								Gram-Po	ositive O	rganisms		0	currence	2										
Hov	v to Use:				-		BETA	STREPT	тососс	US GROUP	В		30	_										
						ENTERO	occus			E. faecalis &	k E. faec	ium)	821											
-Clic	k on the desired organism to hig	hlight								FAECALIS			312											
the	antibiotic sensitivities below.				- 1				ROCOCU	FAECIUM			202 307											
					- 1		CTA			AUREUS (all	,		716											
To	select for multiple organisms, hol	d ctrl			- 1					AUREUS-MRS			242											
-10	select for multiple organisms, no	u cui			- 1					AUREUS-MS			475											
					- 1	ST				SULASE NEG			383											
						STRE	РТОСО	CUS AN	NGINOS	US (MILLERI) GROUI	•	85											
Gran	n-Positive Organisms	n	96	n	%	n	%	n	96	n %	r	%	n	%	n	%	n	%	n	%	n	%	n	%
																								100
ETA STREPTOCOCC				30	40%					2							30	100%					30	
NTEROCOCCUS SPE	(including E. faecalis & E. faecium)	819	67%	30	40%	173	95%			11,	29			75%			316	61%		20%			821	67
ENTEROCOC	(including E. faecalis & E. faecium) CUS FAECALIS	312	100%	30	40%	173 80	95% 96%				5	4 1009	170	75% 99%						20% 18%			821 312	67 88
NTEROCOCCUS SPE ENTEROCOC ENTE	(including E. faecalis & E. faecium) CUS FAECALIS ROCOCCUS FAECALIS-VRE	312 36	100% 100%	30	40%	80	96%				5		170	99%			316 143	61% 100%	175	18%			821 312 36	67° 88°
ENTEROCOCCUS SPP ENTEROCOC ENTE ENTE	(including E. faecalis & E. faecium) CUS FAECALIS ROCOCCUS FAECALIS-VRE ROCOCCUS FAECALIS-VSE	312 36 276	100% 100% 100%	30	40%	80 68	96% 97%				3	4 1009 6 1009	170 5 151	99% 99%			316 143 126	61% 100% 100%	175	18%			821 312 36 276	67 ⁹ 88 ⁹ 0 100
ENTEROCOCUS SPE ENTEROCOC ENTE ENTEROCOC	(including E. faecalis & E. faecium) CUS FAECALIS ROCOCCUS FAECALIS-VRE ROCOCCUS FAECALIS-VSE CUS FAECIUM	312 36 276 202	100% 100% 100% 8%	30	40%	80 68 92	96% 97% 93%				15	4 1009 6 1009 9 999	170 5 151 6 84	99% 99% 24%			316 143 126 118	61% 100% 100% 9%	175 154 90	18% 19% 19%			821 312 36 276 202	67 ⁹ 88 ⁹ 0 100 24 ⁹
ENTEROCOCCUS SPE ENTEROCOC ENTE ENTEROCOC ENTEROCOC	(including E. faecalis & E. faecium) CUS FAECALIS ROCOCCUS FAECALIS-VRE ROCOCCUS FAECALIS-VSE CUS FAECIUM ROCOCCUS FAECIUM-VRE	312 36 276 202 153	100% 100% 100% 8% 1%	30	40%	80 68	96% 97%				3	4 1009 6 1009 9 999	170 5 151 6 84	99% 99%			316 143 126 118 88	61% 100% 100% 9% 0%	175	18%			821 312 36 276 202 154	67° 88° 0° 100° 24°
ENTEROCOCCUS SPP ENTEROCOC ENTE ENTE ENTEROCOC ENTE ENTEROCOC	(including E. faecalis & E. faecium) CUS FAECALIS ROCOCCUS FAECALIS-VRE ROCOCCUS FAECALIS-VSE CUS FAECIUM ROCOCCUS FAECIUM-VRE ROCOCCUS FAECIUM-VSE	312 36 276 202 153 50	100% 100% 100% 8% 1% 30%	30	40%	80 68 92	96% 97% 93%				15 15	4 1009 6 1009 9 999 62 999	170 151 151 84 65	99% 99% 24% 25%			316 143 126 118 88 31	61% 100% 100% 9% 0% 35%	175 154 90 70	18% 19% 19% 17%			821 312 36 276 202 154 49	679 889 00 1000 249 00 1000
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ENTEROCOCUS SPE ENTEROCOC ENTE ENTEROCOC ENTE ENTEROCOC ENTE ENTEROCOC	(Including E. faecalis & E. faecium) CUS FAECALIS ROCOCCUS FAECALIS-VRE ROCOCCUS FAECALIS-VSE CUS FAECIUM ROCOCCUS FAECIUM-VRE ROCOCCUS FAECIUM-VSE CUS SPE UREUS (all)	312 36 276 202 153 50	100% 100% 100% 8% 1% 30%	666	68%	68 92 73	96% 97% 93% 95%				15 15 15 15 4	4 1009 6 1009 9 999 6 999 12 999 14 989 16 1009	151 151 84 65 65 250 50	99% 99% 24% 25%		66%	316 143 126 118 88 31	61% 100% 100% 9% 0% 35%	175 154 90 70 250 569	18% 19% 19% 17% 23% 91%			821 312 36 276 202 154 49 307 252	67 ⁹ 88 ⁸ 0 100 24 ⁹ 0 100 73 ⁹ 100
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ENTEROCOCCUS SPP ENTEROCOC ENTE ENTEROCOC ENTE ENTEROCOC ENTE ENTEROCOC CAPHYLOCOCCUS A STAPHYLOC STAPHYLOCO	(Including E. faecalis & E. faecium) CUS FAECALIS ROCOCCUS FAECALIS-VPE ROCOCCUS FAECALIS-VSE CUS FAECIUM ROCOCCUS FAECIUM-VPE ROCOCCUS FAECIUM-VSE CUS SPP LUREUS (all) LUREUS (all) LUREUS (all)	312 36 276 202 153 50	100% 100% 100% 8% 1% 30%	666	68% 56%	68 92 73 73 61	96% 97% 93% 95% 100%	241	98% 99%	232 3 472 9	15 15 15 15 15 4 4 18% 3	4 1009 6 1009 9 999 6 999 12 999 14 989 16 1009	151 6 84 6 65 6 250 6 50	99% 99% 24% 25% 76% 98%	242	0%	316 143 126 118 88 31	61% 100% 100% 9% 0% 35%	175 154 90 70 250 569 186	18% 19% 19% 17% 23% 91% 85%	241	94%	821 312 36 276 202 154 49 307 252	67 ⁹ 88 ⁸ 0 100 24 ⁹ 0 100 73 ⁹ 100

Presentation Type:

Poster Presentation - Poster Presentation **Subject Category:** Antibiotic Stewardship

Creating an electronic antibiogram using visualization software: Easily updatable and removes the need for yearly manual review

Ashley Dauphin; Christopher McCoy; Robert Bowden; Matthew Lee; Howard Gold and Ryan Chapin

Background: Previously, our hospital manually built a static antibiogram from a surveillance system (VigiLanz) culture report. In 2019, a collaboration between the antimicrobial stewardship team (AST) and the infection control (IC) team set out to leverage data automation to create a dynamic antibiogram. The goal for the antibiogram was the ability to easily distribute and update for hospital staff, with the added ability to perform advanced tracking and surveillance of organism and drug susceptibilities for AST and IC. By having a readily available, accurate, and Clinical and Laboratory Standards Institute (CLSI)-compliant antibiogram, clinicians have the best available data on which to base their empiric antibiotic decisions. Methods: First, assessment of required access to hospital databases and selection of a visualization software (MS Power BI) was performed. Connecting SQL database feeds to Power BI enabled creation of a data model using DAX and M code to comply with the CLSI, generating the first isolate per patient per year. Once a visual antibiogram was created, it was validated against compiled antibiograms using data from the microbiology laboratory middleware (bioMerieux, Observa Integrated Data Management Software). This validation process uncovered some discrepancies between the 2 reference reports due to cascade reporting of susceptibilities. The Observa-derived data were used as the source of truth. The antibiogram prototype was presented to AST/IC members, microbiology laboratory leadership, and other stakeholders to assess functionality. Results: Following feedback and revisions by stakeholders, the new antibiogram was published on a hospital-wide digital platform (Fig. 1). Clinicians may view the antibiogram at any time on desktops from a firewall (or password)-protected intranet. The antibiogram view defaults to the current calendar year and users may interact with the antibiogram rows and columns without disrupting the integrity of the background databases or codes. Each year, simple refreshing of the Power BI antibiogram and changing of the calendar year allows us to easily and accurately update the antibiogram on the hospital-wide digital platform. **Conclusions:** This interdisciplinary collaboration resulted in a new dynamic, CLSI-compliant antibiogram with improved usability, increased visibility, and straightforward updating. In the future, a mobile version of the antibiogram may further enhance accessibility, bring more useful information to providers, and optimize AST/IC guidelines and education.

Disclosures: None

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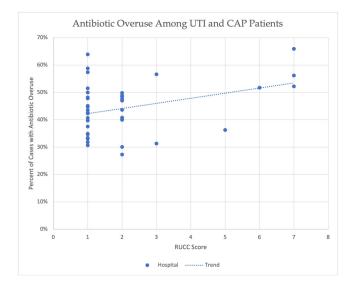
Presentation Type:

Poster Presentation - Poster Presentation **Subject Category:** Antibiotic Stewardship

Identifying the relationship between hospital rurality and antibiotic overuse

Hannah Hardin; Valerie Vaughn; Andrea White; Jennifer Horowitz; Elizabeth McLaughlin; Julia Szymczak; Lindsay Petty; Anurag Malani; Scott Flanders and Tejal Gandhi

Background: Antibiotic overuse and the resulting patient outcomes span all hospitals. However, although antibiotic stewardship can improve antibiotic use, effective stewardship programs require expertise and an infrastructure that are not present in all hospitals. Rural hospitals have less access to resources, infectious disease expertise, and participation in academic research. Thus, we compared antibiotic overuse at discharge between rural and nonrural hospitals for patients diagnosed with community-associated pneumonia (CAP) or urinary tract infection (UTI)—the 2 most common hospital infections. Methods: To determine whether antibiotic overuse at discharge was higher among rural versus nonrural hospitals, we analyzed data from a 41-hospital prospective cohort of patients treated for CAP or UTI between July 1, 2017, and July 30, 2019, in Michigan. Antibiotic overuse was defined as treatment that was unnecessary (ie, patient did not have an infection), excessive (ie, duration >4 days for CAP), or included suboptimal fluoroquinolone use (ie, safer alternative available). Overuse was determined based on patient risk



factors, symptoms, allergies, diagnostic results, and time to stability. Hospital rurality was defined using the Rural-Urban Continuum Codes (RUCC) score. We defined rural as a score ≥4 and very rural as a score of 7–9. We used t tests to compare the mean percentage of patients with antibiotic overuse at discharge between nonrural and rural (and very rural) hospitals. Results: Across 41 hospitals, we included 23,449 patients with CAP or UTI. There were 5 rural (and 3 very rural) hospitals with 2,039 (and 1,082) patients. Antibiotic overuse at discharge was present in 43.1% of patient cases in nonrural hospitals, 52.5% in rural hospitals (P = .04 vs nonrural) and 58.1% in very rural hospitals (P = .007 vs nonrural). Compared to nonrural hospitals, the mean percentage of cases with antibiotic overuse at discharge in rural hospitals was 9.4% higher (15.1% higher in very rural hospitals). Results were similar in a subgroup analysis of only patients with UTI (47.0% in rural vs 37.5% in nonrural, mean difference, 9.5%; P = .03) but were not statistically significant in patients with CAP (53.8% vs 48.0%, respectively; mean difference, 5.8%; P = 0.23). Conclusions: In this retrospective study, rural hospitals—especially very rural hospitals, had higher rates of antibiotic overuse at discharge than nonrural hospitals. Our findings suggest that antibiotic stewardship interventions tailored toward the unique differences in infrastructure, resources, and needs of rural hospitals are essential to community health.

Disclosures: None

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Presentation Type:

Poster Presentation - Poster Presentation Subject Category: Antibiotic Stewardship

Validation of antibiotic stewardship metrics for genitourinary infection management in Veterans Affairs outpatient settings

Jordan Braunfeld; Matthew Samore; Jacob Crook; McKenna Nevers; Kelly Echevarria; Ben Brintz; Matthew Goetz and Karl Madaras-Kelly

Background: Diagnosis and management of suspected urinary tract infection (UTI) in outpatient settings has been shown to be suboptimal. We previously developed a set of stewardship metrics for UTIs based on electronic health record (EHR) data (Antimicrobial Stewardship & Healthcare

Tier	Antibiotic	Cases	Age (SD)	Female	Reviewer	Reviewer
	Prescribed	reviewed (N)		(%)	diagnosed	recommended
					GU infection,	antibiotics, N (%)
					N (%)	
1/2	Yes	33	63.1 (15)	8 (24.2)	23 (69.7)	20 (60.6)
	No	45	64.7 (18.3)	6 (13.3)	3 (6.7)	5 (11.1)
3	Yes	29	71.4 (13.1)	0	2 (6.9)	4 (13.8)
	No	41	71.5 (11.6)	4 (9.8)	1 (2.4)	0

Epidemiology 2022;2 suppl 1:S5-S6. doi:10.1017/ash.2022). A tier-based approach was used to more fully capture antibiotic use associated with genitourinary (GU) symptoms and diagnoses. Herein we report a preliminary analysis of validity and reliability of these metrics based on chart abstraction. Methods: The study cohort consisted of patients who visited Veterans Affairs emergency departments or primary care clinics between 2015 and 2022 and who had a GU diagnosis based on International Classification of Disease, Tenth Revision (ICD-10) codes, divided into 3 categories: tier 1 (antibiotics always indicated), tier 2 (antibiotics sometimes indicated), and tier 3 (antibiotics not indicated). Visits related to urological procedures, nontarget settings, or concomitant non-GU infections were excluded. Cases were randomly sampled for manual review from within 8 strata based on tier, use of antibiotics, and visit type. An infectious disease physician and pharmacist abstracted charts using a standardized data-collection instrument. Clinical judgments regarding diagnosis and treatment were recorded on a Likert scale without knowledge of how the patient was managed. The intraclass correlation coefficient (ICC) was used to estimate interrater reliability. Results: To date, 148 cases have been reviewed (50 by both reviewers). Mean (SD) age was 67.5 (15.3) years and 12.2% were female. In a majority of tier 1 and 2 visits in which antibiotics were given, the reviewers found evidence for GU infection (69.7%) and favored prescribing of antibiotics (60.6%) (Table). In contrast, most patients in the tier 3 category who received antibiotics were judged to have noninfectious conditions (eg, benign prostatic hypertrophy) and to not require antibiotics. In the subset of records examined by both reviewers, the interrater reliability of judgments of whether antibiotics were warranted was good (ICC = .704). **Conclusions:** This preliminary validation provides support for a tier-based approach for stewardship metrics for GU conditions that relies upon electronic data to identify patients for whom antibiotics are generally not indicated.

Disclosures: None

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Understanding clinician perspectives on antibiotic associated adverse events to inform feedback

Jerald Cherian; George Jones; Taylor Helsel; Zunaira Virk; Alejandra Salinas; Suzanne Grieb; Sara Keller; Pranita Tamma and Sara Cosgrove

Background: Feedback regarding antibiotic-associated adverse events (ABX-AEs) may assist clinicians with antibiotic decision making. We

Table. Categorization of Antibiotic-Associated Adverse Events by Degree of Clinical Concern

Prespecified Categorization	Votes (n)*						
Very Concerning	Very	Moderately	Mildly				
very concerning	Concerning	Concerning	Concerning				
Nephrotoxicity – Requiring dialysis	12	-					
Clostridioides difficile infection – Severe	12	-	-1				
Neuropathy	12	-	-				
Stevens-Johnson syndrome	12	-	-				
Anaphylaxis	12	-	-				
DRESS Syndrome	11	1	-				
Moderately Concerning							
Nephrotoxicity – Not requiring dialysis	-	11	1				
Clostridioides difficile infection – Non-severe	-	10	2				
Hepatotoxicity	1	4	4				
Encephalopathy	2	9	-				
Seizures	7	5					
Hemolytic anemia	-	10	-3				
Neutropenia	1	9	-				
Thrombocytopenia	1	8	1				
Prolonged QTc	1	10	-				
Mildly Concerning							
Diarrhea, nausea, or emesis		1	10				
Non-hives rash	-	-	10				
Myositis	-	-	10				