	(46)	(70)	(70)	(70)		
Pseudomonas aeruginosa	R	R	R	92%	R	R	R	86%	80%	86%	100%	99%	99%	75%	74%	R	R	R
- seadomonas deraginosa				(88)				(88)	(88)	(88)	(88)	(73)	(73)	(88)	(88)			

## 2016 Alaska State Antibiogram: Southwest Region

The following tables show the proportion of isolates of various bacterial species that tested susceptible to various antibiotics during 2014 and 2015. These data were aggregated from the antibiograms produced by Alaska hospitals in order to create aggregate regional resistance pattern summaries. These antibiograms can be helpful for health care providers in selecting appropriate "presumptive" antimicrobial therapy for their patients until specific individual laboratory test results are available. They can also be helpful for determining antibiotic stewardship priorities within hospitals and emerging resistance patterns in a broader service area.

- **Methodology:** Individual hospitals prepared their own facility antibiograms, which were shared with the Alaska Section of Epidemiology. Aggregated susceptibility percentages were calculated as the proportion of all tested isolates for the region that were susceptible. Values are only reported when more than one facility provided data for the given species-antibiotic combination. Intrinsic resistance is indicated with an "R", following the guidance of CLSI document M100-S24.
- Multi-Drug Resistant Organisms of Note:
  - o Vancomycin-resistant Staphylococcus aureus (VRSA): no cases of VRSA have ever been reported in Alaska. VRSA is reportable to the Alaska Section of Epidemiology.
  - o Carbapenem-resistant Enterobacteriaceae (CRE): there were no cases of CRE reported in Southwest residents in 2016.
- Legend:
  - o The top value in each square is the percent of isolates of that species that tested susceptible to that antibiotic.
  - o The lower value in each square indicates the number of tested isolates for that bacteria-antibiotic combination.
  - o "R" indicates intrinsic resistance to that antibiotic, while "S" indicates definitional susceptibility.
  - o "NED" indicates that there was Not Enough Data to report the value: either only one facility reported data for that drug-bug combination or <30 isolates were tested.
- **Limitations:** Individual facilities often use different methods to test for antimicrobial susceptibility, different methods to build their antibiograms, and different antibiotics in their pharmacies. These factors limit interpretation of these data. Additionally, antimicrobial susceptibility testing done in the laboratory does not always predict how effective that drug will be when used to treat a patient. Data are not stratified by infection site, which influences antibiotic choice and effectiveness.
- Contributing Facilities: Thanks to the following facilities for providing data in support of this project:
  - o Yukon-Kuskokwim Health Center
  - Bristol Bay Area Health Center

Southwest Region data Species	Penicillin	Ampicillin	Oxacillin	Levofloxacin	Clindamycin	Erythromycin	Vancomycin	Trimethoprim-sulfamethoxazole	Tetracycline	Nitrofurantoin
Total Staphylococcus aureus	10%	NED	57%	66%	97%	47%	99%	99%	97%	NED
	(738)		(738)	(738)	(725)	(725)	(738)	(738)	(738)	
MSSA	17%	NED	S	85%	96%	69%	99%	99%	99%	NED
	(416)			(416)	(406)	(406)	(416)	(416)	(416)	
MRSA	0%	NED	R	41%	98%	19%	99%	100%	94%	NED
	(322)			(322)	(319)	(319)	(322)	(322)	(322)	
Coag-negative	17%		50%	90%	75%	38%	98%	79%	91%	95%
Staphylococcus	(294)		(294)	(294)	(157)	(157)	(294)	(292)	(294)	(137)
Enterococcus faecalis	100%	100%		96%		NED	98%		36%	98%
	(56)	(56)		(56)			(56)		(56)	(50)

Southwest Region data Species	Amoxicillin+ clavulanic acid	Ampicillin	Piperacillin+Tazobactam	Cefazolin	Ceftriaxone	Gentamicin	Ciprofloxacin	Levofloxacin	Trimeth+Sulfa	Tetracycline	Nitrofurantoin
Enterobacter cloacae	0%	0%	94%	0%	91%	100%	100%	100%	97%	91%	NED
	(33)	(33)	(33)	(33)	(33)	(33)	(33)	(33)	(33)	(33)	
Escherichia coli	85%	47%	99%	91%	97%	92%	84%	84%	73%	78%	99%
	(1290)	(1290)	(1290)	(1290)	(1290)	(1290)	(1290)	(1290)	(1290)	(1290)	(1244)
Klebsiella pneumoniae	94%	0%	99%	97%	99%	100%	96%	99%	96%	89%	63%
Nicosiciia pricumoniae	(72)	(72)	(72)	(72)	(72)	(72)	(72)	(72)	(72)	(72)	(56)
Proteus mirabilis	98%	95%	100%	98%	100%	NED	100%	100%	98%	0%	0%
. Toteus IIII ubiis	(56)	(56)	(56)	(56)	(56)		(56)	(56)	(56)	(56)	(56)