# ANTIBIOTIC STEWARDSHIP NEWPORT HOSPITAL

AN ARGUMENT FOR POOLING REGIONAL DATA
WHEN CONSTRUCTING A CUMULATIVE ANTIBIOGRAM

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## Pooling of Antibiogram Data Rationale

- A more statistically powerful data set (narrow 95% confidence interval) by increasing sample size
- More reliable detection of emerging resistant organisms
- Allows for inclusion of a greater number of organisms above the 30 isolate threshold

#### Cumulative Antibiogram

- Offers a guide to empiric antibiotic therapy at an institution ("Use local data; know your antibiogram")
- Can offer clues towards early detection of emerging drug resistant organism
- Can reliably inform infection control mechanisms

# Cumulative Antibiogram CLSI Recommendations

- Analyze and present data at least annually
- Include only species in which there are at least 30 isolates
- Diagnostic, not surveillance cultures
- Include the results for all antimicrobials tested
- Include only the first isolate for the patient tested

#### Newport Hospital

- Our current antibiogram report is a spreadsheet with the sensitivity data on 43 organisms
- Only four organisms meet the criteria for inclusion (at least 30 isolates reported)
- E coli, Enterococcus faecalis, Klebsiella pneumonia, Staph aureus

Just because the sample size is 30 and approximates a "normal distribution", does this necessarily imply that we can derive statistically meaningful conclusions from the data?

### NEWPORT HOSPITAL ANTIBIOGRAM

Organism	N	Sensitivity Levofloxacin
Pseudomonas aeruginosa	30*	80%

#### NEWPORT HOSPITAL ANTIBIOGRAM

Is the 80% sensitivity of Pseudomonas aeruginosa to Levofloxacin a meaningful number?

 Problem: Confidence interval testing tells us with 95% probability that the sensitivity could be represented anywhere between 61% and 92%

Susceptibl	e or	Resistant	Rate
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c 1																			
Sample Size		1	10%	2	0%	30	0%	40	0%	50	%	6	0%	70	)%	80	%	90	0%
	10	0	45	3	56	7	65	12	74	19	81	26	88	35	93	44	97	55	100
	20	1	32	6	44	12	54	19	64	27	73	36	81	46	88	56	94	68	99
	30	2	27	8	39	15	49	23	59	31	69	41	77	5		61	92	73	98
	40	3	24	9	36	17	47	25	57	34	66	43	75	53	83	64	91	76	97
	50	3	22	10	34	18	45	26	55	36	64	45	74	55	82	66	90	78	97
	60	4	21	11	32	19	43	28	53	37	63	47	72	57	81	68	89	79	96
	70	4	20	11	31	20	42	28	52	38	62	48	72	58	80	69	89	80	96
	80	4	19	12	30	20	41	29	52	39	61	48	71	59	80	70	88	81	96
	90	5	18	12	30	21	41	30	51	39	61	49	70	59	79	70	88	82	95
	100	5	18	13	29	21	40	30	50	40	60	50	70	60	79	71	87	82	95
	200	6	15	15	26	24	37	33	47	43	57	53	67	63	76	74	85	85	94
4	400	7	13	16	24	26	35	35	45	45	55	55	65	65	74	76	84	87	93
(	600	8	13	17	23	26	34	36	44	46	54	56	64	66	74	77	83	87	92
10	000	8	12	18	23	27	33	37	43	47	53	57	63	67	73	77	82	88	92

Confidence intervals were calculated using the Clopper-Pearson method.

$$\sum_{k=0}^{k} {n \choose k} p_{UB}^{k} (1 - p_{UB})^{n-k} = \frac{\alpha}{2}$$

$$\sum_{k=x}^{n} {n \choose k} p_{LB}^{k} (1 - p_{LB})^{n-k} = \frac{\alpha}{2}$$

Clopper Pearson method of calculating confidence interval

#### NEWPORT HOSPITAL ANTIBIOGRAM

- If the sample size were N=200, the confidence interval narrows significantly and the 80% sensitivity could be represented anywhere between 74% and 85%.
- More statistically meaningful data?

Susceptibl	e or l	Resistant	Rate

C 1																			
Sample Size		1	10%	2	0%	30	0%	40	0%	50	%	6	0%	70	)%	80	1%	90	0%
	10	0	45	3	56	7	65	12	74	19	81	26	88	35	93	44	97	55	100
	20	1	32	6	44	12	54	19	64	27	73	36	81	46	88	56	94	68	99
	30	2	27	8	39	15	49	23	59	31	69	41	77	51	85	61	92	73	98
	40	3	24	9	36	17	47	25	57	34	66	43	75	53	83	64	91	76	97
	50	3	22	10	34	18	45	26	55	36	64	45	74	55	82	66	90	78	97
	60	4	21	11	32	19	43	28	53	37	63	47	72	57	81	68	89	79	96
	70	4	20	11	31	20	42	28	52	38	62	48	72	58	80	69	89	80	96
	80	4	19	12	30	20	41	29	52	39	61	48	71	59	80	70	88	81	96
	90	5	18	12	30	21	41	30	51	39	61	49	70	59	79	70	88	82	95
1	.00	5	18	13	29	21	40	30	50	40	60	50	70	60	79	71	87	82	95
2	200	6	15	15	26	24	37	33	47	43	57	53	67	63		74	85	85	94
4	100	7	13	16	24	26	35	35	45	45	55	55	65	65	74	76	84	87	93
6	600	8	13	17	23	26	34	36	44	46	54	56	64	66	74	77	83	87	92
10	000	8	12	18	23	27	33	37	43	47	53	57	63	67	73	77	82	88	92

Confidence intervals were calculated using the Clopper-Pearson method.

#### CONSENSUS STATEMENT: CLSI

• Combining data from several facilities located in the same geographic area is another way to circumvent the concern about having a small number of isolates available for analysis.

Analysis and Presentation of Cumulative Antibiograms: A New Consensus Guideline from the Clinical and Laboratory Standards Institute

## ANTIBIOTIC STEWARDSHIP NHHS

- It is likely that with pooling of our antibiogram data with other regional facilities, we could add three other organisms to our data set
- Proteus mirabilis (N=18), Pseudomonas aeruginosa (N=24), Staph epidermidis (N=23)

# ANTIBIOTIC STEWARDSHIP NHHS SURVEILLANCE FOR EMERGING ANTIBIOTIC RESISTANCE

#### EMERGING ANTIBIOTIC RESISTANCE

Organism	N	Sensitivity Ciprofloxacin
Enterococcus faecalis	50*	90%

#### EMERGING ANTIBIOTIC RESISTANCE

• Hypothetical: What If the following year the sensitivity of Enterococcus faecalis to Ciprofloxacin drops from 90% to 80%? Would this be an indication of emerging resistance?

#### DETERMINING EMERGING ANTIBIOTIC RESISTANCE

- Hypothesis testing shows that the sensitivity of Enterococcus faecalis to Ciprofloxacin can vary from 90% to 73% by chance alone when the sample N=50. (p<.05)
- If the sample size were 200, a drop in the sensitivity from 90% to 82% would be statistically significant

Table H2. Table to Use if % Susceptible Decreases

				Sample S	ize			
Initial %S	10	20	50	100	200	400	600	1000
98	-	-	84	90	93	95	95	96
95	-	65	78	85	89	91	92	92
90	30	55	72	78	82	85	86	87
80	20	45	60	66	71	73	75	76
70	10	30	48	55	60	63	64	65
60	0	20	38	45	49	52	54	55
50	0	15	28	35	39	42	44	45
40	NS	5	20	25	30	33	34	35
30	NS	0	12	17	20	23	24	25
20	NS	NS	4	9	12	14	15	16
10	NS	NS	NS	2	4	5	6	7

NS - not significant

## Regional Pooling of Antibiogram Data

- •For a smaller hospital to produce a useful antibiogram with sufficient statistical power for validation, one option is to increase the sample size by pooling data from regional facilities
- CLSI supports this concept

